

http://www.aimspress.com/journal/MBE

MBE, 17(5): 4384–4405. DOI: 10.3934/mbe.2020242 Received: 14 April 2020 Accepted: 08 June 2020 Published: 22 June 2020

#### Research article

# Stability and Hopf bifurcation in a virus model with self-proliferation and delayed activation of immune cells

## Huan Kong, Guohong Zhang\*and Kaifa Wang\*

School of Mathematics and Statistics, Southwest University, Chongqing 400715, China

\* Correspondence: Email: zgh711@swu.edu.cn, kfwang72@163.com.

**Abstract:** A new mathematical model was proposed to study the effect of self-proliferation and delayed activation of immune cells in the process of virus infection. The global stability of the boundary equilibria was obtained by constructing appropriate Lyapunov functional. For positive equilibrium, the conditions of stability and Hopf bifurcation were obtained by taking the delay as the bifurcation parameter. Furthermore, the direction and stability of the Hopf bifurcation are derived by using the theory of normal form and center manifold. These results indicate that self-proliferation intensity can significantly affect the kinetics of viral infection, and the delayed activation of immune cells can induce periodic oscillation scenario. Along with the increase of delay time, numerical simulations give the corresponding bifurcation diagrams under different self-proliferation rates, and verify that there exists stability switch phenomenon under some conditions.

**Keywords:** viral infection; self-proliferation; delayed activation of immune cells ; global stability; Hopf bifurcation; stability switch

#### 1. Introduction

Recently, different kinds of viral infection such as HIV (human immunodeficiency virus), HCV (hepatitis C virus) and HBV (hepatitis B virus) have received many attention. Different mathematical models have been formulated to describe the dynamics of virus population in vivo [1–6]. The basic viral infection model of within-host can be described by the following three dimensional system [7]:

$$\begin{pmatrix}
\frac{dx}{dt} = s - \beta xv - d_1 x, \\
\frac{dy}{dt} = \beta xv - d_2 y, \\
\frac{dv}{dt} = ky - uv.
\end{cases}$$
(1.1)

Here x(t), y(t), and v(t) represent uninfected target cells, infected cells and free virus, respectively. Uninfected cells are produced at a constant rate *s*, die at rate  $d_1x$ , and become infected at rate  $\beta xv$ . Infected cells are produced at rate  $\beta xv$  and die at rate *by*. Free virus are produced from infected cells at rate *ky* and die at rate *uv*. It was showed that, for system (1.1), there exist a critical threshold named as the basic reproduction number of virus  $R_0 = \frac{\beta sk}{d_1d_2}$  to determine its global dynamical behavior [8]. During viral infections, the adaptive immune response is mediated by lymphocytes expressing

During viral infections, the adaptive immune response is mediated by lymphocytes expressing antigen specific receptors, T and B lymphocytes, namely, humoral and cellular immunity. To study the population dynamics of immune response, Nowak et al. [9] introduced the CTL population into system (1.1) and obtain the following four dimensional system:

$$\begin{cases} \frac{dx}{dt} = s - \beta xv - d_1 x, \\ \frac{dy}{dt} = \beta xv - d_2 y - pyz, \\ \frac{dv}{dt} = ky - uv, \\ \frac{dz}{dt} = cyz - d_3 z. \end{cases}$$
(1.2)

Here z(t) denotes CTLs, which is produced at rate cyz because of the stimulation of infected cells, and die at rate  $d_{3}z$ . The infected cells are eliminated by CTLs at rate pyz. Following [9], many authors present and develop mathematical models for the cell-mediated immune response [10–13] and the humoral immunity [14–17].

Recently study indicates that the self-proliferation of immune cells can not be neglected besides the stimulation of infected cells. In order to mimic the spontaneous proliferation of CTLs, a logistic proliferation term for CTLs was incorporated in virus infection models in [18], where a rigorous mathematical analysis of the effect of self-proliferation of CTLs on the dynamics of viral infection is necessary. In order to understand the effect of self-proliferation and delayed activation of immune cells in a virus model, we propose the following virus infection model:

$$\begin{cases} \frac{dx}{dt} = s - \beta xv - d_1 x, \\ \frac{dy}{dt} = \beta xv - d_2 y - pyz, \\ \frac{dv}{dt} = ky - uv, \\ \frac{dz}{dt} = cy(t - \tau)z(t - \tau) + rz(1 - \frac{z}{m}) - d_3 z. \end{cases}$$
(1.3)

Here the logistic proliferation term rz(1 - z/m) describes the self-proliferation of CTLs, in which parameter *r* denotes a per capita self-proliferation rate, and *m* means the capacity of CTLs population. Time delay term  $cy(t - \tau)z(t - \tau)$  represents a sequence of events such as antigenic activation and selection. It has been shown that time delays cannot be ignored in models for viral immune response including intracellular delay during viral infection [19–22] and time delay of CTLs stimulating proliferation [23, 24].

Note that the turnover of virus is much faster than that of infection cells within-host [25, 26]. A plausible quasi steady-state assumption is proposed to mimic the fast time-scale [27, 28]. In other

words, v can be replaced by  $\frac{ky}{u}$  in (1.3). Let  $\hat{\beta} = \frac{\beta k}{u}$  and also note as  $\beta$  to simplify the parameter, system (1.3) can be written as:

$$\begin{cases} \frac{dx}{dt} = s - \beta xy - d_1 x, \\ \frac{dy}{dt} = \beta xy - d_2 y - pyz, \\ \frac{dz}{dt} = cy(t - \tau)z(t - \tau) + rz(1 - \frac{z}{m}) - d_3 z. \end{cases}$$
(1.4)

This paper is organized as follows. In section 2, some preliminary results are obtained, including non-negativity and boundedness of the solution of system (1.4), the existence of equilibria under different self-proliferation intensities of CTLs. Section 3 investigate the local and global stability of the boundary equilibria. Section 4 study the effects of delayed activation of immune cells on the existence of Hopf bifurcation. The direction and stability of Hopf bifurcation is investigated in section 5. Some numerical simulations are given to quantify the impact of self-proliferation and delayed activation of CTLs in section 6. Finally, some conclusions and discusses are presented in section 7.

#### 2. Preliminary results

In this section, we first discuss the non-negativity and boundedness of solutions of system (1.4). For  $\tau > 0$ , let  $C = C([-\tau, 0], \mathbb{R}^3_+)$  denote the Banach space of continuous function mapping the interval  $[-\tau, 0]$  into  $\mathbb{R}^3_+$  with the topology of uniform convergence. The initial conditions are given by

$$x(\xi) = \phi_1(\xi), y(\xi) = \phi_2(\xi), z(\xi) = \phi_3(\xi)$$
(2.1)

with  $\phi_i(\xi) \ge 0, \xi \in [-\tau, 0]$  and  $\phi_i(0) > 0$  (i = 1, 2, 3).

**Theorem 2.1.** *The solutions of system* (1.4) *satisfying the initial conditions* (2.1) *are non-negative and ultimately bounded.* 

*Proof.* We first define the right-hand side function of system (1.4) as

$$G(t, K(t)) = \begin{pmatrix} s - \beta xy - d_1 x \\ \beta xy - d_2 y - pyz \\ cy(t - \tau)z(t - \tau) + rz(1 - \frac{z}{m}) - d_3 z \end{pmatrix},$$

where  $K(t) = (K_1(t), K_2(t), K_3(t))^T$  and  $K_1(t) = x, K_2(t) = y, K_3(t) = z$ . It is obvious that the function G(t, K(t)) is locally Lipschitz and by the standard theory of functional differential equation, we know that there exists a unique solution for a given initial conditions. To prove the non-negativity of solutions of system (1.4), we first prove the non-negativity over the time interval  $[0, \tau]$ . Considering the right-hand side functions of system (1.4) over the time interval  $[0, \tau]$ , we have

$$G(t, K(t)) = \begin{pmatrix} G_1(t, K(t)) \\ G_2(t, K(t)) \\ G_3(t, K(t)) \end{pmatrix}_{K_i(t)=0} = \begin{pmatrix} s \\ 0 \\ cy(t-\tau)z(t-\tau) \end{pmatrix} = \begin{pmatrix} s \\ 0 \\ c\phi_2(\xi)\phi_3(\xi) \end{pmatrix},$$

Mathematical Biosciences and Engineering

of system (1.4) remain nonnegative in  $[0, \tau]$ . Similarly, we can repeat the process over  $[\tau, 2\tau]$  and so on by using the method of steps [29], it can be proved that for any finite interval [0, t], the solutions of system (1.4) remains non-negative.

Now we consider the boundeness of the solutions. Define a new variable  $X(t) = x(t)+y(t)+\frac{p}{c}z(t+\tau)$ , and let  $d = \min\{1, d_1, d_2\}$ . By the non-negativity of the solutions of system (1.4), we have

$$\frac{dX}{dt} = s - d_1 x - d_2 y + \frac{pr}{c} z(t+\tau) \left(1 - \frac{z(t+\tau)}{m}\right) - d_3 z(t+\tau) 
= s - d_1 x - d_2 y - \frac{p}{c} z(t+\tau) + \frac{p}{c} z(t+\tau) \left(1 + r - \frac{rz(t+\tau)}{m}\right) - d_3 z(t+\tau) 
\leq s - d_1 x - d_2 y - \frac{p}{c} z(t+\tau) + \frac{pm(1+r)^2}{4rc} 
\leq s + \frac{pm(1+r)^2}{4rc} - dX(t).$$
(2.2)

Taking  $M = s + \frac{pm(1+r)^2}{4rc}$ , we know  $\limsup_{t\to\infty} X(t) \le \frac{M}{d}$ . So the solutions of system(1.4) are ultimately bounded.

The equilibria of (1.4) are the solutions of the following algebraic equations:

$$\begin{cases} s - \beta xy - d_1 x = 0, \\ \beta xy - d_2 y - pyz = 0, \\ cyz + rz(1 - \frac{z}{m}) - d_3 z = 0. \end{cases}$$
(2.3)

It is easy to see that system (1.4) always has infection-free equilibrium  $E_0^{yz} = (x_0, 0, 0)$ , where  $x_0 = \frac{s}{d_1}$ . According to the definition in [30], we obtain the basic reproduction number of virus  $R_0 = \frac{\beta s}{d_1 d_2}$ . Now we define  $x_1 = \frac{d_2}{\beta}$ ,  $y_1 = \frac{d_1(R_0 - 1)}{\beta}$ ,  $x_2 = \frac{s}{d_1}$ ,  $z_2 = \frac{m(r - d_3)}{r}$ . Based on (2.3), after some simple calculations, it is easy to get the following results.

**Proposition 2.2.** (i) Suppose that  $R_0 > 1$ . The immunity-inactivated infection equilibrium  $E_0^z = (x_1, y_1, 0)$  always exists. Especially, besides  $E_0^z$ , an infection-free but immunity-activated equilibrium  $E_0^y = (x_2, 0, z_2)$  will appear if  $r > d_3$ . (ii) Suppose that  $0 \le r \le d_3$ . If  $R_0 > 1 + \frac{\beta(d_3 - r)}{cd_1}$ , system (1.4) has a unique immunity-activated infection equilibrium  $E_1^* = (x_1^*, y_1^*, z_1^*)$ , where

$$x_1^* = \frac{d_2 + pz_1^*}{\beta}, \qquad y_1^* = \frac{rz_1^* + m(d_3 - r)}{mc}$$

and  $z_1^*$  is the unique positive root of the quadratic equation  $A_1z^2 + B_1z + C_1 = 0$ , in which

$$A_1 = -pr\beta, B_1 = -(d_2r\beta + pmcd_1 + \beta pm(d_3 - r)), C_1 = mcd_1d_2(R_0 - 1) - \beta md_2(d_3 - r))$$

Mathematical Biosciences and Engineering

(iii) Suppose that  $r > d_3$ . If  $R_0 > 1 + \frac{pm(r-d_3)}{rd_2}$ , system (1.4) has a unique immunity-activated infection equilibrium  $E_2^* = (x_2^*, y_2^*, z_2^*)$ , where

$$x_2^* = \frac{s}{d_1 + \beta y_2^*}, \qquad z_2^* = \frac{m(cy_2^* + r - d_3)}{r},$$

and  $y_2^*$  is the unique positive root of the quadratic equation  $A_2y^2 + B_2y + C_2 = 0$ , in which

 $A_2 = pmc\beta, B_2 = pm\beta(r - d_3) + d_2r\beta + pmcd_1, C_2 = -rd_1d_2(R_0 - 1) + pmd_1(r - d_3).$ 

#### 3. Global stability analysis of boundary equilibria

In order to study the stability of equilibrium, we linearize system (1.4) about one equilibrium  $E^* = (x^*, y^*, z^*)$  and obtain the following linear system

$$\begin{pmatrix}
\frac{dx}{dt} = (-\beta y^* - d_1)x - \beta x^* y, \\
\frac{dy}{dt} = \beta y^* x + (\beta x^* - d_2 - pz^*)y - py^* z, \\
\frac{dz}{dt} = cz^* y(t - \tau) + cy^* z(t - \tau) + (r - d_3 - \frac{2rz^*}{m})z.
\end{cases}$$
(3.1)

When r = 0, it follows from the study of Michael Y. Li and Hongying Shu [23] that time delay  $\tau$  does not change the stability of the boundary equilibria.

When r > 0, by using the linear system (3.1) and constructing Lyapunov function, we can obtain the following results.

**Proposition 3.1.** Suppose that  $0 < r < d_3$ . Then we have

(i) The infection-free equilibrium  $E_0^{yz}$  is globally asymptotically stable if  $R_0 < 1$ , and it is unstable when  $R_0 > 1$ .

(ii) The immunity-inactivated infection equilibrium  $E_0^z$  is globally asymptotically stable if  $1 < R_0 < 1 + \frac{\beta(d_3 - r)}{cd_1}$ , and it is unstable when  $R_0 > 1 + \frac{\beta(d_3 - r)}{cd_1}$ .

*Proof.* (i) By the linear system (3.1), we can obtain the characteristic equation of system(1.4) at  $E_0^{yz}$  as

$$(\lambda + d_1)(\lambda - (r - d_3))(\lambda + d_1d_2(1 - R_0)) = 0.$$
(3.2)

So the eigenvalues are

$$\lambda_1 = -d_1 < 0, \lambda_2 = r - d_3 < 0, \lambda_3 = d_1 d_2 (R_0 - 1).$$

As a result, if  $R_0 < 1$ , the infection-free equilibrium  $E_0^{yz}$  is locally asymptotically stable, and  $E_0^{yz}$  is unstable if  $R_0 > 1$ .

In order to obtain the global stability of equilibrium  $E_0^{yz}$ , we consider a Lyapunov function given by

$$L_{1} = x - x_{0} - x_{0} \ln \frac{x}{x_{0}} + y + \frac{p}{c}z + p \int_{-\tau}^{0} y(t+\theta)z(t+\theta)d\theta$$

Mathematical Biosciences and Engineering

Taking the time derivative of  $L_1$  along the solution of system (1.4), we have

$$\begin{aligned} \frac{dL_1}{dt} &= (1 - \frac{x_0}{x})(s - \beta xy - d_1 x) + \beta xy - d_2 y - pyz \\ &+ \frac{p}{c}[cy(t - \tau)z(t - \tau) + rz(1 - \frac{z}{m}) - d_3 z] + pyz - py(t - \tau)z(t - \tau) \\ &= -\frac{d_1(x - x_0)^2}{x} + (\beta x_0 - d_2)y + \frac{p}{c}(r - d_3)z - \frac{prz^2}{cm} \\ &= -\frac{d_1(x - x_0)^2}{x} + d_2(R_0 - 1)y + \frac{p}{c}(r - d_3)z - \frac{prz^2}{cm}. \end{aligned}$$

Notice that  $r < d_3$  and  $R_0 < 1$ . We have  $L'_1 \le 0$  for all x(t), y(t), z(t) > 0, and  $L'_1 = 0$  only if  $x = x_0$ , y = 0 and z = 0. It can be verified that the maximal compact invariant set in  $L'_1 = 0$  is the singleton  $E_0^{yz}$ . By the LaSalle's invariance principle, we know the infection-free equilibrium  $E_0^{yz}$  is globally asymptotically stable.

(ii) The characteristic equation of system(1.4) at  $E_0^z$  is

$$(\lambda^2 + d_1 R_0 \lambda + \beta^2 x_1 y_1) (\lambda - (r - d_3 + c y_1 e^{-\lambda \tau})) = 0.$$
(3.3)

When  $\tau = 0$  and  $R_0 < 1 + \frac{\beta(d_3 - r)}{cd_1}$ , the roots of (3.3) are negative. When  $\tau > 0$ , the local stability of equilibrium  $E_0^z$  is solely determined by

$$\lambda - (r - d_3 + cy_1 e^{-\lambda \tau}) = 0.$$
(3.4)

Let  $\lambda = i\omega(\tau)(\omega(\tau) > 0)$  be the root of Eq (3.4). Separating real and imaginary parts yields

$$\begin{cases} d_3 - r = cy_1 \cos(\omega\tau), \\ \omega = cy_1 \sin(\omega\tau). \end{cases}$$
(3.5)

Squaring and adding Eq (3.5) gives

$$\omega^2 - (cy_1)^2 + (d_3 - r)^2 = 0.$$

Note that

$$(cy_1)^2 - (d_3 - r)^2 = \left(\frac{cd_1(R_0 - 1)}{\beta(d_3 - r)}\right)^2 - 1 < 0$$

if  $R_0 < 1 + \frac{\beta(d_3 - r)}{cd_1}$ . Then we know that (3.3) has no root that can across the imaginary axis, which indicate that the immunity-inactivated infection equilibrium  $E_0^z$  is locally asymptotically stable when  $R_0 < 1 + \frac{\beta(d_3 - r)}{cd_1}$ .

In order to obtain the global stability of  $E_0^z$ , we construct the following Lyapunov functions:

$$L_{2} = x - x_{1} - x_{1} \ln \frac{x}{x_{1}} + y - y_{1} - y_{1} \ln \frac{y}{y_{1}} + \frac{p}{c}z + p \int_{-\tau}^{0} y(t+\theta)z(t+\theta)d\theta.$$

Mathematical Biosciences and Engineering

Taking the time derivative of  $L_2$  along the solution of system(1.4), we get

$$\frac{dL_2}{dt} = (1 - \frac{x_1}{x})(s - \beta xy - d_1x) + (1 - \frac{y_1}{y})(\beta xy - d_2y - pyz) + \frac{p}{c}(cy(t - \tau)z(t - \tau) + rz(1 - \frac{z}{m}) - d_3z) + pyz - py(t - \tau)z(t - \tau) = s - d_1x - s\frac{x_1}{x} + \beta x_1y + d_1x_1 - d_2y - \beta xy_1 + d_2y_1 + py_1z + \frac{p}{c}(r - d_3)z - \frac{prz^2}{cm}.$$

Using  $s = \beta x_1 y_1 + d_1 x_1$  and  $d_2 = \beta x_1$ , we have

$$\frac{dL_2}{dt} = (d_1x_1 + \beta x_1y_1)(2 - \frac{x}{x_1} - \frac{x_1}{x}) + p(y_1 - \frac{d_3 - r}{c})z - \frac{prz^2}{cm}$$
$$= (d_1x_1 + \beta x_1y_1)(2 - \frac{x}{x_1} - \frac{x_1}{x}) + \frac{pd_1}{\beta}(R_0 - 1 - \frac{\beta(d_3 - r)}{cd_1})z - \frac{prz^2}{cm}.$$

Using the LaSalle's invariance principle, we can obtain that  $E_0^z$  is globally asymptotically stable.  $\Box$ 

## **Proposition 3.2.** *Suppose that* $r = d_3$ *. Then we have*

(i) The infection-free equilibrium  $E_0^{yz}$  is globally asymptotically stable if  $R_0 < 1$ , and it is unstable when  $R_0 > 1$ .

(ii) The immunity-inactivated infection equilibrium  $E_0^z$  is unstable as long as it appears, i.e.,  $R_0 > 1$ .

*Proof.* (i) The characteristic equation of system (1.4) at  $E_0^{yz}$  is

$$\lambda(\lambda + d_1)(\lambda + d_1d_2(1 - R_0)) = 0.$$
(3.6)

So the eigenvalues are

$$\lambda_1 = 0, \lambda_2 = -d_1 < 0, \lambda_3 = d_1 d_2 (R_0 - 1).$$

So if  $R_0 < 1$ , the infection-free equilibrium  $E_0^{yz}$  is locally asymptotically stable and if  $R_0 > 1$ , the equilibrium  $E_0^{yz}$  is unstable.

In order to obtain the global stability of  $E_0^{yz}$ , we construct the following Lyapunov functions:

$$L_{3} = x - x_{0} - x_{0} \ln \frac{x}{x_{0}} + y + \frac{p}{c}z + p \int_{-\tau}^{0} y(t+\theta)z(t+\theta)d\theta.$$

Taking the time derivative of  $L_3$  along the solution of system (1.4), we have

$$\frac{dL_3}{dt} = (1 - \frac{x_0}{x})(s - \beta xy - d_1 x) + \beta xy - d_2 y - pyz + \frac{p}{c}(cy(t - \tau)z(t - \tau) - \frac{rz^2}{m}) + pyz - py(t - \tau)z(t - \tau) = -\frac{d_1(x - x_0)^2}{x} + (\beta x_0 - d_2)y - \frac{prz^2}{cm} = -\frac{d_1(x - x_0)^2}{x} + d_2(R_0 - 1)y - \frac{prz^2}{cm}.$$

Using the LaSalle's invariance principle, we can obtain that  $E_0^{yz}$  is globally asymptotically stable.

(ii)The characteristic equation of system (1.4) at  $E_0^z$  is

$$(\lambda^2 + d_1 R_0 \lambda + \beta^2 x_1 y_1)(\lambda - c y_1 e^{-\lambda \tau}) = 0.$$
(3.7)

It can be showed that there exist positive real root for the characteristic Eq (4.3), which indicates that the infection-free equilibrium  $E_0^z$  is unstable. This completes the proof.

# **Proposition 3.3.** *Suppose that* $r > d_3$ *. Then we have*

- (i) The infection-free equilibrium  $E_0^{yz}$  is unstable.
- (ii) The immunity-inactivated infection equilibrium  $E_0^z$  is unstable.

(iii) If  $R_0 < 1 + \frac{pm(r-d_3)}{rd_2}$ , the infection-free equilibrium  $E_0^y$  is locally asymptotically stable, moreover if  $R_0 < 1$ , the infection-free equilibrium  $E_0^y$  is globally asymptotically stable; if  $R_0 > 1 + \frac{pm(r-d_3)}{rd_2}$ ,  $E_0^y$  is unstable.

*Proof.* (i) The characteristic equation of system (1.4) at  $E_0^{yz}$  is

$$(\lambda + d_1)(\lambda - (r - d_3))(\lambda + d_1d_2(1 - R_0)) = 0.$$
(3.8)

So the eigenvalues are

$$\lambda_1 = -d_1, \lambda_2 = r - d_3, \lambda_3 = d_1 d_2 (R_0 - 1).$$

It follows from  $\lambda_2 = r - d_3 > 0$  that the infection-free equilibrium  $E_0^{yz}$  is always unstable. So we can easily get the infection-free equilibrium  $E_0^{yz}$  is always unstable when  $\tau > 0$ .

(ii) The characteristic equation of system (1.4) at  $E_0^z$  is

$$(\lambda^2 + d_1 R_0 \lambda + \beta^2 x_1 y_1) (\lambda - (r - d_3 + c y_1 e^{-\lambda \tau})) = 0.$$
(3.9)

Note that  $r > d_3$ . It can be shown that there exist positive real root for the characteristic equation above, which indicates that the immunity-inactivated infection equilibrium  $E_0^z$  is unstable.

(iii) The characteristic equation of system (1.4) at  $E_0^y$  is

$$(\lambda + d_1)(\lambda + r - d_3)(\lambda - \frac{s\beta}{d_1} + d_2 + \frac{pm(r - d_3)}{r}) = 0.$$
(3.10)

The eigenvalues are

$$\lambda_1 = -d_1, \lambda_2 = -(r-d_3), \lambda_3 = \frac{s\beta}{d_1} - d_2 - \frac{pm(r-d_3)}{r}.$$

Note that

$$\frac{s\beta}{d_1} - d_2 - \frac{pm(r-d_3)}{r} = d_2(R_0 - 1) - \frac{pm(r-d_3)}{r} < 0 \Leftrightarrow R_0 < 1 + \frac{pm(r-d_3)}{rd_2}.$$

Thus, the infection-free equilibrium  $E_0^y$  is locally asymptotically stable if  $R_0 < 1 + \frac{pm(r-d_3)}{rd_2}$ , and  $E_0^y$  is unstable if  $R_0 > 1 + \frac{pm(r-d_3)}{rd_2}$ .

Mathematical Biosciences and Engineering

In order to obtain the global stability of  $E_0^y$ , we construct the following Lyapunov function:

$$L_4 = x - x_2 - x_2 \ln \frac{x}{x_2} + y + \frac{p}{c}(z - z_2 - z_2 \ln \frac{z}{z_2}) + \frac{p}{c}z + p \int_{-\tau}^0 y(t + \theta)z(t + \theta)d\theta.$$

Taking the time derivative of  $L_4$  along the solution of system (1.4) and using a computation process similar to that of Theorem 3.9, we have

$$\begin{aligned} \frac{dL_4}{dt} &= (1 - \frac{x_2}{x})(s - \beta xy - d_1 x) + \beta xy - d_2 y - pyz \\ &+ \frac{p}{c}(1 - \frac{z_2}{z})\Big(cy(t - \tau)z(t - \tau) + rz(1 - \frac{z}{m}) - d_3 z\Big) \\ &= -\frac{d_1(x - x_2)^2}{x} + d_2(R_0 - 1)y - \frac{pz_2y(t - \tau)z(t - \tau)}{z} - \frac{rp}{cm}(z - z_2)^2. \end{aligned}$$

Thus,  $\frac{dL_4}{dt} \le 0$  if  $R_0 < 1$ , and  $\frac{dL_4}{dt} = 0$  only if  $x = x_2, y = 0$  and  $z = z_2$ , i.e., the maximal invariant subset in  $\{(x, y, z) : \frac{dL_4}{dt} |_{(1,4)} = 0\}$  is the singleton  $\{E_0^y\}$ . As a result,  $E_0^y$  is globally asymptotically stable based on the LaSalle's invariance principle. This complete the proof.

*Remark* 3.4. From Proposition 3.1 to Proposition 3.3, we can see that the delay  $\tau$  does not affect the stability of infection-free equilibrium  $E_0^{yz}$ ,  $E_0^y$  and immune-unactivated  $E_0^z$  equilibrium.

#### 4. Stability of positive equilibrium and Hopf bifurcation

In this section, we take the discrete delay  $\tau$  as a bifurcation parameter and show that when the positive equilibrium  $E_2^*$  loses its stability and a Hopf bifurcation appears when the delay  $\tau$  passes through a critical value. We point out there also exists a Hopf bifurcation at positive equilibrium  $E_1^*$  as the delay  $\tau$  passes through a critical value and the proof is similar.

The characteristic equation of system (1.4) at  $E_2^*$  is

$$\lambda^{3} + B_{1}\lambda^{2} + B_{2}\lambda + B_{3} + (C_{1}\lambda^{2} + C_{2}\lambda + C_{3})e^{-\lambda\tau} = 0,$$
(4.1)

where

$$B_{1} = \frac{2rz_{2}^{*}}{m} - r + d_{3} + d_{1} + \beta y_{2}^{*}, \qquad B_{2} = (d_{1} + \beta y_{2}^{*})(\frac{2rz_{2}^{*}}{m} - r + d_{3}) + \beta^{2} x_{2}^{*} y_{2}^{*},$$
  

$$B_{3} = (\frac{2rz_{2}^{*}}{m} - r + d_{3})\beta^{2} x_{2}^{*} y_{2}^{*}, \qquad C_{1} = -cy_{2}^{*},$$
  

$$C_{2} = -(d_{1} + \beta y_{2}^{*})cy_{2}^{*} + pcy_{2}^{*} z_{2}^{*}, \qquad C_{3} = -cy_{2}^{*}\beta^{2} x_{2}^{*} y_{2}^{*} + (d_{1} + \beta y_{2}^{*})pcy_{2}^{*} z_{2}^{*}.$$

When  $\tau = 0$ , the characteristic equation of system (1.4) at  $E_2^*$  is

$$\lambda^3 + A_1\lambda^2 + A_2\lambda + A_3 = 0,$$

where

$$A_1 \equiv B_1 + C_1 = d_1 + \beta y_2^* + c y_2^* + r - d_3 > 0,$$

Mathematical Biosciences and Engineering

$$\begin{split} A_2 &\equiv B_2 + C_2 = (d_1 + \beta y_2^*)(cy_2^* + r - d_3) + \beta^2 x_2^* y_2^* + p c y_2^* z_2^* > 0, \\ A_3 &\equiv B_3 + C_3 = p c y_2^* z_2^* (d_1 + \beta y_2^*) + (cy_2^* + r - d_3) \beta^2 x_2^* y_2^* > 0. \end{split}$$

After some calculations, we have

$$\begin{aligned} A_1 A_2 - A_3 &= (cy_2^* + r - d_3)(pcy_2^* z_2^* \\ &+ (d_1 + \beta y_2^*)^2) + (d_1 + \beta y_2^*)(\beta^2 x_2^* y_2^* + (cy_2^* + r - d_3)^2) > 0 \end{aligned}$$

It follows from the Routh-Hurwitz criterion that  $E_2^*$  is locally asymptotically stable when time delay is absent.

When  $\tau > 0$ , putting  $\lambda = i\omega$  into characteristic Eq (4.1) and separating real and imaginary parts, we have

$$B_1 \omega^2 - B_3 = (C_3 - C_1 \omega^2) \cos(\omega \tau) + C_2 \omega \sin(\omega \tau),$$
(4.2)

$$\omega^3 - B_2\omega = -(C_3 - C_1\omega^2)\sin(\omega\tau) + C_2\omega\cos(\omega\tau).$$
(4.3)

Since  $\sin^2(\omega \tau) + \cos^2(\omega \tau) = 1$ , squaring and adding the two Eqs (4.2) and (4.3), we have

$$\omega^6 + p_1 \omega^4 + p_2 \omega^2 + p_3 = 0, \tag{4.4}$$

where

$$p_1 = B_1^2 - 2B_2 - C_1^2,$$
  

$$p_2 = B_2^2 + 2C_1C_3 - C_2^2 - 2B_1B_3,$$
  

$$p_3 = B_3^2 - C_3^2.$$

Let  $u = \omega^2$ , Eq (4.4) becomes

$$G(u) = u^3 + p_1 u^2 + p_2 u + p_3 = 0.$$
(4.5)

If Eq (4.5) has a positive real root u, the characteristic Eq (4.1) has a purely imaginary root  $i\omega = i\sqrt{u}$ ; otherwise, Eq (4.1) has no purely imaginary root. Note that

$$G'(u) = 3u^2 + 2p_1u + p_2.$$
(4.6)

Let

$$\Delta = p_1^2 - 3p_2$$

Then we know that G(u) is monotonically increasing if  $\Delta \le 0$ , which indicates that Eq (4.5) has no positive root when  $p_3 \ge 0$  and  $\Delta \le 0$ .

If  $\Delta > 0$ , the function G(u) has two critical points

$$u^* = \frac{-p_1 + \sqrt{\Delta}}{3}, \qquad u^{**} = \frac{-p_1 - \sqrt{\Delta}}{3}.$$

Thus we know that Eq (4.5) has unique positive root  $u_0$  with  $G'(u_0) > 0$  if one of the following two conditions hold:

 $(H_1) \Delta > 0, p_3 < 0, u^{**} < 0 \text{ and } u^* > 0;$  $(H_2) \Delta > 0, p_3 < 0, u^* > 0 \text{ and } G(u^{**}) < 0.$ 

Equation (4.5) has two positive roots  $u_1 < u_2$  with  $G'(u_1) < 0$  and  $G'(u_2) > 0$  if the following assumption is satisfied:

 $(H_3) \Delta > 0, p_3 > 0, u^* > 0 \text{ and } G(u^*) < 0.$ 

Let  $\omega_k = \sqrt{u_k}$ , k = 0, 1, 2. It follows from Eqs (4.2) and (4.3) that the value of  $\tau$  associated with the purely imaginary root  $i\omega_k$  should satisfy

$$(B_1\omega_k^2 - B_3)(C_3 - C_1\omega_k^2) + (\omega_k^3 - B_2\omega_k)C_2\omega_k = ((C_3 - C_1\omega_k^2)^2 + C_2^2\omega_k^2)\cos(\omega_k\tau).$$

For k = 0, 1, 2, we define

$$\tau_n^k = \frac{1}{\omega_k} \arccos\left(\frac{(B_1\omega_k^2 - B_3)(C_3 - C_1\omega_k^2) + (\omega_k^3 - B_2\omega_k)C_2\omega_k}{(C_3 - C_1\omega_k^2)^2 + C_2^2\omega_k^2}\right) + \frac{2n\pi}{\omega_k}, n = 0, 1, 2\cdots.$$
(4.7)

Then at increasing sequences of  $\tau$  values,

$$\begin{aligned} \tau_0^0 &< \tau_1^0 < \tau_2^0 < \dots < \tau_n^0 \cdots, \\ \tau_0^1 &< \tau_1^1 < \tau_2^1 < \dots < \tau_n^1 \cdots, \\ \tau_0^2 &< \tau_1^2 < \tau_2^2 < \dots < \tau_n^2 \cdots, \end{aligned}$$

Eq (4.1) has purely imaginary roots  $i\omega_k$ , k = 0, 1, 2.

Now we consider the transversality conditions associated with Hopf bifurcation. Substituting  $\lambda(\tau)$  into Eq (4.1) and differentiating the resulting equation in  $\tau$ , we obtain

$$\left(\frac{d\lambda}{d\tau}\right)^{-1} = \frac{3\lambda^2 + 2B_1\lambda + B_2 + (2C_1\lambda + C_2)e^{-\lambda\tau} + (C_1\lambda^2 + C_2\lambda + C_3) \cdot (-\tau)e^{-\lambda\tau}}{\lambda e^{-\lambda\tau}(C_1\lambda^2 + C_2\lambda + C_3)} = \frac{(3\lambda^2 + 2B_1\lambda + B_2)e^{\lambda\tau}}{\lambda(C_1\lambda^2 + C_2\lambda + C_3)} + \frac{2C_1\lambda + C_2}{\lambda(C_1\lambda^2 + C_2\lambda + C_3)} - \frac{\tau}{\lambda}.$$

which indicates that

$$\begin{aligned} \operatorname{sign} \left\{ \frac{d(\operatorname{Re}\lambda)}{d\tau} \right\} \Big|_{\lambda = i\omega_{k}} &= \operatorname{sign} \left\{ \operatorname{Re} \left( \frac{d\lambda}{d\tau} \right)^{-1} \right\} \Big|_{\lambda = i\omega_{k}} \\ &= \operatorname{sign} \left\{ \frac{3\omega_{k}^{4} + 2(B_{1}^{2} - 2B_{2} - C_{1}^{2})\omega_{k}^{2} + (B_{2}^{2} + 2C_{1}C_{3} - C_{2}^{2} - 2B_{1}B_{3})}{(C_{3} - C_{1}\omega_{k}^{2})^{2} + C_{2}^{2}\omega_{k}^{2}} \right\} \\ &= \operatorname{sign} \left\{ \frac{G'(\omega_{k}^{2})}{(C_{3} - C_{1}\omega_{k}^{2})^{2} + C_{2}^{2}\omega_{k}^{2}} \right\}.\end{aligned}$$

Since  $G'(u_0) > 0$ ,  $G'(u_1) < 0$  and  $G'(u_2) > 0$ , then for n = 0, 1, 2, ..., we have

$$\operatorname{sign}\left\{\frac{d(\operatorname{Re}\lambda)}{d\tau}\Big|_{\tau=\tau_n^k}\right\} = \operatorname{sign}\left\{\operatorname{Re}\left(\frac{d\lambda}{d\tau}\right)^{-1}\Big|_{\tau=\tau_n^k}\right\} > 0, (k=0,2),$$

and

$$\operatorname{sign}\left\{\frac{d(\operatorname{Re}\lambda)}{d\tau}\Big|_{\tau=\tau_n^k}\right\} = \operatorname{sign}\left\{\operatorname{Re}\left(\frac{d\lambda}{d\tau}\right)^{-1}\Big|_{\tau=\tau_n^k}\right\} < 0, (k=1).$$

Mathematical Biosciences and Engineering

Then we know that at each  $\tau_n^0$  and  $\tau_n^2$ ,  $n = 0, 1, 2, \cdots$ , a pair of characteristic roots of Eq (4.1) cross the imaginary axis to the right. At each  $\tau_n^1$ ,  $n = 0, 1, 2, \cdots$ , a pair of characteristic roots of Eq (4.1) cross the imaginary axis to the left. Then the transversality condition required by the Hopf bifurcation theorem is satisfied.

Note that  $\omega_k = \sqrt{u_k}$  and  $u_1 < u_2$ . We have  $\omega_1 < \omega_2$ , which indicates that

$$\tau_n^1 - \tau_{n-1}^1 = \frac{2\pi}{\omega_1} > \frac{2\pi}{\omega_2} = \tau_n^2 - \tau_{n-1}^2, n = 0, 1, 2...$$

Then there exists a positive integer k such that

$$au_0^2 < au_0^1 < au_1^2 < au_1^1 \cdot \dots < au_k^2 < au_{k+1}^2 < au_k^1.$$

We thus obtain the following result.

**Theorem 4.1.** For system (1.4), we have

(*i*) If  $\Delta \leq 0$  and  $p_3 \geq 0$  holds,  $E_2^*$  is asymptotically stable for all  $\tau > 0$ .

(ii) If one of the conditions  $(H_1)$  and  $(H_2)$  holds,  $E_2^*$  is asymptotically stable when  $\tau \in [0, \tau_0^0)$  and unstable when  $\tau > \tau_0^0$ , that is system (1.4) undergoes a Hopf bifurcation at  $E_2^*$  when  $\tau = \tau_0^0$ .

(iii) If the condition (H<sub>3</sub>) holds, system (1.4) undergoes a Hopf bifurcation at  $E_2^*$  along two sequences of  $\tau$  values  $\tau_n^{1,2}$ , n = 0, 1, 2... Furthermore there exists a positive integer k such that  $E_2^*$  is stable when

$$au \in [0, au_0^2) \cup ( au_0^1, au_1^2) \cup \dots \cup ( au_{k-1}^1, au_k^2);$$

 $E_2^*$  is unstable when

$$\tau \in (\tau_0^2, \tau_0^1) \cup (\tau_1^2, \tau_1^1) \cup \dots \cup (\tau_{k-1}^2, \tau_{k-1}^1) \cup (\tau_k^2, +\infty).$$

#### 5. Direction and stability of the Hopf bifurcation

In the previous section, we know that system (1.4) undergoes a Hopf bifurcation at  $E_2^*$  along some sequences of  $\tau$  values. Let  $\tau^*$  be one of the Hopf bifurcation points. As pointed out in Hassard et al. [31], it is interesting to determine the direction, stability and period of these periodic solutions.

Let  $\mu = \tau - \tau^*$ , and then  $\mu$  is new bifurcation parameter of the system. Define

$$X(t) = (x - x_2^*, y - y_2^*, z - z_2^*)^{\mathrm{T}}, X_t(\theta) = X(t + \theta), \theta \in [-\tau, 0].$$

System (1.4) may be written as:

$$X'(t) = L_{\mu}X_{t} + f(X_{t}(.),\mu), \quad L_{\mu}\phi = F_{1}\phi(0) + F_{2}\phi(-\tau), \tag{5.1}$$

where

$$f(\phi,\mu) = \begin{bmatrix} -\beta\phi_1(0)\phi_2(0) \\ \beta\phi_1(0)\phi_2(0) - p\phi_2(0)\phi_3(0) \\ c\phi_2(-\tau)\phi_3(-\tau) - \frac{r}{m}\phi_3^{-2}(0) \end{bmatrix},$$

Mathematical Biosciences and Engineering

$$F_{1} = \begin{bmatrix} -\beta y_{2}^{*} - d_{1} & -\beta x_{2}^{*} & 0\\ \beta y_{2}^{*} & -d_{2} - pz_{2}^{*} & -py_{2}^{*}\\ 0 & 0 & r - d_{3} - \frac{2rz_{2}^{*}}{m} \end{bmatrix}$$

and

$$F_2 = \left[ \begin{array}{ccc} 0 & 0 & 0 \\ 0 & 0 & 0 \\ 0 & cz_2^* & cy_2^* \end{array} \right].$$

By Riesz representation theorem, there exists a matrix  $\eta(\theta, \mu) : [-\tau, 0] \to \mathbb{R}^3$  whose components are bounded variation functions such that

$$L_{\mu}\phi = \int_{-\tau}^{0} d\eta(\theta,\mu)\phi(\theta),$$

where

$$d\eta(\theta,\mu) = F_1 \delta(\theta) d\theta + F_2 \delta(\theta+\tau) d\theta$$

and  $\delta(\theta)$  is the Dirac delta function.

For  $\phi \in C([-\tau, 0], R^3)$ , we define

$$A(\mu)\phi(\theta) = \begin{cases} \frac{d\phi(\theta)}{d\theta}, & \theta \in [-\tau, 0) \\ \int_{-\tau}^{0} d\eta(\xi, \mu)\phi(\xi), & \theta = 0, \end{cases}$$

and

$$R(\mu)\phi(\theta) = \begin{cases} 0, & \theta \in [-\tau, 0), \\ f(\phi, \mu), & \theta = 0. \end{cases}$$

Then system (1.4) is equivalent to the following operator equation

$$X'_{t}(\theta) = A(\mu)X_{t}(\theta) + R(\mu)X_{t}(\theta).$$
(5.2)

,

For  $\varphi \in C([0, \tau], \mathbb{R}^3)$ , we define the adjoint operator of A(0) as  $A^*(0)$ , where

,

$$A^*(0)\varphi(s) = \begin{cases} -\frac{d\varphi(s)}{ds}, & s \in (0,\tau], \\ \int_{-\tau}^0 d\eta^{\mathrm{T}}(\xi,0)\phi(-\xi), & s = 0. \end{cases}$$

For the convenience of research, we simply write *A* for *A*(0), *A*<sup>\*</sup> for *A*<sup>\*</sup>(0), *R* for *R*(0),  $\eta(\theta)$  for  $\eta(\theta, 0)$ , for  $\varphi \in C([0, \tau], R^3)$  and  $\phi \in C([-\tau, 0], R^3)$ .

Define a bilinear form as

$$\langle \varphi, \phi \rangle = \bar{\varphi}^{\mathrm{T}}(0)\phi(0) - \int_{\theta=-\tau}^{0} \int_{\xi=0}^{\theta} \bar{\varphi}^{\mathrm{T}}(\xi-\theta)d\eta(\theta)\phi(\xi)d\xi.$$

Mathematical Biosciences and Engineering

Let  $q(\theta)$  and  $q^*(s)$  to be the eigenvectors of matric A and A<sup>\*</sup> corresponding to eigenvalue  $i\omega_0$  and  $-i\omega_0$ , respectively. Then

$$Aq(\theta) = i\omega_0 q(\theta), \qquad A^*q^*(s) = -i\omega_0 q^*(s).$$

We can choose appropriate  $q(\theta)$  and  $q^*(s)$  such that  $\langle q(\theta), q^*(s) \rangle = 1$ , where

$$q(\theta) = (1, q_2, q_3)^{\mathrm{T}} e^{i\omega_0 \theta}, \quad q^*(s) = D(1, q_2^*, q_3^*)^{\mathrm{T}} e^{i\omega_0 s},$$

and

$$\begin{aligned} q_2 &= \frac{\beta y_2^* + d_1 + i\omega_0}{-\beta x_2^*}, \qquad q_3 = \frac{\beta^2 y_2^* x_2^* + (\beta y_2^* + d_1 + i\omega_0)(d_2 + pz_2^* + i\omega_0)}{p y_2^* \beta x_2^*}, \\ q_2^* &= \frac{\beta y_2^* + d_1 - i\omega_0}{\beta y_2^*}, \qquad q_3^* = \frac{[\beta^2 y_2^* x_2^* + (\beta y_2^* + d_1 - i\omega_0)(d_2 + pz_2^* - i\omega_0)]e^{i\omega_0 \tau^*}}{c\beta y_2^* z_2^*} \end{aligned}$$

Note that

$$< q, q^* > = \bar{D}\bar{q}^{\mathrm{T}}(0)q^*(0) - \int_{\theta=-\tau^*}^{0} \int_{s=0}^{\theta} \bar{D}\bar{q}^{\mathrm{T}}(s-\theta)d\eta(\theta)q^*(s)ds$$

$$= \bar{D}(1+\bar{q}_2q_2^*+\bar{q}_3q_3^* - \int_{\theta=-\tau^*}^{0} \int_{s=0}^{\theta} (1,\bar{q}_2,\bar{q}_3)e^{-(s-\theta)i\omega_0}(1,q_2^*,q_3^*)^{\mathrm{T}}e^{i\omega_0s}dsd\eta(\theta))$$

$$= \bar{D}(1+\bar{q}_2q_2^*+\bar{q}_3q_3^* - \int_{\theta=-\tau^*}^{0} (c\bar{q}_2z^*+cq_3^*y^*)\theta e^{i\omega_0\theta}d\eta(\theta))$$

$$= \bar{D}(1+\bar{q}_2q_2^*+\bar{q}_3q_3^* + (c\bar{q}_2z^*+cq_3^*y^*)q_3^*\tau^*e^{i\omega_0\tau^*}).$$

Then we can choose  $\overline{D}$  as

$$\bar{D} = [1 + \bar{q}_2 q_2^* + \bar{q}_3 q_3^* + (c\bar{q}_2 z_2^* + cq_3^* y_2^*) q_3^* \tau^* e^{i\omega_0 \tau^*}]^{-1}$$

such that  $\langle q(\theta), q^*(s) \rangle = 1$ .

According to the notations in Hassard et al. [31], we need to compute the center manifold  $C_0$  at  $\mu = 0$ . Let  $u_t$  be the solution of Eq (5.1) when  $\mu = 0$  and define

$$Z(t) = \langle q^*, u_t \rangle, \quad W(t,0) = u_t(0) - 2Re\{Z(t)q(\theta)\}.$$
(5.3)

On the center manifold  $C_0$ , we have  $W(t, 0) = W(Z(t), \overline{Z}(t), \theta)$ , where

$$W(Z, \bar{Z}, \theta) = W_{20}(\theta) \frac{Z^2}{2} + W_{11}(\theta) Z \bar{Z} + W_{02}(\theta) \frac{\bar{Z}^2}{2} + \cdots,$$

Z and  $\overline{Z}$  are local coordinates for center manifold  $C_0$  in the direction of  $q^*$  and  $\overline{q}^*$ . Note that if  $X_t$  is real, W is real. We only consider the real solutions. For the solution of the Eq (5.1)  $X_t \in C_0$ , we have

$$\dot{Z}(t) = i\omega Z + \bar{q}^*(\theta) f(0, W(Z, \bar{Z}, \theta)) + 2\operatorname{Re}\{Zq(\theta)\} \stackrel{\scriptscriptstyle \triangle}{=} i\omega Z + \bar{q}^*(0) f_0(Z, \bar{Z}).$$

Now we rewrite this equation as  $\dot{Z}(t) = i\omega Z + g(Z, \bar{Z})$ , where

$$g(Z,\bar{Z}) = \bar{q}^*(0)f_0(Z,\bar{Z}) = g_{20}\frac{Z^2}{2} + g_{11}Z\bar{Z} + g_{02}\frac{\bar{Z}^2}{2} + g_{21}\frac{Z^2\bar{Z}}{2} + \cdots$$
(5.4)

Mathematical Biosciences and Engineering

Note that  $u_t(\theta) = W(t, \theta) + Zq(\theta) + \overline{Z}\overline{q}(\theta)$ . We have

$$\begin{split} g(Z,\bar{Z}) &= \bar{q}^*(0) f_0(Z,\bar{Z}) \\ &= \bar{D} \big\{ -\beta u_{1t}(0) u_{2t}(0) + q_2^* \big( \beta u_{1t}(0) u_{2t}(0) - p u_{2t}(0) u_{3t}(0) \big) + q_3^* \big( c u_{2t}(-\tau) u_{3t}(-\tau) - \frac{r}{m} u_{3t}^2(0) \big) \big\} \\ &= \bar{D} \big\{ (q_2^* - \beta)(Z + \bar{Z} + W_{20}^{(1)}(0) \frac{Z^2}{2} + W_{11}^{(1)}(0) Z\bar{Z} + W_{02}^{(1)}(0) \frac{\bar{Z}^2}{2}) (q_2 Z + \bar{q}_2 \bar{Z} + W_{20}^{(3)}(0) \frac{Z^2}{2} \\ &+ W_{11}^{(3)}(0) Z\bar{Z} + W_{02}^{(3)}(0) \frac{\bar{Z}^2}{2}) - p q_2^* (q_2 Z + \bar{q}_2 \bar{Z} + W_{20}^{(2)}(0) \frac{Z^2}{2} + W_{11}^{(2)}(0) Z\bar{Z} + W_{02}^{(2)}(0) \frac{\bar{Z}^2}{2}) (q_3 Z + \bar{q}_3 \bar{Z} + W_{20}^{(3)}(0) \frac{Z^2}{2} + W_{11}^{(3)}(0) Z\bar{Z} + W_{02}^{(3)}(0) \frac{\bar{Z}^2}{2}) + q_3^* c (q_2 Z e^{-i\omega_0 \tau} + \bar{q}_2 \bar{Z} e^{-i\omega \tau} + W_{20}^{(2)}(-\tau) \frac{Z^2}{2} \\ &+ W_{11}^{(2)}(-\tau) Z\bar{Z} + W_{02}^{(2)}(-\tau) \frac{\bar{Z}^2}{2}) (q_4 Z e^{-i\omega_0 \tau} + \bar{q}_4 \bar{Z} e^{-i\omega \tau} + W_{20}^{(3)}(-\tau) \frac{Z^2}{2} + W_{11}^{(3)}(-\tau) Z\bar{Z} \\ &+ W_{02}^{(3)}(-\tau) \frac{\bar{Z}^2}{2}) - \frac{r}{m} q_3^* (q_3 Z + \bar{q}_3 \bar{Z} + W_{20}^{(3)}(0) \frac{Z^2}{2} + W_{11}^{(3)}(0) Z\bar{Z} + W_{02}^{(3)}(0) \frac{Z^2}{2})^2 \big\}. \end{split}$$

Comparing the coefficients with Eq (5.4), we have

$$g_{20} = 2\bar{D}\{(q_{2}^{*} - \beta)q_{2} + q_{3}^{*}q_{2}q_{3}ce^{-2i\omega\tau} - q_{2}^{*}q_{2}q_{3}p - \frac{r}{m}q_{3}^{*}q_{3}^{2}\},\$$

$$g_{11} = 2\bar{D}\{(q_{2}^{*} - \beta)\operatorname{Re}\{q_{2}\} + q_{3}^{*}c\operatorname{Re}\{\bar{q}_{2}q_{3}\}e^{-2i\omega\tau} - q_{2}^{*}p\operatorname{Re}\{\bar{q}_{2}q_{3}\} - \frac{r}{m}q_{3}^{*}q_{3}\bar{q}_{3}\},\$$

$$g_{02} = 2\bar{D}\{(q_{2}^{*} - \beta)\bar{q}_{2} + q_{3}^{*}\bar{q}_{2}\bar{q}_{3}ce^{-2i\omega\tau} - q_{2}^{*}\bar{q}_{2}\bar{q}_{3}p - \frac{r}{m}q_{3}^{*}\bar{q}_{3}^{2}\},\$$

$$g_{21} = 2\bar{D}\{(q_{2}^{*} - \beta)(W_{11}^{(2)}(0) + \frac{W_{20}^{(2)}(0)}{2} + \frac{\bar{q}_{2}W_{20}^{(1)}(0)}{2} + q_{2}W_{11}^{(1)}(0))\$$

$$+ q_{3}^{*}c(q_{2}e^{-i\omega\tau}W_{11}^{(3)}(-\tau) + q_{3}e^{-i\omega\tau}W_{11}^{(2)}(-\tau) + \frac{\bar{q}_{2}W_{20}^{(3)}(-\tau)}{2}e^{-i\omega\tau} + \frac{\bar{q}_{3}W_{20}^{(2)}(-\tau)}{2}e^{-i\omega\tau})\$$

$$- q_{2}^{*}p(q_{2}W_{11}^{(3)}(0) + \frac{\bar{q}_{2}W_{20}^{(3)}(0)}{2} + \frac{\bar{q}_{3}W_{20}^{(2)}(0)}{2} + q_{3}W_{11}^{(2)}(0))\$$

$$- \frac{r}{m}q_{3}^{*}(2q_{3}W_{11}^{(3)}(0) + \bar{q}_{3}W_{20}^{(3)}(0))\}.$$
(5.5)

It remains to compute  $W_{11}(0)$  and  $W_{20}(0)$  in  $g_{21}$ . From Eqs (5.2) and (5.3), we have

$$\dot{W} = \dot{x}_t - \dot{Z}q - \overline{\dot{Z}q} = \begin{cases} AW - 2\operatorname{Re}\{\bar{q}^*(0)f_0q(\theta)\}, & \theta \in [-1,0), \\ AW - 2\operatorname{Re}\{\bar{q}^*(0)f_0q(0)\} + f_0, & \theta = 0. \end{cases}$$
(5.6)

We rewrite the Eq (5.6) as

$$\dot{W} \stackrel{\scriptscriptstyle \Delta}{=} AW + H(Z, \bar{Z}, \theta) \tag{5.7}$$

where

$$H(Z, \bar{Z}, \theta) = H_{20}(\theta) \frac{Z^2}{2} + H_{11}(\theta) Z \bar{Z} + H_{02}(\theta) \frac{\bar{Z}^2}{2} + \cdots$$

Thus

$$(A - 2i\omega)W_{20}(\theta) = -H_{20}(\theta), \quad AW_{11}(\theta) = -H_{11}(\theta).$$
 (5.8)

Mathematical Biosciences and Engineering

From Eq (5.7), we know that for  $\theta \in [-1, 0)$ 

$$H(Z,\bar{Z},\theta) = -\bar{q}^*(0)f_0q(\theta) - q^*(0)\bar{f}_0\bar{q}(\theta) = -gq(\theta) - \bar{g}\bar{q}(\theta).$$

Comparing the coefficients with Eq (5.8), we obtain

$$H_{20}(\theta) = -g_{20}q(\theta) - \bar{g}_{20}\bar{q}(\theta), \tag{5.9}$$

$$H_{11}(\theta) = -g_{11}q(\theta) - \bar{g}_{11}\bar{q}(\theta).$$
(5.10)

From Eqs (5.8) and (5.9) and the definition of A, we have

$$\dot{W}_{20}(\theta) = 2i\omega W_{20}(\theta) + g_{20}q(\theta) + \bar{g}_{02}\bar{q}(\theta),$$

where  $q(\theta) = (1, q_2, q_3)^{\mathrm{T}} e^{i\theta\omega\tau}$ . Hence

$$W_{20}(\theta) = \frac{ig_{20}}{\omega}e^{i\omega\theta}q(0) + \frac{i\bar{g}_{02}}{3\omega}e^{-i\omega\theta}\bar{q}(0) + E_{20}e^{2i\omega\theta}$$

where

$$E_{20} = 2(2i\omega - F_1 - F_2 e^{-2i\omega\tau})^{-1} \begin{bmatrix} -\beta q_2 \\ \beta q_2 - p q_2 q_3 \\ c q_2 q_3 e^{-2i\omega\tau} - \frac{r}{m} q_3^2 \end{bmatrix}.$$

Similarly, from Eqs (5.8) and (5.10) we obtain

$$W_{11}(\theta) = \frac{ig_{11}}{\omega}e^{i\omega\theta}q(0) + \frac{i\bar{g}_{11}}{\omega}e^{-i\omega\theta}\bar{q}(0) + E_{11},$$

where

$$E_{11} = 2(-F_1 - F_2)^{-1} \begin{bmatrix} -\beta Re\{q_2\} \\ \beta Re\{q_2\} - pRe\{q_2\bar{q}_3\} \\ cRe\{q_2\bar{q}_3\} - \frac{r}{m}\{q_3^2\} \end{bmatrix}.$$

So far, we have calculated  $g_{20}$ ,  $g_{11}$ ,  $g_{02}$ ,  $g_{21}$  in Eq (5.5) and then we can obtain

$$c_{1}(0) = \frac{i}{2\omega} \left( g_{11}g_{20} - 2 |g_{11}|^{2} - \frac{|g_{02}|^{2}}{3} \right) + \frac{g_{21}}{2},$$

$$v_{2} = -\frac{\operatorname{Re}(c_{1}(0))}{\operatorname{Re}(\lambda'(\tau^{*}))},$$

$$\beta_{2} = 2\operatorname{Re}(c_{1}(0)),$$

$$T_{2} = \frac{-(\operatorname{Im}\{c_{1}(0)\} + v_{2}\operatorname{Im}\{\lambda'(\tau^{*})\})}{\omega}.$$
(5.11)

It is well known that  $v_2$  and  $\beta_2$  will determine the direction and stability of the Hopf bifurcation, and  $T_2$  determines the period of the bifurcated periodic solutions, respectively. In particular, the Hopf bifurcation is supercritical (subcritical) if  $v_2 > 0(v_2 < 0)$ , and the bifurcated periodic solutions exist for  $\tau > \tau_0(\tau < \tau_0)$ . The bifurcated periodic solutions are stable (unstable) if  $\beta_2 < 0$  ( $\beta_2 > 0$ ) and the period will become longer (shorter) if  $T_2 > 0$  ( $T_2 < 0$ ).

#### 6. Numerical simulations

In this section, we carry out some numerical simulations to display some qualitative behaviours of system (1.4). Table 1 lists the values or ranges of parameters for system (1.4) referring to [21].

We first study the effect of logistic growth on the dynamics of system (1.4). According to the ranges of parameters in Table 1, we take the following parameters

$$s = 10, \beta = 0.02, d_1 = 0.2, d_2 = 0.5, d_3 = 0.2, p = 0.05, c = 0.4, m = 20.$$
 (6.1)

Figure 1 presents the bifurcation diagrams of the solutions for system (1.4) with respect to  $\tau$  and different logistic growth rate r. One can see that the positive equilibrium  $E_1^*$  of system (1.4) is local asymptotically stable when  $\tau < \tau^* = 0.8$  and the bifurcated periodic solutions occurs through Hopf bifurcations when  $\tau > \tau^*$ . Similarly, the positive equilibrium  $E_2^* = (49.42, 0.12, 9.77)$  of system (1.4) is local asymptotically stable when  $\tau = 8 < \tau^* = 8.655$  and the bifurcated periodic solutions occur through Hopf bifurcations when  $\tau = 9 > \tau^*$ . At the same time, one can note that, except r = 0, the values of Hopf bifurcation points increase with the increase of r. On the other hand, the amplitude of the bifurcated periodic solution increase with the increase of time delay  $\tau$  and decrease with the increase of r.

Parameters	Descriptions	Ranges	Units
S	Source rate of uninfected cell	1–10	cells mL <sup>-1</sup> day <sup>-1</sup>
β	Virus-to-cell infection rate	0.00025-0.5	mL virion <sup>-1</sup> day <sup>-1</sup>
р	Predation rate of infection cell by CTLs	$1-4.048 \times 10^{-4}$	mL cell <sup>-1</sup> day <sup>-1</sup>
r	Breath rate of CTLs	0.0051-3.912	mL cell <sup>-1</sup> day <sup>-1</sup>
т	Capacity of immune cells	6.25-235999.9	mL cell <sup><math>-1</math></sup>
С	Development rate of CTLs	0.0051-3.912	mL cell <sup>-1</sup> day <sup>-1</sup>
$d_1$	Death rate of uninfected cell	0.007-0.1	$day^{-1}$
$d_2$	Death rate of infected cell	0.2–0.3	$day^{-1}$
$d_3$	Death rate of immune cell	0.001-0.3	day <sup>-1</sup>

Table 1. Meanings and units of parameters.

Figure 2 presents phase diagrams of the solutions for system (1.4) with r = 0.3 and different values of  $\tau$ . One can see that the positive equilibrium  $E_2^* = (49.42, 0.12, 9.77)$  is local asymptotically stable when  $\tau = 8 < \tau^* = 8.655$  and the bifurcated periodic solutions occur through Hopf bifurcations when  $\tau = 9 > \tau^*$ . Furthermore, using the given parameter values (6.1), we can obtain  $c_1(0) = -19.651 +$ 26.769*i* by some calculations and then we know that  $\mu_2 > 0$ ,  $\beta_2 < 0$ ,  $T_2 < 0$  by (5.11). According to the conclusion in [31], we know that the Hopf bifurcation at  $\tau^* = 8.655$  is supercritical, and the bifurcating periodic solution is stable.

In order to investigate the stability switch of equilibrium, we take the following parameters

$$s = 10, \beta = 0.2, d_1 = 0.2, d_2 = 0.3, d_3 = 0.2, p = 0.05, c = 0.5, r = 1.6, m = 50.$$
 (6.2)

We obtain there exists positive equilibrium  $E_2^* = (18.8775, 1.6487, 69.5102)$  and the characteristic equation of system (1.4) have two pure virtual roots  $\omega_1 = 0.4464$  and  $\omega_2 = 1.4748$ . Then by (4.7) we

get

$$\tau_0^1 = 0.8365, \tau_1^1 = 8.1911, \tau_2^1 = 12.4513;$$

and

$$\tau_0^2 = 3.9309, \tau_1^2 = 14.9113.$$

Noting that  $\tau_1^2 > \tau_2^1$ , we know that  $E_2^*$  is asymptotically stable when  $\tau \in [0, \tau_0^1) \cup (\tau_0^2, \tau_1^1)$  and is unstable when  $\tau \in (\tau_0^1, \tau_0^2) \cup (\tau_1^1, +\infty)$ . Figure 3 presents time series diagrams of the solutions for system (1.4) with different values of  $\tau$ . One can see that there exists stability switch of equilibrium  $E_2^*$  as  $\tau$  increases and some periodic solutions are bifurcated by Hopf bifurcation.



**Figure 1.** Bifurcation diagrams of system (1.4) with respect to  $\tau$  and different *r*. Parameter values are given by (6.1).



Figure 2. Phase diagrams of the solutions for system (1.4) with r = 0.3 and different values of  $\tau$ . The other parameters values are given by (6.1).



**Figure 3.** Time series diagrams of the solutions for system (1.4) showing stability switch with increase of  $\tau$ . Parameter values are given by (6.2).

	<b>a</b> 11.1	
Case	Conditions	Equilibria and its stability
r = 0	$R_0 < 1$	$E_0^{yz}$ is GAS
	$1 < R_0 < 1 + \frac{\beta d_3}{cd_1}$	$E_0^{yz}$ is US, $E_0^z$ is GAS
	$R_0 > 1 + \frac{\beta d_3}{c d_1}$	$E_0^{yz}$ and $E_0^z$ are US, $E^*$ (Hopf)
$r < d_3$	$R_0 < 1$	$E_0^{yz}$ is GAS
	$1 < R_0 < 1 + \frac{\beta(d_3 - r)}{cd_1}$	$E_0^{yz}$ is US, $E_0^z$ is GAS
	$R_0 > 1 + \frac{\beta(d_3 - r)}{cd_1}$	$E_0^{yz}$ and $E_0^z$ are US, $E^*$ (Hopf)
$r = d_3$	$R_0 < 1$	$E_0^{yz}$ is GAS
	$R_0 > 1$	$E_0^{\tilde{y}_z}$ and $E_0^z$ are US, $E^*$ (Hopf)
$r > d_3$	$R_0 < 1$	$E_0^{yz}$ is US, $E_0^y$ is GAS
	$1 < R_0 < 1 + \frac{pm(r-d_3)}{rd_2}$	$E_0^{yz}$ and $E_0^z$ are US, $E_0^y$ is GAS
	$R_0 > 1 + \frac{pm(r-d_3)}{rd_2}$	$E_0^{yz}$ , $E_0^z$ and $E_0^y$ are US, $E^*$ (Hopf)

**Table 2.** Global properties of system (1.4) with  $\tau > 0$ .

Note: GAS means globally asymptotically stable; US means unstable.

## 7. Conclusions and discussions

In this paper, a viral infection model with self-proliferation of cytotoxic T lymphocytes (CTLs) and activation time delay of immune cells is proposed. We mainly focus on two topics. First, we study the global dynamics of system (1.4) through constructing appropriate Lyapunov functions. Then we examine the impact of the activation time delay of immune cells on the existence of periodic solutions. The global dynamical properties of system (1.4) can be summarized in the following

Table 2. Numerical simulations verify the theoretical analyses. In particular, we find that for different values of the delay in CTL response, the system can stabilize at the positive equilibrium when the delay is small, or stabilize at a stable periodic oscillation when the delay is large. The amplitudes of bifurcated periodic solutions increase with the increase of activation time delay of immune cells and decrease with the increase of r (Figure 1). It is also found that there exists stability switch phenomenon under some conditions (Figure 3). The results indicate that the self-proliferation intensity and activation time delay of immune cells can significantly affect the kinetics of viral infection.

Some aspects of the viral infection problem remain to be studied in the future. For instance, we plan to extend our analysis to global Hopf bifurcation analysis. Some other factors that influence the dynamics of viral infection including the heterogeneity of space and movement of cells may be investigated.

## Acknowledgments

The authors would like to thank the reviewers and the editor for their careful reading, helpful comments and suggestions that greatly improved the paper. This work is supported by the National Natural Science Foundation of China (Grant Nos. 11701472, 11771448, 11871403).

## **Conflict of interest**

The authors declare that they have no conflict of interest.

# References

- J. E. Schmitz, M. J. Kuroda, S. Santra, V. G. Sasseville, M. A. Simon, M. A. Lifton, et al, Control of viremia in simian immunodeficiency virus infection by CD8+ lymphocytes, *Science*, 283 (1999), 857–860.
- 2. L. C. Wang, M. Y. Li, Mathematical analysis of the global dynamics of a model for HIV infection of CD4+ T cells, *Math. Biosci.*, **200** (2006), 44–57.
- 3. X. Y. Song, A. U. Neumann, Global stability and periodic solution of the viral dynamics, *J. Math. Anal. Appl.*, **329** (2007), 281–297.
- 4. R. J. De Boer, A. S. Perelson, Target cell limited and immune control models of HIV infection: A comparison, *J. Theoret. Biol.*, **190** (1998), 201–214.
- 5. Y. Nakata, Global dynamics of a cell mediated immunity in viral infection models with distributed delays, *J. Math. Anal. Appl.*, **375** (2011), 14–27.
- J. L. Wang, J. M. Pang, T. Kuniya, Global threshold dynamics in a five-dimensional virus model with cell-mediated, humoral immune responses and distributed delays, *Appl. Math. Comput.*, 241 (2014), 298–316.
- 7. M. A. Nowak, S. Bonhoefier, A. M. Hill, R. Boehme, H. C. Thomas, H. McDade, Viral dynamics in hepatitis B virus infection, *Proc. Natl. Acad. Sci. USA*, **93** (1996), 4398–4402.

- 8. A. Korobeinikov, S. Giles, Global properties of basic virus dynamics models, *Bull. Math. Biol.*, **66** (2004), 879–883.
- 9. M. A. Nowak, C. R. M. Bangham, Population dynamics of immune responses to persistent viruses, *Science*, **272** (1996), 74–79.
- 10. H. Zhu, X. Zou, Dynamics of a HIV-1 Infection model with cell-mediated immune response and intracellular delay, *Discr. Cont. Dyn. Syst. Ser. B*, **12** (2009), 511–524.
- 11. K. Wang, W. Wang, H. Pang, X. Liu, Complex dynamic behavior in a viral model with delayed immune response, *Physica D*, **226** (2007), 197–208.
- 12. Yukihiko Nakata, Global dynamics of a cell mediated immunity in viral infection models with distributed delays, *J. Math. Anal. Appl.*, **375** (2011), 14–27.
- H. Gomez-Acevedo, M. Y. Li, S. Jacobson, Multi-stability in a model for CTL response to HTLV-I infection and its consequences in HAM/TSP development and prevention, *Bull. Math. Biol.*, 72 (2010), 681–696.
- 14. R. M. Anderson, R. M. May, S. Gupta, Non-linear phenomena in host-parasite interactions, *Parasitology*, **99** (1989), 59–79.
- 15. A. Murase, T. Sasaki, T. Kajiwara, Stability analysis of pathogen-immune interaction dynamics, *J. Math. Biol.*, **51** (2005), 247–267.
- 16. C. Chiyaka, W. Garira, S. Dube, Modelling immune response and drug therapy in human malaria infection, *Comput. Math. Method. Med.*, **9** (2008), 143–163.
- 17. A. S. Perelson, Modelling viral and immune system dynamics, *Nature Rev. Immunol.*, **2** (2002), 28–36.
- 18. A. Korobeinikov, Immune response and within-host viral evolution:Immune response can accelerate evolution, *J. Theor. Biol.*, **456** (2018),74–83.
- 19. H. Q. Zhang, H. Chen, C. C Jiang, K. F. Wang, Effect of explicit dynamics of free virus and intracellular delay, *Chaos, Solitons Fractals*, **104** (2017), 827–834.
- 20. Y. Wang, J. Liu, J. M. Heffernan, Viral dynamics of an HTLV-I infection model with intracellular delay and CTL immune response delay, *J. Math. Anal. Appl.*, **459** (2018), 506–527.
- 21. K. Allali, S. Harroudi, D. F. M. Torre, Analysis and optimal control of an intracellular delayed HIV model with CTL immune response, *Math. Comput. Sci.*, **12** (2018), 111–127.
- 22. H. J. Liu, J. F. Zhang, Dynamics of two time delays differential equation model to HIV latent infection, *Physica A*, **514** (2019), 384–395.
- 23. M. Y. Li, H. Shu, Multiple stable periodic oscillations in a mathematical model of CTL response to HTLV-I infection, *Bull. Math. Biol.*, **73** (2011), 1774–1793.
- 24. D. W. Huang, X. Zhang, Y. F. Guo, H. L. Wang, Analysis of an HIV infection model with treatment sand delayed immune response, *Appl. Math. Model.*, **40** (2016), 3081–3089.
- 25. D. Wodarz, J. P. Christensen, A. R. Thomsen, The importance of lytic and nonlytic immune responses in viral infections, *Trends Immunol.*, **23** (2002), 194–200.

- 26. C. Bartholdy, J. P. Christensen, D. Wodarz, A. R. Thomsen, Persistent virus infection despite chronic cytotoxic T-lymphocyte activation in Gamma interferon-deficient mice infected with lymphocytic chroniomeningitis virus, *J. Virology*, **74** (2000), 10304–10311.
- 27. K. Wang, Y. Kuang, Fluctuation and extinction dynamics in host-microparasite systems, *Comm. Pure Appl. Anal.*, **10** (2011), 1537–1548.
- 28. S. Bonhoeffer, J. M. Coffin, M. A. Nowak, Human immunodeficiency virus drug therapy and virus load, *J. Virology*, **71** (1997), 3275–3278.
- 29. M. Nagumo, Uber die lage der integralkurven gewohnlicher differentialgleichungen, *Proc. Phys. Math. Soc.*, **24** (1942), 551–559.
- 30. P. van den Driessche, J. Watmough, Reproduction numbers and sub-threshold endemic equilibria for compartmental models of disease transmission, *Math. Biosci.*, **180** (2002), 29–48.
- 31. B. Hassard, D. Kazarinoff, Y. Wan, *Theory and Applications of Hopf Bifurcation*, Cambridge: Cambridge University Press, 1981.



 $\bigcirc$  2020 the Author(s), licensee AIMS Press. This is an open access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/4.0)