

Global dynamics of humoral immunity Chikungunya virus with two routes of infection and Holling type-II



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Abstract

In this work, we analyze the global dynamics of within-host Chikungunya virus (CHIKV) infection model with humoral immune response. We incorporate two modes of infections, attaching a CHIKV to a host monocyte, and contacting an infected monocyte with an uninfected monocyte. The infection incident rate is given by Holling type-II. The basic reproduction number \mathcal{R}_0 is used to prove that the CHIKV-free equilibrium E_0 is globally asymptotically stable when $\mathcal{R}_0 \leq 1$ and the infected equilibrium E_1 is globally asymptotically stable when $\mathcal{R}_0 > 1$. Numerical simulations have been performed to confirm the theoretical results.

Keywords: Chikungunya virus, holling type-II, global stability, Lyapunov function, viral and cellular infections.

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1. Introduction

Mosquito is one of the dangerous insect throughout the world. It can carry and spread viruses to humans and animals causes many of deaths every year. A great efforts has been paid to develop and analyze mathematical models that describe the population dynamics of mosquito-borne diseases such as Zika [2, 4, 7], dengue [1, 27, 43, 48], malaria [3, 5, 6, 36], yellow fever [40] and chikungunya [8–10, 34, 37–39, 46]. Chikungunya virus (CHIKV) is transmitted to humans by infected *Aedes albopictus* and *Aedes aegypti* mosquito. CHIKV causes severe joint and muscle pain, fever, rash, headache, nausea and fatigue. Wang and Liu [45] have proposed and studied a within-host CHIKV dynamics model which contains four compartments, uninfected-monocytes (s), infected monocytes (y), free CHIKV particles (p) and antibodies (x). The model has been extended in [13, 14] by considering general CHIKV-monocyte incidence rate. In [13, 14, 45] it has been assumed that the uninfected monocyte becomes infected by contacting with CHIKV (CHIKV-to-monocyte transmission). Long and Heise [35] have reported that the CHIKV can also spread by infected-to-monocyte transmission. Mathematical models of different viruses with both cellular and

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viral infections have been studied in several works [24, 25, 31–33, 41, 44, 47]. In a very recent work, Elaiw et al. [15] have studied the dynamics of CHIKV model with two routes of infection, however, they did not consider the holling-II. The aim of the present paper is to propose and analyze a CHIKV dynamics model where the infection rate is given by Holling type-II incidence. The proposed model is given as:

$$\dot{s}(t) = \beta - \delta s(t) - \frac{\eta_1 s(t)p(t)}{1 + \omega s(t)} - \frac{\eta_2 s(t)y(t)}{1 + \omega s(t)}, \tag{1.1}$$

$$\dot{y}(t) = \frac{\eta_1 s(t)p(t)}{1 + \omega s(t)} + \frac{\eta_2 s(t)y(t)}{1 + \omega s(t)} - \epsilon y(t), \tag{1.2}$$

$$\dot{p}(t) = \pi y(t) - cp(t) - rx(t)p(t), \tag{1.3}$$

$$\dot{x}(t) = \lambda + \rho x(t)p(t) - mx(t). \tag{1.4}$$

The uninfected monocytes are generated monocytes by rate β , die with rate $\delta s(t)$ and be infected by CHIKV and infected monocytes with rate $\frac{\eta_1 s(t)p(t)}{1 + \omega s(t)} + \frac{\eta_2 s(t)y(t)}{1 + \omega s(t)}$, where ω is the uninfected monocyte Holling type-II constant, and η_1 , and η_2 are the incidence rate constants. Constants ϵ , c , and m represent, respectively, the death rate constants of the infected monocytes, CHIKV, and antibodies. Constant π is the production rate constant of the CHIKV from infected monocytes. Antibodies attack the CHIKV at rate $rx(t)p(t)$. Once antigen is encountered, the antibodies expand at a constant rate λ and proliferate at rate $\rho x(t)p(t)$. All the parameters of the model are positive.

1.1. Basic properties

The following lemma establishes the nonnegativity and boundedness of the solutions of system (1.1)-(1.4).

Lemma 1.1. *There exist $M_1, M_2, M_3 > 0$, such that the following compact set is positively invariant for system (1.1)-(1.4);*

$$\Gamma = \{(s, y, p, x) \in \mathbb{R}_{\geq 0}^4 : 0 \leq s, y \leq M_1, 0 \leq p \leq M_2, 0 \leq x \leq M_3\}.$$

Proof. We have

$$\begin{aligned} \dot{s} |_{s=0} &= \beta > 0, \\ \dot{y} |_{y=0} &= \frac{\eta_1 sp}{1 + \omega s} \geq 0 \text{ for all } s, p \geq 0, \\ \dot{p} |_{p=0} &= \pi y \geq 0 \text{ for all } y \geq 0, \\ \dot{x} |_{x=0} &= \lambda > 0. \end{aligned}$$

This shows that $(s(t), y(t), p(t), x(t)) \in \mathbb{R}_{\geq 0}^4$ with $(s(0), y(0), p(0), x(0)) \in \mathbb{R}_{\geq 0}^4$. Let us define

$$H_1(t) = s(t) + y(t), \qquad H_2(t) = p(t) + \frac{r}{\rho}x(t).$$

Then from Eqs. (1.1)-(1.4) we get

$$\dot{H}_1(t) = \beta - \delta s(t) - \epsilon y(t) \leq \beta - \sigma_1(s(t) + y(t)) = \beta - \sigma_1 H_1(t),$$

where, $\sigma_1 = \min\{\delta, \epsilon\}$. Hence $H_1(t) \leq M_1$, if $H_1(0) \leq M_1$, where $M_1 = \frac{\beta}{\sigma_1}$. It follows that $0 \leq s(t), y(t) \leq M_1$ if $0 \leq s(0) + y(0) \leq M_1$. Moreover, we have

$$\dot{H}_2(t) = \pi y(t) - cp(t) + \frac{r}{\rho}\lambda - \frac{mr}{\rho}x(t) \leq \pi M_1 + \frac{r}{\rho}\lambda - \sigma_2 \left(p(t) + \frac{r}{\rho}x(t) \right) = \pi M_1 + \frac{r}{\rho}\lambda - \sigma_2 H_2(t),$$

where, $\sigma_2 = \min\{c, m\}$. Hence $H_2(t) \leq M_2$, if $H_2(0) \leq M_2$, where $M_2 = \frac{\pi M_1 + \frac{r}{\rho}\lambda}{\sigma_2}$. Since $p(t)$ and $x(t)$ are all non-negative, then $0 \leq p(t) \leq M_2$ and $0 < x(t) \leq M_3$ if $0 < p(0) + \frac{r}{\rho}x(0) \leq M_2$, where $M_3 = \frac{\rho M_2}{r}$. \square

1.2. Equilibria

We define the basic reproduction number as:

$$\mathcal{R}_0 = \frac{(\eta_1\pi m + \eta_2cm + \eta_2r\lambda)\beta}{\epsilon(cm + r\lambda)(\delta + \beta\omega)}$$

Lemma 1.2. Consider system (1.1)-(1.4), then

- if $\mathcal{R}_0 \leq 1$, then there exists only one equilibrium $E_0 \in \Gamma$, and
- if $\mathcal{R}_0 > 1$, then there exist two equilibria $E_0 \in \Gamma$ and $E_1 \in \overset{\circ}{\Gamma}$, where $\overset{\circ}{\Gamma}$ is the interior of Γ .

Proof. Let $E(s, y, p, x)$ be any equilibrium satisfying

$$0 = \beta - \delta s - \frac{\eta_1 sp}{1 + \omega s} - \frac{\eta_2 sy}{1 + \omega s}, \tag{1.5}$$

$$0 = \frac{\eta_1 sp}{1 + \omega s} + \frac{\eta_2 sy}{1 + \omega s} - \epsilon y, \tag{1.6}$$

$$0 = \pi y - cp - rxp, \tag{1.7}$$

$$0 = \lambda + \rho xp - mx. \tag{1.8}$$

By solving Eqs. (1.5)-(1.8) we get two equilibria a CHIKV-free equilibrium $E_0 = (s_0, 0, 0, x_0)$, where $s_0 = \frac{\beta}{\delta}$ and $x_0 = \frac{\lambda}{m}$. Moreover, we have

$$\frac{C_1 p^3 + C_2 p^2 + C_3 p + C_4}{\bar{C}_1 p + \bar{C}_2} = 0,$$

where

$$\begin{aligned} C_1 &= c\epsilon\rho^2(-\pi\eta_1 - c\eta_2 + c\epsilon\omega), \\ C_2 &= C_{21} + C_{22} + C_{23} + C_{24} + C_{25}, \\ C_3 &= C_{31} + C_{32} + C_{33} + C_{34} + C_{35} + C_{36}, \\ C_4 &= m\pi(\pi\pi\beta\eta_1 - cm(\delta\epsilon - \beta\eta_2 + \beta\epsilon\omega) - r\lambda(\delta\epsilon - \beta\eta_2 + \beta\epsilon\omega)), \\ \bar{C}_1 &= \bar{C}_{11} + \bar{C}_{12}, \\ \bar{C}_2 &= \bar{C}_{21} + \bar{C}_{22}, \end{aligned}$$

and

$$\begin{aligned} C_{21} &= \rho\pi\eta_1(r\epsilon\lambda + \pi\beta\rho), & C_{22} &= 2\rho c^2 m\epsilon(\eta_2 - \epsilon\omega), & C_{23} &= 2\rho c m\pi\epsilon\eta_1, \\ C_{24} &= 2\rho c r\epsilon\lambda(\eta_2 - \epsilon\omega), & C_{25} &= -\pi\rho^2 c(\delta\epsilon - \beta\eta_2 + \beta\epsilon\omega), & C_{31} &= -m\pi\eta_1(r\epsilon\lambda + 2\pi\beta\rho), \\ C_{32} &= c^2 m^2 \epsilon(-\eta_2 + \epsilon\omega), & C_{33} &= r^2 \lambda^2 \epsilon(-\eta_2 + \epsilon\omega), & C_{34} &= r\lambda\pi\rho(\delta\epsilon - \beta\eta_2 + \beta\epsilon\omega), \\ C_{35} &= cm(-m\pi\epsilon\eta_1 + 2r\epsilon\lambda(-\eta_2 + \epsilon\omega)), & C_{36} &= 2cm\pi\rho(\delta\epsilon - \beta\eta_2 + \beta\epsilon\omega), & \bar{C}_{11} &= \pi r r\lambda(\eta_2 - \epsilon\omega), \\ \bar{C}_{12} &= 2\pi\rho m(\pi\eta_1 + c(\eta_2 - \epsilon\omega)), & \bar{C}_{21} &= -m\pi(\pi\eta_1 + cm(\eta_2 - \epsilon\omega)), & \bar{C}_{22} &= -m\pi r\lambda(\eta_2 - \epsilon\omega). \end{aligned}$$

Let define a function $X(p)$ as:

$$X(p) = \frac{C_1 p^3 + C_2 p^2 + C_3 p + C_4}{\bar{C}_1 p + \bar{C}_2} = 0,$$

we obtain

$$X(0) = \frac{\beta(cme + r\epsilon\lambda)(\delta + \beta\omega)(\mathcal{R}_0 - 1)}{\epsilon(cm + r\lambda)(\delta + \beta\omega)(\mathcal{R}_0 - 1) + \epsilon\delta(cm + r\lambda)}, \quad \lim_{p \rightarrow (\frac{m}{\rho})^-} X(p) = -\frac{m r \epsilon \lambda}{\rho^2} < 0.$$

Therefore, if $\mathcal{R}_0 > 1$ then $X(0) > 0$ and there exists $p_1 \in (0, \frac{m}{\rho})$ such that $X(p_1) = 0$. It follows from Eqs. (1.6)-(1.8) that

$$x_1 = \frac{\lambda}{m - \rho p_1} > 0, \quad y_1 = \frac{p_1(c + rx_1)}{\pi} > 0,$$

$$s_1 = \frac{-\delta - p_1\eta_1 - y_1\eta_2 + \beta\omega + \sqrt{4\beta\delta\omega + (-\delta - p_1\eta_1 - y_1\eta_2 + \beta\omega)^2}}{2\delta\omega} > 0.$$

Therefore, if $\mathcal{R}_0 > 1$, then the system has an infected equilibrium $E_1 = (s_1, y_1, p_1, x_1)$. Now we show that $E_0 \in \Gamma$ and $E_1 \in \overset{\circ}{\Gamma}$. Clearly, $E_0 \in \Gamma$. From the equilibria conditions of E_1 we have

$$\beta = \delta s_1 + \frac{\eta_1 s_1 p_1}{1 + \omega s_1} + \frac{\eta_2 s_1 y_1}{1 + \omega s_1} \Rightarrow \delta s_1 + \epsilon y_1 = \beta \Rightarrow 0 < s_1 < \frac{\beta}{\delta} \leq M_1, 0 < y_1 < \frac{\beta}{\epsilon} \leq M_1.$$

Moreover, from Eqs. (1.7) and (1.8) we have

$$cp_1 = \pi y_1 + \frac{r}{\rho}\lambda - \frac{mr}{\rho}x_1 \Rightarrow cp_1 + \frac{mr}{\rho}x_1 = \pi y_1 + \frac{r}{\rho}\lambda < \pi M_1 + \frac{r}{\rho}\lambda,$$

$$p_1 < \frac{\pi M_1 + \frac{r}{\rho}\lambda}{c} \leq M_2, \quad x_1 < \frac{\rho}{r} \frac{\pi M_1 + \frac{r}{\rho}\lambda}{m} \leq \frac{\rho M_2}{r} = M_3.$$

It follows that, $E_1 \in \overset{\circ}{\Gamma}$. □

2. Global properties

To investigate the global stability of the equilibria we construct Lyapunov functions using the method presented [30] and followed by [11, 12, 15–24, 26, 28, 29, 42]. Define $F(v) = v - 1 - \ln v$.

Theorem 2.1. *For system (1.1)-(1.4), if $\mathcal{R}_0 \leq 1$, then E_0 is globally asymptotically stable in Γ .*

Proof. Let $\mathcal{R}_0 \leq 1$ and construct a Lyapunov function $U_0(s, y, p, x)$ as:

$$U_0(s, y, p, x) = s - s_0 - \int_{s_0}^s \frac{s_0(1 + \omega\theta)}{\theta(1 + \omega s_0)} d\theta + y + \frac{\eta_1 s_0}{(c + rx_0)(1 + \omega s_0)} p + \frac{r\eta_1 s_0}{\rho(c + rx_0)(1 + \omega s_0)} x_0 F\left(\frac{x}{x_0}\right).$$

Clearly, $U_0(s, y, p, x) > 0$ for all $s, y, p, x > 0$ and $U_0(s_0, 0, 0, x_0) = 0$. Calculating $\frac{dU_0}{dt}$ along system (1.1)-(1.4) we obtain

$$\begin{aligned} \frac{dU_0}{dt} &= \left(1 - \frac{s_0(1 + \omega s)}{s(1 + \omega s_0)}\right) \left(\beta - \delta s - \frac{\eta_1 sp}{1 + \omega s} - \frac{\eta_2 sy}{1 + \omega s}\right) + \frac{\eta_1 sp}{1 + \omega s} + \frac{\eta_2 sy}{1 + \omega s} - \epsilon y \\ &\quad + \frac{\eta_1 s_0}{(c + rx_0)(1 + \omega s_0)} \left(\pi y - cp - rxp\right) + \frac{r\eta_1 s_0}{\rho(c + rx_0)(1 + \omega s_0)} \left(1 - \frac{x_0}{x}\right) \left(\lambda + \rho xp - mx\right) \\ &= \left(1 - \frac{s_0(1 + \omega s)}{s(1 + \omega s_0)}\right) \left(\beta - \delta s\right) + \frac{\eta_2 s_0 y}{1 + \omega s_0} - \epsilon y + \frac{\eta_1 s_0}{(c + rx_0)(1 + \omega s_0)} \pi y \\ &\quad + \frac{r\eta_1 s_0}{\rho(c + rx_0)(1 + \omega s_0)} \left(1 - \frac{x_0}{x}\right) \left(\lambda - mx\right). \end{aligned}$$

Substituting $\beta = \delta s_0$ and $\lambda = mx_0$ we get

$$\begin{aligned} \frac{dU_0}{dt} &= -\delta \frac{(s - s_0)^2}{s(1 + \omega s_0)} + \epsilon \left(\frac{\eta_2 s_0}{\epsilon(1 + \omega s_0)} + \frac{\eta_1 s_0 \pi}{\epsilon(c + rx_0)(1 + \omega s_0)} - 1\right) y - \frac{r\eta_1 s_0 m}{\rho(c + rx_0)(1 + \omega s_0)} \frac{(x - x_0)^2}{x} \\ &= -\delta \frac{(s - s_0)^2}{s(1 + \omega s_0)} - \frac{r\eta_1 s_0 m}{\rho(c + rx_0)(1 + \omega s_0)} \frac{(x - x_0)^2}{x} + \epsilon(\mathcal{R}_0 - 1)y. \end{aligned}$$

If $\mathcal{R}_0 \leq 1$, then $\frac{dU_0}{dt} \leq 0$ for all $s, y, p, x > 0$ and $\frac{dU_0}{dt} = 0$ when $s = s_0, x = x_0$ and $y = 0$. It can be easily shown that $\frac{dU_0}{dt} = 0$ at E_0 . Applying LaSalle’s invariance principle, we get E_0 is globally asymptotically stable when $\mathcal{R}_0 \leq 1$. □

Theorem 2.2. For system (1.1)-(1.4), if $\mathcal{R}_0 > 1$, then E_1 is globally asymptotically stable in $\overset{\circ}{\Gamma}$.

Proof. Let a function $U_1(s, y, p, x)$ be defined as:

$$U_1(s, y, p, x) = s - s_1 - \int_{s_1}^s \frac{s_1(1 + \omega\theta)}{\theta(1 + \omega s_1)} d\theta + y_1 F\left(\frac{y}{y_1}\right) + \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} p_1 F\left(\frac{p}{p_1}\right) + \frac{\eta_1 s_1 p_1}{\rho \pi y_1(1 + \omega s_1)} x_1 F\left(\frac{x}{x_1}\right).$$

Clearly, $U_1(s, y, p, x) > 0$ for all $s, y, p, x > 0$ and $U_1(s_1, y_1, p_1, x_1) = 0$. Calculating $\frac{dU_1}{dt}$ along the trajectories of (1.1)-(1.4) we obtain

$$\begin{aligned} \frac{dU_1}{dt} &= \left(1 - \frac{s_1(1 + \omega s)}{s(1 + \omega s_1)}\right) \left(\beta - \delta s - \frac{\eta_1 s p}{1 + \omega s} - \frac{\eta_2 s y}{1 + \omega s}\right) + \left(1 - \frac{y_1}{y}\right) \left(\frac{\eta_1 s p}{1 + \omega s} + \frac{\eta_2 s y}{1 + \omega s} - \epsilon y\right) \\ &\quad + \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} \left(1 - \frac{p_1}{p}\right) (\pi y - c p - r x p) + \frac{\eta_1 s_1 p_1}{\rho \pi y_1(1 + \omega s_1)} \left(1 - \frac{x_1}{x}\right) (\lambda + \rho x p - m x) \\ &= \left(1 - \frac{s_1(1 + \omega s)}{s(1 + \omega s_1)}\right) (\beta - \delta s) + \frac{\eta_1 s_1 p}{1 + \omega s_1} + \frac{\eta_2 s_1 y}{1 + \omega s_1} - \frac{\eta_1 s p}{1 + \omega s} \frac{y_1}{y} - \frac{\eta_2 s y}{1 + \omega s} \frac{y_1}{y} - \epsilon y + \epsilon y_1 \\ &\quad + \frac{\eta_1 s_1 p_1}{1 + \omega s_1} \frac{y}{y_1} - \frac{\eta_1 s_1 p_1}{1 + \omega s_1} \frac{p_1 y}{p y_1} - \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} c p + \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} c p_1 + \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} r x p_1 \\ &\quad - \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} x_1 p + \frac{\eta_1 s_1 p_1}{\rho \pi y_1(1 + \omega s_1)} \left(1 - \frac{x_1}{x}\right) (\lambda - m x). \end{aligned}$$

Applying the equilibrium conditions for E_1

$$\beta = \delta s_1 + \frac{\eta_1 s_1 p_1}{1 + \omega s_1} + \frac{\eta_2 s_1 y_1}{1 + \omega s_1}, \quad \epsilon y_1 = \frac{\eta_1 s_1 p_1}{1 + \omega s_1} + \frac{\eta_2 s_1 y_1}{1 + \omega s_1}, \quad c p_1 = \pi y_1 - r x_1 p_1, \quad \lambda = m x_1 - \rho x_1 p_1.$$

we get

$$\begin{aligned} \frac{dU_1}{dt} &= -\delta \frac{(s - s_1)^2}{s(1 + \omega s_1)} + \left(1 - \frac{s_1(1 + \omega s)}{s(1 + \omega s_1)}\right) \left(\frac{\eta_1 s_1 p_1}{1 + \omega s_1} + \frac{\eta_2 s_1 y_1}{1 + \omega s_1}\right) \\ &\quad - \frac{\eta_1 s_1 p_1}{1 + \omega s_1} \frac{s p y_1(1 + \omega s_1)}{s_1 p_1 y(1 + \omega s)} - \frac{\eta_2 s_1 y_1}{1 + \omega s_1} \frac{s(1 + \omega s_1)}{s_1(1 + \omega s)} + \frac{\eta_1 s_1 p_1}{1 + \omega s_1} + \frac{\eta_2 s_1 y_1}{1 + \omega s_1} \\ &\quad - \frac{\eta_1 s_1 p_1}{1 + \omega s_1} \frac{p_1 y}{p y_1} + \frac{\eta_1 s_1 p_1}{1 + \omega s_1} - 2 \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} r x_1 p_1 + \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} r x p_1 \\ &\quad + \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} r x_1 p_1 \frac{x_1}{x} - \frac{\eta_1 s_1 p_1 m}{\rho \pi y_1(1 + \omega s_1)} \frac{(x - x_1)^2}{x}. \end{aligned} \tag{2.1}$$

Eq. (2.1) can be simplified as:

$$\begin{aligned} \frac{dU_1}{dt} &= -\delta \frac{(s - s_1)^2}{s(1 + \omega s_1)} + \frac{\eta_1 s_1 p_1}{1 + \omega s_1} \left[3 - \frac{s_1(1 + \omega s)}{s(1 + \omega s_1)} - \frac{s p y_1(1 + \omega s_1)}{s_1 p_1 y(1 + \omega s)} - \frac{p_1 y}{p y_1}\right] \\ &\quad + \frac{\eta_2 s_1 y_1}{1 + \omega s_1} \left[2 - \frac{s_1(1 + \omega s)}{s(1 + \omega s_1)} - \frac{s(1 + \omega s_1)}{s_1(1 + \omega s)}\right] - \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} r x_1 p_1 \left[2 - \frac{x}{x_1} - \frac{x_1}{x}\right] \\ &\quad - \frac{\eta_1 s_1 p_1 m}{\rho \pi y_1(1 + \omega s_1)} \frac{(x - x_1)^2}{x} \\ &= -\delta \frac{(s - s_1)^2}{s(1 + \omega s_1)} - \frac{\eta_1 s_1 p_1}{\pi y_1(1 + \omega s_1)} \frac{r \lambda}{\rho x_1} \frac{(x - x_1)^2}{x} + \frac{\eta_1 s_1 p_1}{1 + \omega s_1} \left[3 - \frac{s_1(1 + \omega s)}{s(1 + \omega s_1)} - \frac{s p y_1(1 + \omega s_1)}{s_1 p_1 y(1 + \omega s)} - \frac{p_1 y}{p y_1}\right] \\ &\quad + \frac{\eta_2 s_1 y_1}{1 + \omega s_1} \left[2 - \frac{s_1(1 + \omega s)}{s(1 + \omega s_1)} - \frac{s(1 + \omega s_1)}{s_1(1 + \omega s)}\right]. \end{aligned}$$

Using the rule

$$\frac{1}{n} \sum_{i=1}^n a_i \geq \sqrt[n]{\prod_{i=1}^n a_i}, \quad \text{where, } a_i \geq 0, i = 1, 2, \dots, n,$$

we get

$$\frac{1}{3} \left(\frac{s_1(1 + \omega s)}{s(1 + \omega s_1)} + \frac{s p y_1(1 + \omega s_1)}{s_1 p_1 y(1 + \omega s)} + \frac{p_1 y}{p y_1}\right) \geq 1, \quad \frac{1}{2} \left(\frac{s_1(1 + \omega s)}{s(1 + \omega s_1)} + \frac{s(1 + \omega s_1)}{s_1(1 + \omega s)}\right) \geq 1.$$

Therefore, $\frac{dU_1}{dt} \leq 0$ for all $s, y, p, x > 0$ and $\frac{dU_1}{dt} = 0$ if and only if $s = s_1, y = y_1, p = p_1$ and $x = x_1$. It follows that the global stability of E_1 is induced from LaSalle’s invariance principle. \square

3. Numerical simulations

Using the values in Table 1, we consider two cases as follows:

Case 1: We simulate system (1.1)-(1.4) with the following initial conditions:

IC1: $s(0) = 14.0, y(0) = 1.0, p(0) = 1.5,$ and $x(0) = 1.5;$

IC2: $s(0) = 8.0, y(0) = 2.0, p(0) = 3.0,$ and $x(0) = 4.0;$

IC3: $s(0) = 4.0, y(0) = 3.5, p(0) = 6.0,$ and $x(0) = 7.0.$

We fix $\omega = 0.09$ and consider the values of η_1 and η_2 as following sets:

Set (I): We let $\eta_1 = \eta_2 = 0.001$. Computing $\mathcal{R}_0 = 0.0857 < 1$, Figures (1)-(4) show that, $E_0 = (s_0, 0, 0, x_0)$ is globally asymptotically stable, where $s_0 = \frac{\beta}{\delta} = 20$ and $x_0 = \frac{\lambda}{m} = 1.4$, which agrees with the result of (2.1).

Set (II): We choose $\eta_1 = \eta_2 = 0.05$. Calculating $\mathcal{R}_0 = 4.2857 > 1$, we compute the equilibria as $E_0(20.0, 0, 0, 1.4)$ and $E_1 = (6.66, 2.66, 3.73, 5.51)$. We have observed that in Figures (1)-(4), when $\mathcal{R}_0 > 1$, the solution of the system tend to E_1 for IC1-IC3 and (2.2) is confirmed.

Case 2. We fixed the value $\eta_1 = \eta_2 = 0.06$, by using the following initial conditions $s(0) = 7, y(0) = 2.0, p(0) = 3.0,$ and $x(0) = 4.0$, we can see from Figures (5)-(8) that the evolution of the system’s states with different values of ω . We have observed that $\mathcal{R}_0 > 1$, and the trajectory of the system converges to the equilibrium E_1 for smaller values of ω e.g. $\omega = 0.0, 0.2, 0.4$. Whereas, $\mathcal{R}_0 \leq 1$, and the system has one equilibrium E_0 when ω become larger e.g. $\omega = 2, 5$. Let ω^{ct} be the critical value of the parameter ω , such that

$$\mathcal{R}_0 = \frac{(\eta_1\pi m + \eta_2cm + \eta_2r\lambda)\beta}{\epsilon(cm + r\lambda)(\delta + \beta\omega^{ct})} = 1.$$

Using the data given in Table 1, we obtain $\omega^{ct} = 0.67$. The variation of \mathcal{R}_0 w.r.t. ω are listed in Table 2. We can observed that as ω is increased then \mathcal{R}_0 is decreased. Moreover, we have the following cases:

- (i) if $0 \leq \omega < 0.67$, then E_1 exists and it is globally asymptotically stable,
- (ii) if $\omega \geq 0.67$, then E_0 is globally asymptotically stable.

Table 1: The value of the parameters of model (1.1)-(1.4).

Parameter	Value	Parameter	Value
β	2	δ	0.1
η_1	varied	η_2	varied
π	4	c	0.1
r	0.5	λ	1.4
m	1	ρ	0.2
ω	varied	ϵ	0.5

Table 2: The value of \mathcal{R}_0 for different values of ω .

ω	Equilibria	\mathcal{R}_0
0.0	(3.77, 3.24, 3.91, 6.43)	14.3997
0.2	(8.93, 2.21, 3.54, 4.80)	2.8800
0.4	(15.93, 0.08, 2.25, 2.60)	1.6000
0.67	(20.00, 0.00, 0.00, 1.40)	1.00
1	(20.00, 0.00, 0.00, 1.40)	0.6857
5	(20.00, 0.00, 0.00, 1.40)	0.1426

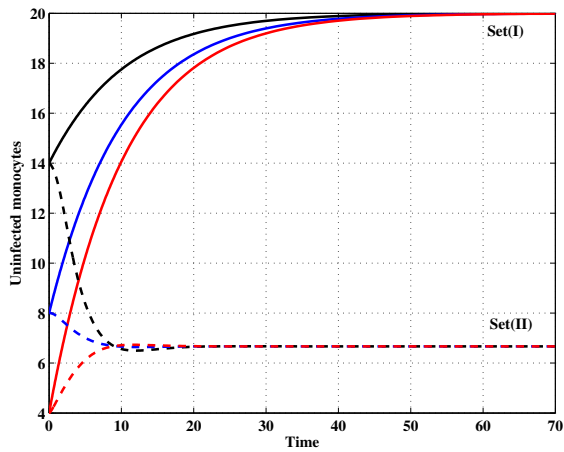


Figure 1: Uninfected monocytes.

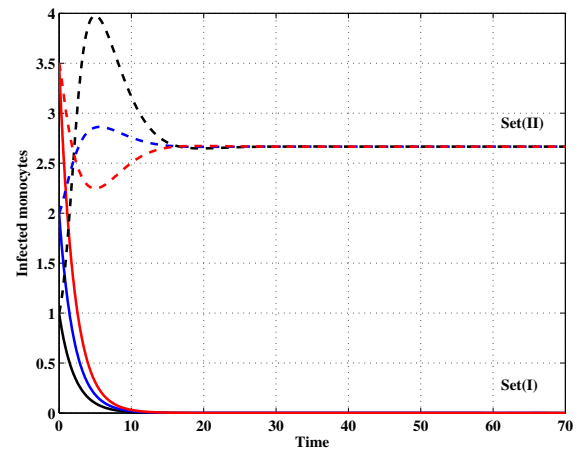


Figure 2: Infected monocytes.

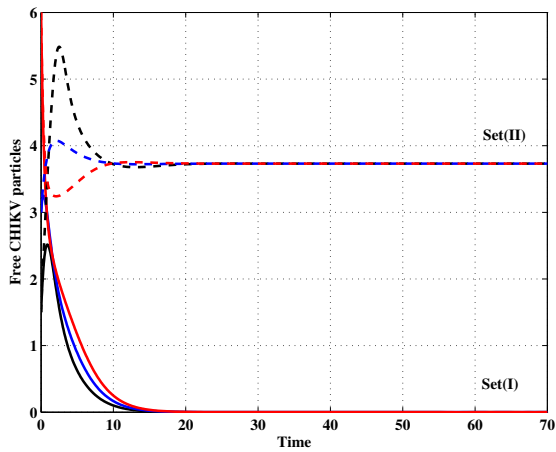


Figure 3: Free CHIKV particles.

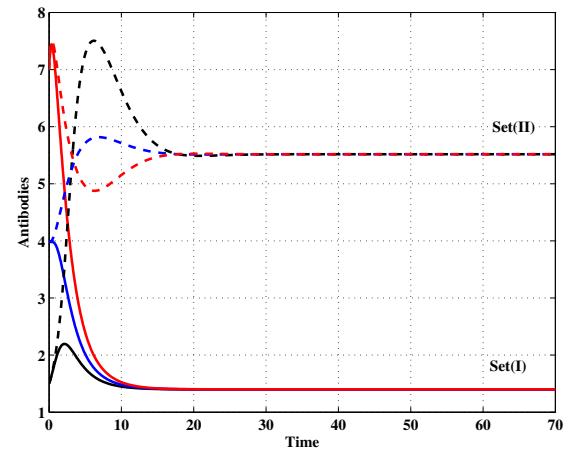


Figure 4: Antibodies.

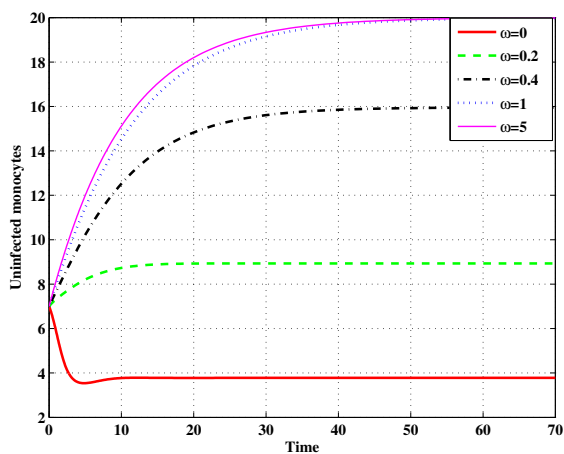


Figure 5: Uninfected monocytes.

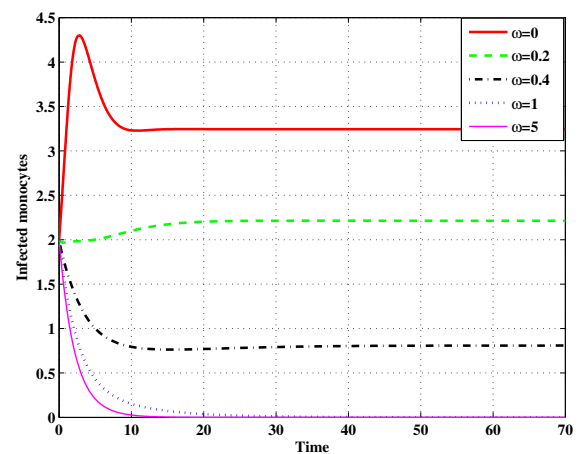


Figure 6: Infected monocytes.

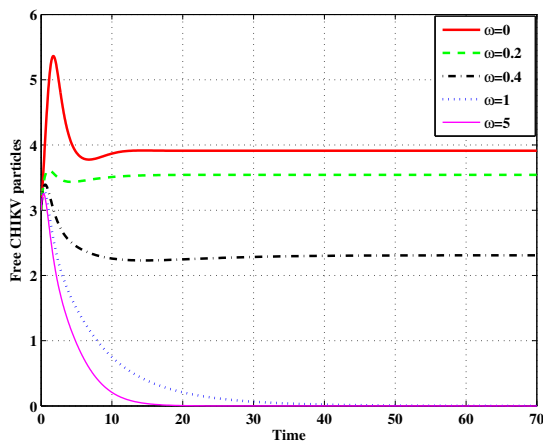


Figure 7: Free CHIKV particles.

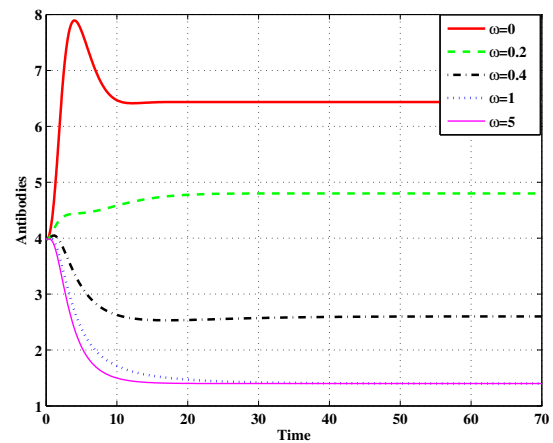


Figure 8: Antibodies.

References

- [1] A. Abdelrazec, J. Belair, C. H. Shan, H. P. Zhu, *Modeling the spread and control of dengue with limited public health resources*, *Math. Biosci.*, **271** (2016), 136–145. 1
- [2] F. B. Augusto, S. Bewick, W. F. Fagan, *Mathematical model of Zika virus with vertical transmission*, *Infectious Disease Modelling*, **2** (2017), 244–267. 1
- [3] E. Beretta, V. Capasso, D. G. Garao, *A mathematical model for malaria transmission with asymptomatic carriers and two age groups in the human population*, *Math. Biosci.*, **300** (2018), 87–101. 1
- [4] E. Bonyah, K. O. Okosun, *Mathematical modeling of Zika virus*, *Asian Pacific J. Tropical Disease*, **6** (2016), 673–679. 1
- [5] N. Chitnis, J. M. Cushing, J. M. Hyman, *Bifurcation analysis of a mathematical model for malaria transmission*, *SIAM J. Appl. Math.*, **67** (2006), 24–45. 1
- [6] N. Chitnis, J. M. Hyman, J. M. Cushing, *Determining important parameters in the spread of malaria through the sensitivity analysis of a mathematical model*, *Bull. Math. Biol.*, **70** (2008), 1272–1296. 1
- [7] E. Dantas, M. Tosin, A. Cunha, *Calibration of a SEIR-SEI epidemic model to describe the Zika virus outbreak in Brazil*, *Appl. Math. Comput.*, **338** (2018), 249–259. 1
- [8] Y. Dumont, F. Chiroleu, *Vector control for the chikungunya disease*, *Math. Biosci. Eng.*, **7** (2010), 313–345. 1
- [9] Y. Dumont, F. Chiroleu, C. Domerg, *On a temporal model for the chikungunya disease: modeling, theory and numerics*, *Math. Biosci.*, **213** (2008), 80–91.
- [10] Y. Dumont, J. M. Tchenche, *Mathematical studies on the sterile insect technique for the chikungunya disease and aedes albopictus*, *J. Math. Biol.*, **65** (2012), 809–854. 1
- [11] A. M. Elaiw, *Global properties of a class of HIV models*, *Nonlinear Anal. Real World Appl.*, **11** (2010), 2253–2263. 2
- [12] A. M. Elaiw, *Global properties of a class of virus infection models with multitarget cells*, *Nonlinear Dynam.*, **69** (2012), 423–435. 2
- [13] A. M. Elaiw, T. O. Alade, S. M. Alsulami, *Analysis of latent CHIKV dynamics models with general incidence rate and time delays*, *J. Biol. Dyn.*, **12** (2018), 700–730. 1
- [14] A. M. Elaiw, T. O. Alade, S. M. Alsulami, *Analysis of within-host CHIKV dynamics models with general incidence rate*, *Int. J. Biomath.*, **11** (2018), 25 pages. 1
- [15] A. M. Elaiw, S. E. Almalki, A. D. Hobiny, *Stability of CHIKV infection models with CHIKV-monocyte and infected-monocyte saturated incidences*, *AIP Advances*, **9** (2019), 12 pages. 1, 2
- [16] A. M. Elaiw, N. A. Almualllem, *Global properties of delayed-HIV dynamics models with differential drug efficacy in cocirculating target cells*, *Appl. Math. Comput.*, **265** (2015), 1067–1089.
- [17] A. M. Elaiw, N. A. Almualllem, *Global dynamics of delay-distributed HIV infection models with differential drug efficacy in cocirculating target cells*, *Math. Methods Appl. Sci.*, **39** (2016), 4–31.
- [18] A. M. Elaiw, N. H. AlShamrani, *Global stability of humoral immunity virus dynamics models with nonlinear infection rate and removal*, *Nonlinear Anal. Real World Appl.*, **26** (2015), 161–190.
- [19] A. M. Elaiw, N. H. AlShamrani, *Stability of a general delay-distributed virus dynamics model with multi-staged infected progression and immune response*, *Math. Methods Appl. Sci.*, **40** (2017), 699–719.
- [20] A. M. Elaiw, N. H. AlShamrani, *Stability of an adaptive immunity pathogen dynamics model with latency and multiple delays*, *Math. Methods Appl. Sci.*, **41** (2018), 6645–6672.
- [21] A. M. Elaiw, S. A. Azoz, *Global properties of a class of HIV infection models with Beddington-DeAngelis functional response*, *Math. Methods Appl. Sci.*, **36** (2013), 383–394.

- [22] A. M. Elaiw, E. K. Elnahary, A. A. Raezah, *Effect of cellular reservoirs and delays on the global dynamics of HIV*, Adv. Difference Equ., **2018** (2018), 36 pages.
- [23] A. M. Elaiw, I. Hassanien, S. A. Azoz, *Global stability of HIV infection models with intracellular delays*, J. Korean Math. Soc., **49** (2012), 779–794.
- [24] A. M. Elaiw, A. A. Raezah, *Stability of general virus dynamics models with both cellular and viral infections and delays*, Math. Methods Appl. Sci., **40** (2017), 5863–5880. 1, 2
- [25] A. M. Elaiw, A. A. Raezah, B. S. Alofi, *Dynamics of delayed pathogen infection models with pathogenic and cellular infections and immune impairment*, AIP Advances, **8** (2018), 14 pages. 1
- [26] A. M. Elaiw, A. A. Raezah, S. A. Azoz, *Stability of delayed HIV dynamics models with two latent reservoirs and immune impairment*, Adv. Difference Equ., **2018** (2018), 25 pages. 2
- [27] L. Esteva, C. Vargas, *A model for dengue disease with variable human population*, J. Math. Biol., **38** (1999), 220–240. 1
- [28] A. D. Hobiny, A. M. Elaiw, A. A. Almatrafi, *Stability of delayed pathogen dynamics models with latency and two routes of infection*, Adv. Difference Equ., **2018** (2018), 26 pages. 2
- [29] G. Huang, Y. Takeuchi, W. Ma, *Lyapunov functionals for delay differential equations model of viral infections*, SIAM J. Appl. Math., **70** (2010), 2693–2708. 2
- [30] A. Korobeinikov, *Global properties of basic virus dynamics models*, Bull. Math. Biol., **66** (2004), 879–883. 2
- [31] X. L. Lai, X. F. Zou, *Modelling HIV-1 virus dynamics with both virus-to-cell infection and cell-to-cell transmission*, SIAM J. Appl. Math., **74** (2014), 898–917. 1
- [32] X. L. Lai, X. F. Zou, *Modeling cell-to-cell spread of HIV-1 with logistic target cell growth*, J. Math. Anal. Appl., **426** (2015), 563–584.
- [33] F. Li, J. L. Wang, *Analysis of an HIV infection model with logistic target cell growth and cell-to-cell transmission*, Chaos Solitons Fractals, **81** (2015), 136–145. 1
- [34] X. Z. Liu, P. Stechliniski, *Application of control strategies to a seasonal model of chikungunya disease*, Appl. Math. Model., **39** (2015), 3194–3220. 1
- [35] K. M. Long, M. T. Heise, *Protective and pathogenic responses to chikungunya virus infection*, Curr. Trop. Med. Rep., **2** (2015), 13–21. 1
- [36] S. Mandal, R. R. Sarkar, S. Sinha, *Mathematical models of malaria-a review*, Malaria J., **10** (2011), 19 pages. 1
- [37] C. A. Manore, K. S. Hickmann, S. Xu, H. J. Wearing, J. M. Hyman, *Comparing dengue and chikungunya emergence and endemic transmission in A. aegypti and A. albopictus*, J. Theoret. Biol., **356** (2014), 174–191. 1
- [38] D. Moulay, M. Aziz-Alaoui, M. Cadivel, *The chikungunya disease: modeling, vector and transmission global dynamics*, Math. Biosci., **229** (2011), 50–63.
- [39] D. Moulay, M. Aziz-Alaoui, H.-D. Kwon, *Optimal control of chikungunya disease: larvae reduction, treatment and prevention*, Math. Biosci. Eng., **9** (2012), 369–392. 1
- [40] S. M. Raimundo, M. Amaku, E. Massad, *Equilibrium analysis of a yellow fever dynamical model with vaccination*, Comput. Math. Methods Med., **2015** (2015), 12 pages. 1
- [41] H. Y. Shu, Y. M. Chen, L. Wang, *Impacts of the cell-free and cell-to-cell infection modes on viral dynamics*, J. Dynam. Differential Equations, **30** (2018), 1817–1836. 1
- [42] H. Y. Shu, L. Wang, J. Watmough, *Global stability of a nonlinear viral infection model with infinitely distributed intracellular delays and CTL immune responses*, SIAM J. Appl. Math., **73** (2013), 1280–1302. 2
- [43] J. J. Tewaa, J. L. Dimi, S. Bowong, *Lyapunov functions for a dengue disease transmission model*, Chaos Solitons Fractals, **39** (2009), 936–941. 1
- [44] J. L. Wang, J. Y. Lang, X. F. Zou, *Analysis of an age structured HIV infection model with virus-to-cell infection and cell-to-cell transmission*, Nonlinear Anal. Real World Appl., **34** (2017), 75–96. 1
- [45] Y. Wang, X. N. Liu, *Stability and Hopf bifurcation of a within-host chikungunya virus infection model with two delays*, Math. Comput. Simulation, **138** (2017), 31–48. 1
- [46] L. Yakob, A. C. Clements, *A mathematical model of chikungunya dynamics and control: the major epidemic on Reunion Island*, PLoS One, **8** (2013), 6 pages. 1
- [47] Y. Yang, L. Zou, S. G. Ruan, *Global dynamics of a delayed within-host viral infection model with both virus-to-cell and cell-to-cell transmissions*, Math. Biosci., **270** (2015), 183–191. 1
- [48] M. Zhu, Y. Xu, *A time-periodic dengue fever model in a heterogeneous environment*, Math. Comput. Simulation, **155** (2019), 115–129. 1