

The Baseline and Elimination Model Equations, Derivations and Parameter Values

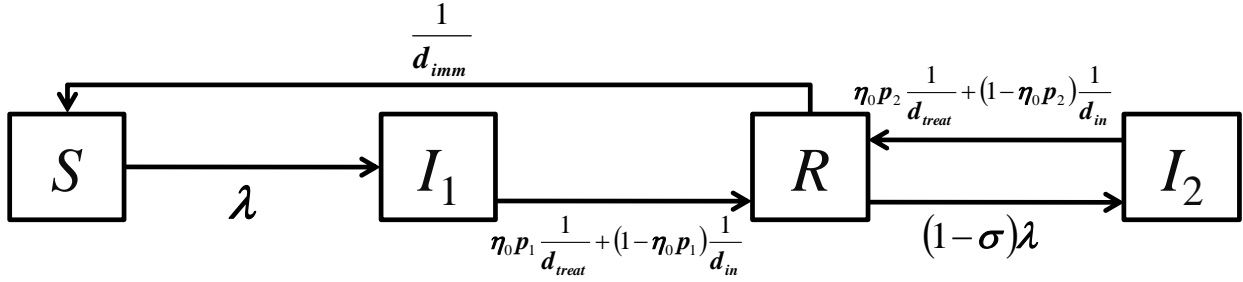
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Baseline model

The diagram shows the compartmental structure of the model with time dependent variables: S (uninfected and non-immune); I_1 (infected with no prior immunity); R (uninfected with immunity); I_2 (infected with prior immunity). Model output variables are: C (clinical infections); P (proportion of people receiving treatment or selective pressure); C_c (cumulative clinical infections); P_c (cumulative selective pressure).



The equations below describe this system:

$$\begin{aligned}
 S' &= \frac{N}{L} - \left(\lambda + \frac{1}{L} \right) S + \frac{1}{d_{imm}} R \\
 I_1' &= \lambda S - \left(\eta_0 p_1 \frac{1}{d_{treat}} + (1 - \eta_0 p_1) \frac{1}{d_{in}} + \frac{1}{L} \right) I_1 \\
 I_2' &= \lambda R - \left(\eta_0 p_2 \frac{1}{d_{treat}} + (1 - \eta_0 p_2) \frac{1}{d_{in}} + \frac{1}{L} \right) I_2 \\
 R' &= \left(\eta_0 p_1 \frac{1}{d_{treat}} + (1 - \eta_0 p_1) \frac{1}{d_{in}} \right) I_1 + \left(\eta_0 p_2 \frac{1}{d_{treat}} + (1 - \eta_0 p_2) \frac{1}{d_{in}} \right) I_2 - \left[\lambda + \frac{1}{d_{imm}} + \frac{1}{L} \right] R \\
 \lambda &= R_0 \left(\frac{1}{L} + \frac{1}{d_{treat}} \right) \frac{(I_1 + I_2)}{N} & p_{res} &= \frac{\frac{1}{d_{treat}} - \frac{1}{d_{treat0}}}{\frac{1}{d_{in}} - \frac{1}{d_{treat0}}} \\
 C &= p_1 I_1 + p_2 I_2 & \tau' &= -k P p_{res} \left(\tau - \frac{1}{d_{in}} \right) \\
 P &= \frac{\eta_0 (p_1 I_1 + p_2 I_2)}{I_1 + I_2} & d_{treat} &= \frac{1}{\tau} \\
 C_c' &= C \\
 P_c' &= P
 \end{aligned}$$

The initial conditions of the variables S , I_1 , I_2 and R are given by the equilibrium solution of the system with $d_{treat} = d_{treat0}$. The initial conditions of C_c and P_c are zero and the initial condition of τ is $(1/d_{treat0})$.

Parameter values

| Symbol | Definition | Value | Source |
|--------------|---|-----------------------|---|
| N | Population size | 10^6 | |
| L | Average life expectancy | 50 years | |
| d_{imm} | Average duration of immunity in the absence of re-exposure | 1 year | [1] |
| d_{in} | Average duration of untreated infection | 0.5 year | Assumed |
| d_{treat0} | Average duration of treated sensitive infection | 2 wks (figure 5) | [2] |
| | | 4 wks (other figures) | Assumed |
| p_1 | Average proportion of infected individuals without preexisting immunity that have clinical malaria | 0.87 | [1] & $p_1 = \frac{d_{treat}(d_{in} - 14.12)}{d_{treat}d_{in} - \eta_0}$ |
| p_2 | Average proportion of infected individuals that have clinical malaria who have experienced infection previously within a year | 0.08 | [1] & $p_1 = \frac{d_{treat}(d_{in} - 2.23)}{d_{treat}d_{in} - \eta_0}$ |
| k | coefficient of speed of spread of resistance | 0 - 0.1 | Assumed |
| η_0 | percentage of individuals with clinical infection that receive treatment | 60 | Assumed |
| R_0 | Basic reproduction number | 2.4 – 63 | corresponds to parasite prevalence of 20% to 98% |

Given presumptive treatment of clinical cases at a coverage of 60%, $k=0.04$ indicates that the percentage of resistant infections would increase from 0% to about 50% in 20 to 40 years depending on transmission intensity. $k=0.08$ indicates that the percentage of resistant infections would increase from 0% to about 50% in 5 to 20 years depending on transmission intensity

Elimination model

This model is produced from two linked copies of the basic model with the presence or absence of the subscript V representing individuals who have received or not received effective vaccine respectively. The additional variables h , m and x represent respectively rates of receiving routine treatment, receiving treatment as part of MSAT and vaccination.

$$\begin{aligned}
S' &= (1 - c_b) \frac{N}{L} - \left(\pi \lambda + \frac{1}{L} \right) S + \frac{1}{d_{imm}} R - x(t) S + y S_V \\
I_1' &= \pi \lambda S - \left(\frac{(\eta p_1 + m)}{d_{treat}} + \frac{(1 - \eta p_1 - m)}{d_{in}} + \frac{1}{L} + \frac{m}{d_{treat}} \right) I_1 - x(t) I_1 + y I_{1V} - m(t) I_1 \\
I_2' &= \pi \lambda R - \left(\frac{(\eta p_2 + m)}{d_{treat}} + \frac{(1 - \eta p_2 - m)}{d_{in}} + \frac{1}{L} + \frac{m}{d_{treat}} \right) I_2 - x(t) I_2 + y I_{2V} - m(t) I_2 \\
R' &= \left(\frac{(\eta p_1 + m)}{d_{treat}} + \frac{(1 - \eta p_1 - m)}{d_{in}} + \frac{m}{d_{treat}} \right) I_1 + \left(\frac{(\eta p_2 + m)}{d_{treat}} + \frac{(1 - \eta p_2 - m)}{d_{in}} + \frac{m}{d_{treat}} \right) I_2 \\
&\quad - \left[\pi \lambda + \frac{1}{d_{imm}} + \frac{1}{L} \right] R - x(t) R + y R_V \\
S_V' &= c_b \frac{N}{L} - \left(\rho \pi \lambda + \frac{1}{L} \right) S_V + \frac{1}{d_{imm}} R_V + x(t) S - y S_V \\
I_{1V}' &= \rho \pi \lambda S_V - \left(\frac{(\eta p_1 + m)}{d_{treat}} + \frac{(1 - \eta p_1 - m)}{d_{in}} + \frac{1}{L} + \frac{m}{d_{treat}} \right) I_{1V} + x(t) I_1 - y I_{1V} \\
I_{2V}' &= \rho \pi \lambda R_V - \left(\frac{(\eta p_2 + m)}{d_{treat}} + \frac{(1 - \eta p_2 - m)}{d_{in}} + \frac{1}{L} + \frac{m}{d_{treat}} \right) I_{2V} + x(t) I_2 - y I_{2V} \\
R_V' &= \left(\frac{(\eta p_1 + m)}{d_{treat}} + \frac{(1 - \eta p_1 - m)}{d_{in}} + \frac{1}{L} + \frac{m}{d_{treat}} \right) I_{1V} + \left(\frac{(\eta p_2 + m)}{d_{treat}} + \frac{(1 - \eta p_2 - m)}{d_{in}} + \frac{1}{L} + \frac{m}{d_{treat}} \right) I_{2V} \\
&\quad - \left[\rho \pi \lambda + \frac{1}{d_{imm}} + \frac{1}{L} \right] R_V + x(t) R - y R_V \\
\lambda &= \frac{R_0 \left(\frac{1}{L} + \frac{1}{d_{treat}} \right) (I_1 + I_2 + I_{1V} + I_{2V})}{N} \\
C &= p_1 (I_1 + I_{1V}) + p_2 (I_2 + I_{2V}) \\
P &= \frac{\eta p_1 (I_1 + I_{1V}) + \eta p_2 (I_2 + I_{2V}) + m (I_1 + I_{1V} + I_2 + I_{2V})}{(I_1 + I_{1V} + I_2 + I_{2V})} \\
C_c' &= C \\
P_c' &= P \\
p_{res} &= \frac{\frac{1}{d_{treat}} - \frac{1}{d_{treat0}}}{\frac{1}{d_{in}} - \frac{1}{d_{treat0}}} \\
\tau' &= -k P p_{res} \left(\tau - \frac{1}{d_{in}} \right) \\
d_{treat} &= \frac{1}{\tau} \\
\eta(t) &= \begin{cases} 0 & \text{for } \text{mod}(t, 1) \geq \frac{9}{12} \\ \eta_0 & \text{for } \text{mod}(t, 1) < \frac{9}{12} \end{cases} \\
x(t) &= \begin{cases} -4 \ln \left(\frac{100 - c}{100} \right) & \text{for } \text{mod}(t, 1) \geq \frac{9}{12} \\ 0 & \text{for } \text{mod}(t, 1) < \frac{9}{12} \end{cases} \\
m(t) &= \begin{cases} 1 - \exp \left(4 \ln(1 - m_{sat}) \text{mod} \left(t - \frac{9}{12} \right) \right) & \text{for } \text{mod}(t, 1) \geq \frac{9}{12} \\ 0 & \text{for } \text{mod}(t, 1) < \frac{9}{12} \end{cases}
\end{aligned}$$

The initial conditions are as for the baseline model for the non-vaccine variables. The initial conditions for the vaccine variables are all zero. To derive the formula for the rate of vaccination $x(t)$, we consider a population being vaccinated at a constant rate q given an initial condition of 100%. Then the differential equation describing the percentage of vaccinated individuals, n , is given by $n' = q(100 - n)$ with solution $n(t) = 100 - 100e^{-qt}$. Thus to achieve a coverage, c , in 3 months (i.e. $\frac{1}{4}$ years) $n\left(\frac{1}{4}\right) = 100 - 100e^{-\frac{q}{4}} = c$, thus $q = -4 \ln\left(\frac{100 - c}{100}\right)$, which should be the function for $x(t)$ during the period of vaccination each year.

Parameter values

| Symbol | Definition | Value | Source |
|-----------|---|--------------------------|--------|
| ρ | proportional vaccine efficacy | 0 to 1 | [3] |
| π | reduction in force of infection due to increased vector control | 0.3 | [4] |
| c | annual vaccine coverage | 80 % | |
| y | rate of loss of vaccine effect | 1- 10 year ⁻¹ | [3] |
| m_{sat} | proportion of infected people treated during MSAT | 0.8 | [2] |

References

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