

ASYMPTOTIC STABILITY OF SPR_SODE MODEL FOR DENGUE

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ABSTRACT

An ordinary differential equation with stochastic parameters, called SPR_SODE model for the spread of dengue fever was considered to analyze further. It was defined the set of stochastic equations and a reproductive number R_0 . This R_0 was defined for mosquito as well as human parameters. In this paper, the asymptotic stability of the disease-free equilibrium point of the above said model was discussed.

KEYWORDS: SPR_SODE Model, Stochastic, Scaled Variables, Asymptotic Stability, Reproductive Number

INTRODUCTION

Dengue is one of the diseases which is in worldwide and comes under infectious diseases. The work of a carrier (i.e) the medium for transmitting is performed by the mosquito, "Aedes Ageypti [4, Gantmacher.F.R., 1977]. There are so many models for such infectious diseases. We need a separate model for such a special disease like dengue fever for better results. In this work, the SPR_SODE model [2,3, Dhevarajan.S, et.al, 2013] (SPR_Stochastic Ordinary differential model) is considered to analyze further. Asymptotically stable equilibrium points or equilibrium solutions can be defined as the equilibrium solutions in which solutions that start "near" them move toward the equilibrium solution [1, Boyd et.al, 1999].

SPR_SODE MODEL

All the notions of SPR_SODE model [2,3, Dhevarajan.S, et.al, 2013] are taken for further analysis without any change and the same model is given below.

$$\begin{aligned} \frac{d}{dt} \Big[SS_h \Big] &= \wp_h + b_h \Big[\Big[TP \Big]_h \Big] + L_h \Big[\Big[RC \Big]_h \Big] - \tau_h(t) \Big[SS \Big]_h - \Omega_h \Big[\Big[TP \Big]_h \Big] \Big[SS \Big]_h \\ \frac{d}{dt} \Big[EX_h \Big] &= \tau_h(t) \Big[SS \Big]_h - \xi_h \Big[EX \Big]_h - \Omega_h \Big(\Big[TP \Big]_h \Big) \Big[EX \Big]_h \\ \frac{d}{dt} \Big[IF_h \Big] &= \xi_h \Big[EX \Big]_h - \Big[RC \Big]_h \Big[IF \Big]_h - \Omega_h \Big(\Big[TP \Big]_h \Big) \Big[RC \Big]_h \\ \frac{d}{dt} \Big[RC_h \Big] &= \theta_h \Big[IF \Big]_h - L_h \Big[RC \Big]_h - \Omega_h \Big(\Big[TP \Big]_h \Big) \Big[IF \Big]_h - \eta \Big[IF \Big]_h \\ \frac{d}{dt} \Big[SS_m \Big] &= \Big[BIR \Big]_m \Big[TP \Big]_m - \tau_m(t) \Big[SS \Big]_m - \Omega_m \Big(\Big[TP \Big]_m \Big) \Big[SS \Big]_m \\ \frac{d}{dt} \Big[EX_m \Big] &= \tau_m(t) \Big[SS \Big]_m - \xi_m \Big[EX \Big]_m - \Omega_m \Big(\Big[TP \Big]_m \Big) \Big[EX \Big]_m \\ \frac{d}{dt} \Big[IF_m \Big] &= \xi_m \Big[EX \Big]_m - \Omega_m \Big(\Big[TP \Big]_m \Big) \Big[IF \Big]_m \end{aligned}$$

By converting (A) to fractional quantities and denoting each scaled population by small letters, one can get,

$$\begin{aligned} \frac{d}{dt}[ex]_{h} &= \frac{\phi_{h}\phi_{m}P_{mh}[if]_{m}}{\phi_{m}[TP]_{m} + \phi_{h}[TP]_{h}} \cdot [TP]_{m} \cdot [1-[ex]_{h} - [if]_{h} - [re]_{h}] - \left[\xi_{h} + [BIR]_{h} + \frac{\partial}{[TP]_{h}}\right] [ex]_{h} + \eta_{h}[if]_{h}[ex]_{h} \\ \frac{d}{dt}[if]_{h} &= \xi_{h}[ex]_{h} - \left(\theta_{h} + [BIR]_{h} + \frac{\partial}{[TP]_{h}}\right) [if]_{h} + \eta_{h}[if]_{h}^{2} \\ \frac{d}{dt}[rc]_{h} &= \theta_{h}[if]_{h} - \left(L_{h} + [BIR]_{h} \frac{\partial}{[TP]_{h}}\right) [rc]_{h} + \eta_{h}[if]_{h}[TP]_{h} \\ \frac{d}{dt}[TP]_{h} &= \theta_{h}[if]_{h} - \left(L_{h} + [BIR]_{h} \frac{\partial}{[TP]_{h}}\right) [rc]_{h} + \eta_{h}[if]_{h}[TP]_{h} \\ \frac{d}{dt}[TP]_{h} &= \delta_{h} + \theta_{h}[TP]_{h} - ([DID]_{h} + [DDD]_{h}[TP]_{h}) [TP]_{h} - \eta_{h}[if]_{h}[TP]_{h} \\ \frac{d}{dt}[ex]_{m} &= \frac{\phi_{h}\phi_{m}}{\phi_{m}[TP]_{m} + \phi_{h}[TP]_{h}} \cdot [TP]_{h} [P_{mh}[if]_{h} + P_{mh}[rc]_{h}] [1-[ex]_{h} - [if]_{h}] - [\xi_{h} + [BIR]_{m}][ex]_{m} \\ \frac{d}{dt}[if]_{m} &= \xi_{m}[ex]_{m} - [BIR]_{m}[if]_{m} \\ \frac{d}{dt}[TP]_{m} &= [BIR]_{m}[TP]_{m} - ([DID]_{m} + [DDD]_{m}[TP]_{m})[TP]_{m} \end{aligned} \tag{B}$$

where \Re_{hm} and \Re_{mh} can be written in mathematical notation as,

$$\Re_{hm} = \frac{\xi_{m}}{\xi_{m} + [DID]_{m} + [DDD]_{m} [TP]_{m}^{*}} \frac{\phi_{h} \phi_{m} P_{mh} [TP]_{h}^{*}}{\phi_{m} [TP]_{m}^{*} + \phi_{h} [TP]_{h}^{*}} P_{mh} [[DID]_{m} + [DDD]_{m} [TP]_{m}^{*}]^{-1}$$

$$\Re_{mh} = \frac{\xi_{h}}{\xi_{h} + [DID]_{h} + [DDD]_{h} [TP]_{h}^{*}} \frac{\phi_{h} \phi_{m} P_{mh} [TP]_{m}^{*}}{\phi_{h} [TP]_{h}^{*} + \phi_{m} [TP]_{m}^{*}} \left(\theta_{h} + \eta_{h} + [DID]_{h} + [DDD]_{h} [TP]_{h}^{*}\right)^{-1}$$

$$\cdot \left[P_{mh} + \overline{P_{mh}} \cdot \theta_{h} \left(\eta_{h} + [DID]_{m} + [DDD]_{m} [TP]_{h}^{*}\right)^{-1}\right]$$
(D)

ASYMPTOTIC STABILITY

The Jacobian of the dengue model (B) evaluated at X_{nodis} is of the form

$$J = \begin{pmatrix} J_{11} & 0 & 0 & 0 & J_{16} & 0 \\ J_{21} & J_{22} & 0 & 0 & 0 & 0 & 0 \\ 0 & J_{12} & J_{13} & 0 & 0 & 0 & 0 \\ 0 & J_{42} & 0 & J_{44} & 0 & 0 & 0 \\ 0 & J_{52} & 0 & J_{54} & J_{55} & 0 & 0 \\ 0 & 0 & 0 & 0 & J_{65} & J_{66} & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & J_{77} \end{pmatrix}$$
(1)

Now our interest to find out the eigen values of (1). Let the eigen values be λ_i , i=1, 2, 3,...,7. The fourth and seventh columns of the jacobian given by (1) corresponding to the total human and mosquito populations, contain only the diagonal terms. The diagonal terms of the Jacobian (1) provide two of the eigen values say, λ_1 and λ_2 and can be defined by,

$$\lambda_{1} = [BIR]_{h} - [DID]_{h} - 2[DDD]_{h} [TP]_{h}^{*} = -\sqrt{[BIR]_{h} - [DID]_{h}}^{2} + 4[DDD]_{h} \mathcal{O}_{h}$$
(2)

$$\lambda_2 = [BIR]_m - [DID]_m - 2[DDD]_m [TP]_m^* = -[[BIR]_m - [DID]_m]$$
(3)

Let us conclude from the earlier assumption that, both λ_1 and λ_2 are always negative since $[BIR]_m > [DID]_m$. Let us create another matrix by excluding the fourth and seventh rows and columns of the Jacobian (1). By solving the characteristic equation of the matrix formed now, gives the other five eigen values. Now by defining

$$U_{i}'s \text{ as } U_{1} = \xi_{h} + \left[BIR\right]_{h} + \frac{\delta \rho_{h}}{\left[TP\right]_{h}^{*}}, U_{2} = \theta_{h} + \eta_{h} + \left[BIR\right]_{h} + \frac{\delta \rho_{h}}{\left[TP\right]_{h}^{*}}, U_{3} = L_{h} + \eta_{h} + \left[BIR\right]_{h} + \frac{\delta \rho_{h}}{\left[TP\right]_{h}^{*}}, U_{4} = \xi_{m} + \left[BIR\right]_{m}, U_{5} = \left[BIR\right]_{m}, U_{6} = \left[MB\right]_{h}^{*} \cdot P_{mh}, U_{7} = \xi_{h}, U_{8} = \left[MB\right]_{m}^{*} \cdot P_{hm}, U_{9} = \xi_{m}, U_{10} = \theta_{h}, U_{10} = \theta_{$$

 $U_{11} = \left[MB \right]_{m}^{*}$. $\overline{P_{hm}}$, Where, W_{i} 's can be written as,

$$\begin{split} W_{5} &= 1 \; ; \; W_{4} = U_{1} + U_{2} + U_{3} + U_{4} + U_{5} \; ; \; W_{4} = U_{1} + U_{2} + U_{3} + U_{4} + U_{5} \\ W_{3} &= U_{1}U_{2} + U_{1}U_{3} + U_{1}U_{4} + U_{1}U_{5} + U_{2}U_{3} + U_{2}U_{4} + U_{2}U_{5} + U_{3}U_{4} + U_{3}U_{5} + U_{4}U_{5}, \\ W_{2} &= U_{1}U_{2}U_{3} + U_{1}U_{2}U_{4} + U_{1}U_{2}U_{5} + U_{1}U_{3}U_{4} + U_{1}U_{3}U_{5} + U_{1}U_{4}U_{5} + U_{2}U_{3}U_{4} + U_{2}U_{3}U_{5} \\ &\quad + U_{2}U_{4}U_{5} + U_{3}U_{4}U_{5}, \\ W_{1} &= U_{1}U_{2}U_{3}U_{4} + U_{1}U_{2}U_{3}U_{5} + U_{1}U_{2}U_{4}U_{5} + U_{1}U_{3}U_{4}U_{5} + U_{2}U_{3}U_{4}U_{5} - U_{6}U_{7}U_{8}U_{9}, \\ W_{0} &= U_{1}U_{2}U_{3}U_{4}U_{5} - U_{3}U_{6}U_{7}U_{8}U_{9} - U_{6}U_{7}U_{9}U_{10}U_{11} \text{ With} \\ W_{5}\lambda^{5} + W_{4}\lambda^{4} + W_{3}\lambda^{3} + W_{2}\lambda^{2} + W_{1}\lambda + A_{0} = 0 \end{split}$$

$$(4)$$

Now, we have to find out the signs of the solutions of (4). The Liénard – Chipart criterion [6] gives, any of the following four conditions is necessary and sufficient in order that all roots of a polynomial $a_n x^n + a_{n-1} x^{n-1} + ... + a_1 x + a_0 = 0$ with real coefficients have negative real parts:

$$a_n > 0, a_{n-2} > 0, \dots, \Delta_1 > 0, \Delta_3 > 0, \dots; a_n > 0, a_{n-2} > 0, \dots, \Delta_2 > 0, \Delta_4 > 0, \dots;$$

 $a_n > 0, a_{n-1} > 0, a_{n-3} > 0, \dots, \Delta_1 > 0, \Delta_3 > 0, \dots; a_n > 0, a_{n-1} > 0, a_{n-3} > 0, \dots, \Delta_2 > 0, \Delta_1 > 0, \dots, where$

 Δ_i be the principal minors with order i, with i = 1, 2, 3,...,n. For the use the Routh–Hurwitz criterion, First it is to prove that when $R_0 < 1$, all roots of (4) have negative real part. The Routh–Hurwitz criterion [5, section 1.6-6(b)] states that for a real algebraic equation

$$a_{n}x^{n} + a_{n-1}x^{n-1} + \dots + a_{1}x + a_{0} = 0$$
(5)

Given $a_n > 0$, all roots have negative real part if and only if $a_n = V_0$, $a_{n-1} = V_1$,

$$\mathbf{V}_{2} = \begin{vmatrix} a_{n-1} & a_{n} \\ a_{n-3} & a_{n-2} \end{vmatrix}, \quad \mathbf{V}_{3} = \begin{vmatrix} a_{n-1} & a_{n} & 0 \\ a_{n-3} & a_{n-2} & a_{n-1} \\ a_{n-5} & a_{n-4} & a_{n-3} \end{vmatrix}, \quad \dots \\ \mathbf{V}_{n} = \begin{vmatrix} a_{n-1} & \dots & 0 \\ \dots & \dots & \dots \\ 0 & \dots & a_{0} \end{vmatrix}$$
 are all positive, with $a_{i} = 0$ for $i < 0$. This is true if

and only if all a_i and either all even-numbered V_k or all odd-numbered V_k are positive (6, Li'enard–Chipart test). By Korn and Korn [5] in section 1.6-6(c) state Descartes's rule of sign as the number of positive real roots of a real algebraic equation (5) is equal to the number; N_a , of sign changes in the sequence, $a_n, a_{n-1}, ..., a_1, a_0$ of coefficients, where the departure terms are ignored, or it is less than N_a by a positive even integer.

Now, it is to show that when $R_0 < 1$, all the coefficients, W_i , of the characteristic equation (4), and V_0 , V_2 , and V_4 , are positive, hence by the Routh–Hurwitz criterion one can say that, all the eigen values of (1) have negative real part. Now, it is to show that when $R_0 > 1$, there is one and only one sign change in the sequence W_5 , W_4 ... W_0 hence, by Descartes's rule of sign along with positive real part, there is only one eigen value, and also the disease free equilibrium point is unstable. The expression for R_0^2 in (C) can be written, in terms of Ui, as

$$R_{0}^{2} = \frac{U_{3}U_{6}U_{7}U_{8}U_{9} + U_{6}U_{7}U_{9}U_{10}U_{11}}{U_{1}U_{2}U_{3}U_{4}U_{5}}$$
(6)

For
$$R_0 < 1$$
, by (6), $U_3 U_6 U_7 U_8 U_9 + U_6 U_7 U_9 U_{10} U_{11} < U_1 U_2 U_3 U_4 U_5$ (7)

$$U_{6}U_{7}U_{8}U_{9} < U_{1}U_{2}U_{4}U_{5}$$
(8)

As all the $B_{i'}$'s are positive, W_5, W_4, W_3 and W_2 are always positive. From (8) we see that $W_1 > 0$, and from (7) we see that $W_0 > 0$. Thus, for $R_0 < 1$, all Wi are positive. We now show that the even-numbered V_k are positive for $R_0 < 1$. For the fifth-degree polynomial (4), $V_0 = W_5$, which is always positive. $V_2 = W_3 W_4 - W_2 W_5$, which we can show to be a positive sum of products of U_i 's, so $V_2 > 0$. Lastly, $V_4 = W_1[W_2W_3W_4 - (W_1W_4^2 + W_2^2 W_5)] - W_0[W_3(W_3W_4 - W_2W_5) - (2W_1W_4W_5 - W_0W_2^5)].$ For convenience let us use the notations T_1 and T_2 such that $T_1 = [W_2 W_3 W_4 - (W_1 W_4^2 + W_2^2 W_5)]$ and $T_2 = [W_3 (W_3 W_4 - W_2 W_5) - (2W_1 W_4 W_5 - W_0 W_5^5)]$ respectively. Where $T_1 > 0$ and $T_2 > 0$. Hence $V_4 = W_1 T_1 - W_0 T_2$. Let us define $T_2^{[1]} = T_2 + U_6 U_7 U_9 U_{10} U_{11}$. $T_2^{[i]} > T_2$ and $W_0 > 0$ for $V_4^{[1]} = W_1 T_1 - W_0 T_2^{[1]}, V_4 > V_4^{[1]}$. Similarly, As let us define $W_{0}^{[1]} = W_{0} - U_{3}U_{6}U_{7}U_{8}U_{9} - U_{6}U_{7}U_{9}U_{10}U_{11}. \quad \text{As } W_{4} > W_{4}^{[1]} \quad \text{and} \quad T_{2}^{[1]} > 0, \quad \text{for} \quad V_{4}^{[2]} = W_{1}T_{1} - W_{0}^{[1]}T_{2}^{[1]},$ $V_{4}^{[1]} > V_{4}^{[2]}$ at last, let us define $W_{1}^{[1]} = W_{1} - U_{1}U_{2}U_{3}U_{5} + U_{6}U_{7}U_{8}U_{9}$, As $W_{1}^{[1]} < W_{1}$ (for $R_{0} < 1$) and $T_{1} > 0$ For $V_{a}^{[3]} = W_{a}^{[1]}T_{1} - W_{a}^{[1]}T_{2}^{[1]}, \quad V_{a}^{[2]} > V_{a}^{[3]}.$ It can be shown that $V_{a}^{[3]}$ Is a sum of positive terms, so $V_{A}^{[3]} > 0$ $V_{A}^{[1]} > V_{A}^{[2]} > V_{A}^{[3]}$, $V_{4} > 0$. Thus, for $R_{0} < 1$, all roots of (4) have negative real parts. When $R_{0} > 1$,

55

sequence, W_5 , W_4 , W_3 , W_2 , W_1 has exactly one sign change. Thus, by Descartes's rule of sign, (4) has one positive real root when $R_0 > 1$.

CONCLUSIONS

A stochastic ordinary differential equation called SPR_SODE model for the spread of dengue fever is analyzed. For our model, the disease-free equilibrium point, X_{nodis} , is locally asymptotically stable if $R_0 < 1$ and unstable if $R_0 > 1$. If $R_0 < 1$, on average each infected individual infects less than one other individual, and the disease dies out. If $R_0 > 1$, on average each infected individual, infects more than one other individual, so one can expect the disease to spread. The Jacobian of (B) at X_{nodis} has one eigen value equal to 0 at $R_0 = 1$.

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