

Text S1. Mathematical Analysis of the model

Change in T_0 affects only parameters λ and p of the model system

We scale the variables T_l , T_h , and I in the model system as $\hat{T}_l = T_l/T_0$, $\hat{T}_h = T_h/T_0$, and $\hat{I} = I/T_0$. Then the model system reduces to

$$\begin{aligned} \frac{d\hat{T}_l}{dt} &= \frac{\lambda}{T_0} + q\hat{T}_h - d\hat{T}_l - r\hat{T}_l - \beta_l V\hat{T}_l, & \hat{T}_l(0) &= \frac{T_{l0}}{T_0}, \\ \frac{d\hat{T}_h}{dt} &= r\hat{T}_l - d\hat{T}_h - \beta_h V\hat{T}_h - q\hat{T}_h, & \hat{T}_h(0) &= \frac{T_{h0}}{T_0}, \\ \frac{d\hat{I}}{dt} &= \beta_l V\hat{T}_l + \beta_h V\hat{T}_h - \delta\hat{I}, & \hat{I}(0) &= \frac{I_0}{T_0}, \\ \frac{dV}{dt} &= pT_0\hat{I} - cV, & V(0) &= V_0. \end{aligned}$$

Now, applying transformations $\lambda \rightarrow \lambda T_0$ and $p \rightarrow p/T_0$, we obtain the original system. Hence, change in T_0 affects only parameters λ and p of the model system.

Calculation of the basic reproduction number using the next generation matrix

In this section, we derive R_0 , the reproduction number, using the next generation matrix method [1-3]. It is easy to see that the model system has exactly one infection-free equilibrium $E^0 = (T_l^0, T_h^0, I^0, V^0) = \left(\frac{\lambda(d+q)}{d(d+r+q)}, \frac{\lambda r}{d(d+r+q)}, 0, 0 \right)$, and the equations for the infected cell and virus compartments of the linearized system of the model at E^0 take the form:

$$\frac{dI}{dt} = -\delta I + \frac{\lambda}{d(d+r+q)} [\beta_l(d+q) + \beta_h r] V, \quad (S1)$$

$$\frac{dV}{dt} = pI - cV. \quad (S2)$$

We now introduce the matrices:

$$F = \begin{pmatrix} 0 & \frac{\lambda}{d(d+r+q)} [\beta_l(d+q) + \beta_h r] \\ 0 & 0 \end{pmatrix},$$

and

$$V = \begin{pmatrix} \delta & 0 \\ -p & c \end{pmatrix}.$$

These expressions give

$$F\mathcal{V}^{-1} = \begin{pmatrix} \frac{\lambda p}{\delta c d(d+r+q)} [\beta_l(d+q) + \beta_h r] & \frac{\lambda}{c d(d+r+q)} [\beta_l(d+q) + \beta_h r] \\ 0 & 0 \end{pmatrix}.$$

Then R_0 corresponds to the spectral radius of $F\mathcal{V}^{-1}$:

$$R_0 = \rho(F\mathcal{V}^{-1}) = \frac{\lambda p}{\delta c d(d+r+q)} [\beta_l(d+q) + \beta_h r].$$

Proof of “ E^0 is locally asymptotically stable if $R_0 < 1$ and unstable if $R_0 > 1$ ”

Jacobian of the model system linearized around E^0 is:

$$J^0 = \begin{pmatrix} X_{2 \times 2} & Y_{2 \times 2} \\ 0_{2 \times 2} & Z_{2 \times 2} \end{pmatrix},$$

where

$$X = \begin{pmatrix} -d-r & q \\ r & -d-q \end{pmatrix},$$

$$Y = \begin{pmatrix} 0 & \frac{-\beta_l \lambda (d+q)}{d(d+r+q)} \\ 0 & \frac{-\beta_h \lambda r}{d(d+r+q)} \end{pmatrix},$$

$$Z = \begin{pmatrix} -\delta & \frac{\lambda}{d(d+r+q)} [\beta_l(d+q) + \beta_h r] \\ p & -c \end{pmatrix}.$$

Here, eigenvalues of J^0 are given by eigenvalues of the matrices X and Z .

Since $\text{tr}(X) = -(2d+r+q) < 0$ and $\det(X) = d(d+r+q) > 0$, both eigenvalues of X are negative.

Also,

$$\text{tr}(Z) = -(\delta + c) < 0.$$

$$\det(Z) = \delta c [1 - R_0] > 0 \quad \text{iff} \quad R_0 < 1.$$

This implies that eigenvalues of Z , and so eigenvalues of J^0 , are negative if and only if $R_0 < 1$. Hence, E^0 is locally asymptotically stable if $R_0 < 1$ and unstable if $R_0 > 1$.

Proof of “ E^* exists if $R_0 > 1$, and E^* , if it exists, is locally asymptotically stable”

For the analysis of the infected steady state during morphine conditioning, we assume $\beta_l \ll \beta_h$ and $q \ll r$ as given by our parameter estimates to simplify the model. In this case, E^* is given by

$$E^* = (T_l^*, T_h^*, I^*, V^*) = \left(\frac{\lambda}{d+r}, \frac{\delta c}{\beta_h p}, \frac{dc}{\beta_h p} (R_0 - 1), \frac{d}{\beta_h} (R_0 - 1) \right).$$

Clearly, E^* exists if $R_0 > 1$. Jacobian of the model system linearized around E^* is:

$$J^* = \begin{pmatrix} -d-r & 0_{1 \times 3} \\ A_{3 \times 1} & B_{3 \times 3} \end{pmatrix},$$

where

$$A = \begin{pmatrix} r \\ 0 \\ 0 \end{pmatrix}, \quad B = \begin{pmatrix} -dR_0 & 0 & -\delta c/p \\ d(R_0 - 1) & -\delta & \delta c/p \\ 0 & p & -c \end{pmatrix}.$$

One eigenvalue of J^* is $-(d+r) < 0$, and other three eigenvalues are given by eigenvalues of the matrix B . The eigenvalue, ξ , of the matrix B is obtained by solving $\det(B - \xi I_{3 \times 3}) = 0$.

$$\text{i.e., } \begin{vmatrix} -dR_0 - \xi & 0 & -\delta c/p \\ d(R_0 - 1) & -\delta - \xi & \delta c/p \\ 0 & p & -c - \xi \end{vmatrix} = 0.$$

This implies

$$a_3 \xi^3 + a_2 \xi^2 + a_1 \xi + a_0 = 0,$$

where $a_0 = \delta c d (R_0 - 1)$, $a_1 = d R_0 (\delta + c)$, $a_2 = d R_0 + \delta + c$, $a_3 = 1$. If $R_0 > 1$, each $a_i > 0$, $i = 0, 1, 2, 3$. Also, it is easy to show that $a_2 a_1 - a_3 a_0 > 0$. Therefore, by Routh-Hurwitz criterion, all eigenvalues of B , and so all eigenvalues of J^* , are negative. Hence, if $R_0 > 1$, E^* exists and is locally asymptotically stable.

References

1. Diekmann O, Heesterbeek JA, Roberts MG (2009) The construction of next-generation matrices for compartmental epidemic models. *J R Soc Interface* 7: 873–885.
2. van den Driessche P, Watmough J (2002) Reproduction numbers and sub-threshold endemic equilibria for compartmental models of disease transmission. *Math Biosci* 180: 29–48.
3. Heffernan JM, Smith RJ, Wahl LM (2005) Perspectives on the basic reproductive ratio. *J R Soc Interface* 2: 281–293.