

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

MBE, 19(2): 1154–1173. DOI: 10.3934/mbe.2022053 Received: 21 October 2021 Accepted: 24 November 2021 Published: 30 November 2021

Research article

Stability and bifurcation analysis of a tumor-immune system with two delays and diffusion

Yuting Ding, Gaoyang Liu and Yong An*

College of Science, Northeast Forestry University, Harbin, 150040, China

* Correspondence: E-mail: anyong@nefu.edu.cn.

Abstract: A tumor-immune system with diffusion and delays is proposed in this paper. First, we investigate the impact of delay on the stability of nonnegative equilibrium for the model with a single delay, and the system undergoes Hopf bifurcation when delay passes through some critical values. We obtain the normal form of Hopf bifurcation by applying the multiple time scales method for determining the stability and direction of bifurcating periodic solutions. Then, we study the tumor-immune model with two delays, and show the conditions under which the nontrivial equilibria are locally asymptotically stable. Thus, we can restrain the diffusion of tumor cells by controlling the time delay associated with the time of tumor cell proliferation and the time of immune cells recognizing tumor cells. Finally, numerical simulations are presented to illustrate our analytic results.

Keywords: reaction-diffusion equation; tumor-immune model; delay; Hopf bifurcation

1. Introduction

Cancer, as the second-most common fatal disease, remains a serious threat to human health, and millions of people suffer from cancer around the world. Cancer cells proliferate and spread rapidly, which causes many difficulties in treatment. The function of the immune system is to prevent tumor cells from invading the body, and the body restricts the growth of tumor cells. In view of molecular cell biology, the immune system has a natural defense for the body, since it can recognize and destroy tumor cells [1,2]. It is obvious that immune system treatment is effective for the treatment of cancer and has more advantages than traditional treatment methods, such as chemotherapy, operation, and so on. Just owing to the character of the immune system, immunotherapy arises gradually, and immunotherapy has become the primary treatment instead of other methods in recent decades. Due to the complexity associated with the function of the immune system in the process of tumor proliferation, pathologists and clinicians think that treating malignant tumors is still very hard, such that human beings cannot overcome it even today. It is necessary to construct a mathematical model for understanding the mechanism of the immune system, numerous linear and nonlinear mathematical models endowed with functional response have been considered by mathematicians, which play a crucial role in predicting or controlling tumor growth. It is also necessary to study the mechanisms underlying the interaction between the tumorimmune system to understand immunotherapy. Further development of immunotherapy depends on a better comprehension of the interaction between the tumor system and immune system. Tumor cells can escape the monitoring of the immune system to proliferate infinitely, which is called immune suppression. Many researchers are devoted to removing immune suppression so that cancer cells can be controlled by the immune system. Mathematical models, for example, ordinary differential equation (ODE), partial differential equation (PDE) and partial functional differential equation (PFDE), are extensively used to investigate the dynamics of the immune system. Many researches study how the immune system influences tumor cells by modeling and analyzing mathematical models. Experts attempt to solve the problem based on theoretical analysis and experimental data. There is no doubt that constructing mathematical models provides us with a new view to recognize the tumor-immune system.

In recent years, immunotherapy models have been extensively utilized to predict the behaviors of tumor cells. Ref. [3] constructed a tumor-immune system with two distributed delays to describe the interactions between tumor cells and immune cells. Ref. [4] proposed a tumor-immune model with time delay, and explained that the time delay of the immune reaction could change the stability of equilibrium. Ref. [5] analyzed a two-dimensional tumor-immune system with two delays. By using bifurcation theory, the stability of equilibrium and the existence of Hopf bifurcation were given when the time delays were regarded as the bifurcation parameters. Relevant scholars proposed all kinds of differential equations [6-8]. Refs. [9, 10] showed that different sizes of delays could cause a change in equilibrium stability. Refs. [11-13] studied the various dynamic properties associated with the changing time delay between tumor cells and immune cells. Ref. [14] studied a tumor-immune system in view of optimal control, and showed the method associated with minimizing the cost of immunechemotherapy and reducing the load of tumor cells. Ref. [15] investigated a tumour-immune system with delays and fractional-order, and provided the necessary and sufficient conditions for stability of the steady states and Hopf bifurcating periodic solutions. Ref. [16] presented a delay differential equation to analyze the dynamics between effector cells and tumor cells. The existence and stability of possible steady states and the local stability of Hopf bifurcating periodic solutions are discussed. Das et al. studied a series of studies associated with tumor-immune interaction. For example, Ref. [17] studied the deterministic and stochastic dynamics of tumor-immune interactions. Refs. [18, 19] considered the dynamics of time-delayed tumor-immune systems. Refs. [20, 21] compared the deterministic and noise-induced chaotic dynamics, and described the growth and proliferation process of tumor cells. The analysis showed that the onset of chaos in the system can be predicted. Refs. [22, 23] established the optimal treatment strategies that maximize the number of immune-effector cells, minimized the number of cancer cells, and detrimental effects caused by the amount of drugs.

The motivation of this paper is as follows. First, Ref. [24] showed that there existed a time delay during the immune system recognition of tumor cells. Ref. [25] pointed out that delay played an important role in the interaction between the immune system and tumor cells. The different delays could cause different phenomena of the tumor-immune system, which is useful for understanding the tumor-immune system and providing treatment for tumors. Second, Ref. [26] indicated that tumor cells

could spread in the body and revealed some factors causing the diffusion of tumor cells. Therefore, it is necessary and reasonable to introduce delay and diffusion into the tumor-immune system. Third, Ref. [27] considered some results on the nonlinear dynamics of delayed differential equation models describing the interaction between tumor cells and effector cells of the immune system, however, it did not consider the influence of diffusion of tumor cells and immune cells. For the immune reaction of organisms, it must be enough time to identify nonself cells, and the growth of tumor cells also needs some time. Based on the model of Ref. [27], we introduce two delays and diffusion, and propose a tumor-immune system with two delays and diffusion.

The paper is organized as follows. In Section 2, we establish a partial functional differential equation (PFDE) associated with tumor-immune system, and study the dynamics of this system. In Section 3, the stability of nonnegative equilibrium and the existence of Hopf bifurcation are presented. The normal form of Hopf bifurcation is deduced in Section 4. Numerical simulations to test our findings are shown in Section 5. In Section 6, we give a conclusion for our work.

2. Mathematical modeling

The essential mechanisms of interaction between tumor cells and immune cells is shown as follows (see Figure 1).



Figure 1. Scheme of essential mechanisms of interaction between tumor cells and immune cells.

Immune cells assault tumor cells, and the multiplication of immune cells is stimulated by the appearance of tumor cells. The body can not recognize tumor cells immediately, and there exists a time delay τ_1 for responding to the entrance of nonself cells, and the response time may be short, however, it cannot be ignored. Meanwhile, there also exists a time delay τ_2 when tumor cells are recognized by immune cells. Furthermore, tumor cells and immune cells diffuse in the body under conditions of limited resources and space, and the growth and interaction between tumor cells and immune cells not only depend on time, but are also affected and restricted by the living spatial environment. Hence, it is also indispensable to investigate the complex interaction caused by spatial factors. Considering the

$$\begin{cases} \frac{\partial u(x,t)}{\partial t} = d_1 \Delta u(x,t) + ru(x,t)(1 - \frac{u(x,t-\tau_1)}{\alpha}) - mu(x,t)v(x,t), \quad t > 0, x \in \Omega, \\ \frac{\partial v(x,t)}{\partial t} = d_2 \Delta v(x,t) + \beta u(x,t-\tau_2)v(x,t-\tau_2) - dv(x,t) - pu(x,t)v(x,t), \quad t > 0, x \in \Omega, \\ \frac{\partial u(x,t)}{\partial x} = \frac{\partial v(x,t)}{\partial x} = 0, \quad t \ge 0, \ x \in \partial\Omega, \\ u(x,t) = u_0(x,t) \ge 0, \ v(x,t) = v_0(x,t) \ge 0, (t,x) \in [-\tau,0] \times \bar{\Omega}. \end{cases}$$
(2.1)

where $\Omega = [0, \pi]$ is a bounded open domain in \mathbb{R}^N $(N \ge 1)$ with a smooth boundary $\partial\Omega$, and Δ denotes the Laplacian operator in \mathbb{R}^N $(N \ge 1)$. *u* and *v* are the numbers of tumor cells and immune cells, respectively. The description of the parameters is given in Table 1, and these parameters are all positive. We also assume that $u_0, v_0 \in \mathbb{C} = C([-\tau, 0], X)$ and *X* is defined by $X = \{u, v \in W^{2,2}(\Omega) : \frac{\partial u(x,t)}{\partial x} = \frac{\partial v(x,t)}{\partial x} = 0, x \in \partial\Omega\}$. There exists a time delay for producing a proper immune response in host cells, and considering the time delay of tumor growth τ_1 , the growth of tumor cells follows $ru(1 - \frac{u(x,t-\tau_1)}{\alpha})$. $\beta u(x,t-\tau_2)v(x,t-\tau_2)$ means that immune cells are activated due to the stimulus of tumor cells.

Parameter	description
d_1	Diffusion rate of tumor cells
d_2	Diffusion rate of immune cells
r	Growth rate of malignant cells
α	Environmental carrying capacity
m	Death rate of tumor cells
р	Death rate of immune cells
d	Intrinsic death rate of the immune cells
eta	Activating rate of the immune cells
$ au_1$	The time delay of tumor cells proliferation
$ au_2$	The time of immune cells recognizing tumor cells

Table 1. The description of parameters.

3. Bifurcation Analysis

System (2.1) has two boundary equilibria $E_0 = (0, 0), E_1 = (\alpha, 0)$, and it has a unique positive constant steady equilibrium $E^* = (u^*, v^*) = (\frac{d}{\beta-p}, \frac{r}{m}(1 - \frac{d}{\alpha(\beta-p)}))$ when $d < \alpha(\beta - p)$ and $\beta > p$. For convenience, we show this condition associated with the existence of positive equilibrium E^* by using the following assumption.

$$(H_1): d < \alpha(\beta - p), \beta > p.$$

Denote $U(x, t) = (u(x, t), v(x, t))^{T}$, then linearized system for Eq. (2.1) can be rewritten as a differential equation at equilibrium $E = (u_0, v_0)$ (here $(u_0, v_0) = (0, 0)$, $(\alpha, 0)$, (u^*, v^*)):

$$\frac{\partial U(x,t)}{\partial t} = D\Delta U(x,t) + AU(x,t) + BU(x,t-\tau_1) + CU(x,t-\tau_2)$$

Mathematical Biosciences and Engineering

where

$$D = \begin{pmatrix} d_1 & 0 \\ 0 & d_2 \end{pmatrix}, \quad A = \begin{pmatrix} r - \frac{ru_0}{\alpha} - mv_0 & -mu_0 \\ -pv_0 & -d - pu_0 \end{pmatrix}, \quad B = \begin{pmatrix} -\frac{ru_0}{\alpha} & 0 \\ 0 & 0 \end{pmatrix}, \quad C = \begin{pmatrix} 0 & 0 \\ \beta v_0 & \beta u_0 \end{pmatrix}.$$

Hence, the characteristic equation of (2.1) at (u_0, v_0) is given as follows:

$$\lambda^{2} + A_{n}\lambda + (B_{n}\lambda + C_{n})e^{-\lambda\tau_{1}} + (D_{n}\lambda + E_{n})e^{-\lambda\tau_{2}} + F_{n}e^{-\lambda(\tau_{1}+\tau_{2})} + G_{n} = 0, \ n = 0, 1, 2, \cdots,$$
(3.1)

where

$$\begin{aligned} A_n &= d + d_1 n^2 + d_2 n^2 + p u_0 - r + \frac{r u_0}{\alpha} + m v_0, \ B_n &= \frac{r u_0}{\alpha}, \ C_n &= \frac{r u_0}{\alpha} (d_2 n^2 + d + p u_0), \\ D_n &= -\beta u_0, \ E_n &= m u_0^2 \beta - \beta u_0 (d_1 n^2 - r + \frac{r u_0}{\alpha} + m v_0), \\ F_n &= -\frac{r u_0^2 \beta}{\alpha}, \ G_n &= (d_1 n^2 - r + \frac{r u_0}{\alpha} + m v_0) (d_2 n^2 + d + p u_0) - m p u_0 v_0. \end{aligned}$$

3.1. The case for $\tau_1 = 0$, $\tau_2 = 0$

When $\tau_1 = 0$, $\tau_2 = 0$, Eq. (3.1) for equilibrium $E_0 = (0, 0)$ becomes

$$\lambda^{2} + (d_{1}n^{2} + d_{2}n^{2} + d - r)\lambda + (d_{1}n^{2} - r)(d_{2}n^{2} + d) = 0, \ n = 0, 1, 2, \cdots$$
(3.2)

For Eq. (3.1) with n = 0, the product of two eigenvalues -rd < 0, obviously, equilibrium $E_0 = (0, 0)$ is always unstable.

When $\tau_1 = 0$, $\tau_2 = 0$, Eq. (3.1) for equilibrium $E_1 = (\alpha, 0)$ becomes

$$\lambda^{2} + (d_{1}n^{2} + d_{2}n^{2} + d + p\alpha + r - \beta\alpha)\lambda + (d_{1}n^{2} + r)(d_{2}n^{2} + d + p\alpha - \beta\alpha) + m\alpha^{2}\beta = 0.$$
(3.3)

When (H_1) does not hold,

$$\begin{aligned} &d_1n^2 + d_2n^2 + d + p\alpha + r - \beta\alpha \ge d + p\alpha + r - \beta\alpha > d + p\alpha - \beta\alpha > 0, \\ &(d_1n^2 + r)(d_2n^2 + d + p\alpha - \beta\alpha) + m\alpha^2\beta > 0, \end{aligned}$$

thus, equilibrium $E_1 = (\alpha, 0)$ is always locally asymptotically stable when (H_1) does not hold.

When (H_1) holds, there also exists positive equilibrium E_2 . The stability analysis for $E_1 = (\alpha, 0)$ is similar to $E^* = (u^*, v^*)$, and we only show the general stability results for $E = (u_0, v_0)$ (here $(u_0, v_0) = (\alpha, 0), (u^*, v^*)$). Thus, we show following assumption.

$$(H_2): d + pu_0 + \frac{2ru_0}{\alpha} - r + mv_0 - \beta u_0 > 0.$$

When $\tau_1 = 0$, $\tau_2 = 0$, Eq. (3.1) becomes

$$\lambda^{2} + (A_{n} + B_{n} + D_{n})\lambda + C_{n} + E_{n} + F_{n} + G_{n} = 0, \ n = 0, 1, 2, \cdots,$$

note that under (H_2) , $A_n + B_n + D_n \ge A_0 + B_0 + D_0 = d + pu_0 + \frac{2ru_0}{\alpha} - r + mv_0 - \beta u_0 > 0$. Denote

$$\mathfrak{F}(n) = C_n + E_n + F_n + G_n = \xi_2 n^4 + \xi_1 n^2 + \xi_0, \tag{3.4}$$

Mathematical Biosciences and Engineering

where

$$\begin{split} \xi_2 &= d_1 d_2, \\ \xi_1 &= d_2 (\frac{2ru_0}{\alpha} - r + mv_0) - \beta u_0 d_1 + d_1 (d + pu_0), \\ \xi_0 &= \frac{ru_0}{\alpha} (d + pu_0) + mu_0^2 \beta - \frac{ru_0^2 \beta}{\alpha} - mpu_0 v_0 - (\frac{ru_0}{\alpha} - r + mv_0) \beta u_0. \end{split}$$

Thus, we obtain following theorem.

Theorem 3.1. For system (2.1) with $\tau_1 = 0$, $\tau_2 = 0$, the stability results for equilibria are given as follows.

(1) Equilibrium $E_0 = (0, 0)$ is always unstable.

(2) Equilibrium $E_1 = (\alpha, 0)$ is always locally asymptotically stable when (H_1) does not hold.

(3) When (H_1) and (H_2) hold, the stability results for equilibrium E_1 (or E^*) are given as follows:

(3-1) If $\xi_0 < 0$, equilibrium E_1 (or E^*) is unstable.

(3-2) If $\xi_0 > 0$ and $-\frac{\xi_1}{2\xi_2} < 0$, equilibrium E_1 (or E^*) is locally asymptotically stable. (3-3) If $\xi_0 > 0$, $-\frac{\xi_1}{2\xi_2} > 0$, $\mathfrak{F}(\lfloor -\frac{\xi_1}{2\xi_2} \rfloor) > 0$ and $\mathfrak{F}(\lfloor -\frac{\xi_1}{2\xi_2} \rfloor + 1) > 0$, equilibrium E_1 (or E^*) is locally asymptotically stable

 $(3-4) If \xi_0 > 0, -\frac{\xi_1}{2\xi_2} > 0, and \mathfrak{F}(\lfloor -\frac{\xi_1}{2\xi_2} \rfloor) < 0 \text{ or } \mathfrak{F}(\lfloor -\frac{\xi_1}{2\xi_2} \rfloor + 1) < 0, equilibrium E_1 (or E^*) is unstable.$

3.2. The case for $\tau_2 = 0, \tau_1 \neq 0$

The stability analysis for $E_1 = (\alpha, 0)$ is similar to $E^* = (u^*, v^*)$, thus, we only show the general stability results for $E = (u_0, v_0)$ (here $(u_0, v_0) = (\alpha, 0), (u^*, v^*)$).

When $\tau_2 = 0$, $\tau_1 \neq 0$, the characteristic equation of system (2.1) is given as follows:

$$\lambda^{2} + L_{1}\lambda + (L_{3}\lambda + L_{4})e^{-\lambda\tau_{1}} + L_{2} = 0, \qquad n = 0, 1, 2, \cdots,$$
(3.5)

where $L_1 = A_n + D_n$, $L_2 = E_n + G_n$, $L_3 = B_n$, $L_4 = C_n + F_n$, with

$$A_{n} = d + d_{1}n^{2} + d_{2}n^{2} + pu_{0} - r + \frac{ru_{0}}{\alpha} + mv_{0}, \ B_{n} = \frac{ru_{0}}{\alpha}, \ C_{n} = \frac{ru_{0}}{\alpha}(d_{2}n^{2} + d + pu_{0}),$$
$$D_{n} = -\beta u_{0}, \ E_{n} = mu_{0}^{2}\beta - \beta u_{0}(d_{1}n^{2} - r + \frac{ru_{0}}{\alpha} + mv_{0}),$$
$$F_{n} = -\frac{ru_{0}^{2}\beta}{\alpha}, \ G_{n} = (d_{1}n^{2} - r + \frac{ru_{0}}{\alpha} + mv_{0})(d_{2}n^{2} + d + pu_{0}) - mpu_{0}v_{0}.$$

We might suppose that $\pm i\omega$ ($\omega > 0$) are a pair of purely imaginary roots of Eq. (3.5). Substituting them into Eq. (3.5) and separating the real and imaginary parts, we obtain

$$\begin{cases} \omega^2 - L_2 = L_3 \omega \sin(\omega \tau_1) + L_4 \cos(\omega \tau_1), \\ - L_1 \omega = L_3 \omega \cos(\omega \tau_1) - L_4 \sin(\omega \tau_1), \end{cases}$$

thus, $R := \sin(\omega \tau_1) = \frac{(\omega^2 - L_2)L_3 \omega + L_1 L_4 \omega}{L_3^2 \omega^2 + L_4^2}$, $Q := \cos(\omega \tau_1) = \frac{(\omega^2 - L_2)L_4 - L_1 L_3 \omega^2}{L_3^2 \omega^2 + L_4^2}$, which implies that

$$\omega^4 - (L_3^2 + 2L_2 - L_1^2)\omega^2 + L_2^2 - L_4^2 = 0.$$

Mathematical Biosciences and Engineering

Let $z = \omega^2$, then the above equation can be rewritten in the following form

$$h(z) = z^{2} - (L_{3}^{2} + 2L_{2} - L_{1}^{2})z + L_{2}^{2} - L_{4}^{2}.$$
(3.6)

Under assumption

$$(H_3): L_3^2 - L_1^2 + 2L_2 < 0, \quad L_2^2 - L_4^2 > 0,$$

Eq. (3.6) has no positive roots.

Under assumption

$$(H_4): L_2^2 - L_4^2 < 0,$$

Eq. (3.6) has one unique positive root ω_0 .

Under assumption

$$(H_5): L_3^2 - L_1^2 + 2L_2 > 0, \quad L_2^2 - L_4^2 > 0, \quad (L_3^2 - L_1^2 + 2L_2)^2 - 4(L_2^2 - L_4^2)^2 > 0,$$

Eq. (3.6) has two positive roots $\omega_{1,2}$, and the critical time delay can be determined as,

$$\tau_{1l}^{(j)} = \begin{cases} \frac{1}{\omega_l} [\arccos(Q) + 2j\pi], & R \ge 0, \\ \frac{1}{\omega_l} [2\pi - \arccos(Q) + 2j\pi], & R < 0, \end{cases}$$
(3.7)

where $R := \sin(\omega \tau_1) = \frac{(\omega^2 - L_2)L_3\omega + L_1L_4\omega}{L_3^2\omega^2 + L_4^2}$, $Q := \cos(\omega \tau_1) = \frac{(\omega^2 - L_2)L_4 - L_1L_3\omega^2}{L_3^2\omega^2 + L_4^2}$, $l = 0, 1, 2; j = 0, 1, 2, \cdots$. Then the following transversality conditions yield,

$$\operatorname{Re}\left[\frac{\mathrm{d}(\lambda)}{\mathrm{d}\tau_{1}}\right]\Big|_{\tau_{1}=\tau_{1l}^{(j)}}^{-1} = \operatorname{Re}\left(\frac{(2\lambda+L_{1})\mathrm{e}^{\lambda\tau_{1}}}{\lambda(L_{3}\lambda+L_{4})} + \frac{L_{3}}{\lambda(L_{3}\lambda+L_{4})}\right) = \frac{z}{\Delta}h'(z),$$

where $\Delta = L_2^2 z + L_3^2 z^2$, $l = 0, 1, 2; j = 0, 1, 2, \cdots$. Obviously, under (H_1) and (H_4) , h(z) = 0 has one positive real root z_0 and $h'(z_0) > 0$; under (H_1) and (H_5) , h(z) = 0 has two positive real roots z_1 and z_2 . Suppose $z_1 < z_2$, then $h'(z_1) < 0$ and $h'(z_2) > 0$.

Theorem 3.2. For system (2.1) with $\tau_2 = 0$, $\tau_1 > 0$, the following conclusions hold when assumptions (H_1), (H_2) and the stability conditions of Theorem 3.1 (3-2) or (3-3) hold,

(1) If (H_3) is satisfied, then equilibrium E_1 (or E^*) is locally asymptotically stable for all $\tau_1 > 0$. (2) If (H_4) is satisfied, then equilibrium equilibrium E_1 (or E^*) is locally asymptotically stable for all $0 < \tau_1 < \tau_{10}^{(0)}$, and unstable for $\tau_1 > \tau_{10}^{(0)}$, furthermore, system (2.1) undergoes a Hopf bifurcation at equilibrium E_1 (or E^*) when $\tau_1 = \tau_{10}^{(j)}$, $j = 0, 1, 2, \cdots$.

(3) If (H₅) is satisfied, then there is a positive integer m such that the equilibrium E_1 (or E^*) is locally asymptotically stable when $\tau_1 \in [0, \tau_{12}^{(0)}) \bigcup \bigcup_{k=0}^{(m-1)} (\tau_{11}^{(k)}, \tau_{12}^{(k+1)})$, and unstable $\tau_1 \in \bigcup_{k=0}^{(m-1)} (\tau_{12}^{(k)}, \tau_{11}^{(k)}) \bigcup (\tau_{12}^{(m)}, +\infty)$, furthermore, system (2.1) undergoes a Hopf bifurcation at equilibrium E_1 (or E^*) when $\tau_1 = \tau_{11}^{(j)}$, $l = 1, 2; j = 0, 1, 2, \cdots$.

Mathematical Biosciences and Engineering

3.3. The case for $\tau_2 \neq 0, \tau_1 \neq 0$

The first stable interval mentioned in Theorem 3.2, on which the equilibrium E_1 (or E^*) is locally asymptotically stable, is denoted by stable region *I*. That is, assumptions (H_1) , (H_2) and the stability conditions of Theorem 3.1 (3-2) or (3-3) hold, and if (H_3) also holds, $I = \{\tau_1 | \tau_1 > 0\}$. If (H_4) also holds, $I = \{\tau_1 | 0 < \tau_1 < \tau_{10}^{(0)}\}$. If (H_5) holds, $I = \{\tau_1 | 0 < \tau_1 < \tau_{12}^{(0)}\}$. Next, we choose $\tau_1 = \tau_1^* \in I$. Regarding τ_2 as a parameter, the characteristic equation of (2.1) at (u_0, v_0) (here $(u_0, v_0) = (\alpha, 0), (u^*, v^*)$) is rewritten as follows:

$$\lambda^{2} + (A_{n} + B_{n} e^{-\lambda \tau_{1}^{*}})\lambda + (D_{n}\lambda + E_{n})e^{-\lambda \tau_{2}} + F_{n} e^{-\lambda \tau_{1}^{*}} + C_{n} e^{-\lambda \tau_{1}^{*}} + G_{n} = 0,$$

where

$$A_{n} = d + d_{1}n^{2} + d_{2}n^{2} + pu_{0} - r + \frac{ru_{0}}{\alpha} + mv_{0}, B_{n} = \frac{ru_{0}}{\alpha}, C_{n} = \frac{ru_{0}}{\alpha}(d_{2}n^{2} + d + pu_{0})$$
$$D_{n} = -\beta u_{0}, E_{n} = mu_{0}^{2}\beta - \beta u_{0}(d_{1}n^{2} - r + \frac{ru_{0}}{\alpha} + mv_{0}),$$
$$F_{n} = -\frac{ru_{0}^{2}\beta}{\alpha}, G_{n} = (d_{1}n^{2} - r + \frac{ru_{0}}{\alpha} + mv_{0})(d_{2}n^{2} + d + pu_{0}) - mpu_{0}v_{0}.$$

Letting $\lambda = i\omega(\tau_2)(\omega > 0)$ be the root of the above equation, then separate the real and imaginary parts for above equation,

$$\omega^2 - G_n - C_n \cos(\omega \tau_1^*) - B_n \omega \sin(\omega \tau_1^*) = E_n \cos(\omega \tau_2) + D_n \omega \sin(\omega \tau_2) + F_n \cos(\omega (\tau_1^* + \tau_2)),$$

$$A_n \omega + C_n \sin(\omega \tau_1^*) + B_n \omega \cos(\omega \tau_1^*) = -D_n \omega \cos(\omega \tau_2) + E_n \sin(\omega \tau_2) + F_n \sin(\omega (\tau_1^* + \tau_2)),$$

which leads to,

$$F(\omega) = \omega^4 + (B_n^2 - 2G_n + A_n^2 - D_n^2)\omega^2 + 2(B_nG_n\omega - A_nC_n\omega - B_n\omega^3 + D_nF_n\omega)\sin(\omega\tau_1^*) + G_n^2 + C_n^2 - E_n^2 - F_n^2 + 2(C_nG_n + A_nB_n\omega^2 - C_n\omega^2 - E_nF_n)\cos(\omega\tau_1^*) = 0.$$

Suppose

$$(H_6): (G_n + C_n)^2 - (E_n + F_n)^2 < 0,$$

then, we get $F(0) = G_n^2 + C_n^2 - E_n^2 - F_n^2 + 2C_nG_n - 2E_nF_n < 0$, $F(+\infty) > 0$. Hence, $F(\omega) = 0$ has definite positive roots ω_{2i} , i = 0, 1, 2. For every fixed ω_i , there is a sequence of $\tau_{2i}^{(j)}$, $j = 0, 1, 2, \cdots$, defined by

$$\tau_{2i}^{(j)} = \begin{cases} \frac{1}{\omega_{2i}} \arccos[Q+2j\pi], \ P \ge 0, \\ \frac{1}{\omega_{2i}} \arccos[Q+2j\pi], \ P < 0, \end{cases}$$
(3.8)

where $j = 0, 1, 2, \cdots$,

$$Q = \cos(\omega\tau_2) = \frac{f(E_n + F_n \cos(\omega\tau_1^*)) - g(D_n\omega - F_n \sin(\omega\tau_1^*))}{(E_n + F_n \cos(\omega\tau_1^*))^2 + (D_n\omega - F_n \sin(\omega\tau_1^*))^2},$$

$$P = \sin(\omega\tau_2) = \frac{g(E_n + F_n \cos(\omega\tau_1^*)) + f(D_n\omega - F_n \sin(\omega\tau_1^*))}{(E_n + F_n \cos(\omega\tau_1^*))^2 + (D_n\omega - F_n \sin(\omega\tau_1^*))^2},$$

Mathematical Biosciences and Engineering

with

$$f = \omega^2 - B_n \omega \sin(\omega \tau_1^*) - C_n \cos(\omega \tau_1^*) - G_n, \ g = A_n \omega + B_n \cos(\omega \tau_1^*) \omega + C_n \sin(\omega \tau_1^*).$$

Let $\tau_{20} = \min \tau_{2i}^{(j)}$, $i = 0, 1, 2, j = 0, 1, 2, \cdots$, when $\tau_2 = \tau_{20}$, Eq. (3.1) has a pair of purely imaginary roots $\pm i\omega_{20}$ for $\tau_1^* \in I$. Assume

$$(H_7): \operatorname{Re}\left[\frac{\mathrm{d}\lambda}{\mathrm{d}\tau_2}\right]_{\tau_2=\tau_{20}^{(j)}} \neq 0,$$

then we have,

Theorem 3.3. Assuming that (H_1) , (H_6) and (H_7) are satisfied, and one of the assumptions $(H_3), (H_4)$ and (H_5) holds, then for $\tau_1^* \in I$, system (2.1) undergoes a Hopf bifurcation at $E = (u_0, v_0)$ when $\tau_2 = \tau_{20}^{(j)}$, here $(u_0, v_0) = (\alpha, 0)$ or (u^*, v^*) with $u^* = \frac{d}{\beta - p}$, $v^* = \frac{r}{m}(1 - \frac{d}{\alpha(\beta - p)})$, and the equilibrium E is locally asymptotically stable when $\tau_2 \in [0, \tau_{20})$.

4. Normal formal of Hopf bifurcation

In this section, we derive the normal formal of Hopf bifurcation at $E^* = (u^*, v^*)$ when $\tau_2 = 0$, $\tau_1 \neq 0$ for the system (2.1). We denote the critical value $\tau = \tau_2 = \tau_c$. Define $\overline{u}(x, t) = u(x, \tau t) - u^*$, $\overline{v}(x, t) = v(x, \tau t) - v^*$, and drop the bar for convenience, the system (2.1) can be rewritten as,

$$\begin{cases} \frac{\partial u(x,t)}{\partial t} = \tau [d_1 \triangle u(x,t) - \frac{ru(x,t-1)}{\alpha} (u^* + u(x,t)) - \frac{r}{\alpha} u^* u(x,t) + ru(x,t) \\ - m(u(x,t)v(x,t) + u^*v(x,t) + v^*u(x,t))], \quad t > 0, x \in \Omega, \\ \frac{\partial v(x,t)}{\partial t} = \tau [d_2 \triangle v(x,t) + (\beta - p)(u(x,t)v(x,t) + u(x,t)v^* + u^*v(x,t)) - dv(x,t)], \quad t > 0, x \in \Omega, \\ \frac{\partial u(x,t)}{\partial x} = \frac{\partial v(x,t)}{\partial x} = 0, \quad t \ge 0, \ x \in \partial\Omega, \\ u(x,t) = u_0(x,t) \ge 0, \ v(x,t) = v_0(x,t) \ge 0, (t,x) \in [-\tau,0] \times \bar{\Omega}. \end{cases}$$

$$(4.1)$$

Let $h = (h_{11}, h_{12})^T$ be the eigenvector of the linear operator corresponding to the eigenvalue $i\omega\tau$, and let $h^* = (h_{21}, h_{22})^T$ be the normalized eigenvector of the adjoint operator of the linear operator corresponding to the eigenvalues $-i\omega\tau$ satisfying the inner product $\langle \overline{h^*}, h \rangle = \overline{h^*}^T \cdot h = 1$. By a simple calculation, we get

$$h = (h_{11}, h_{12})^{T} = (1, \frac{(\beta - p)v^{*}}{i\omega + d_{2}n^{2} + d + pu^{*} - \beta u^{*}})^{T},$$

$$h^{*} = (h_{21}, h_{22})^{T} = l(\frac{i\omega - d_{2}n^{2} - d - pu^{*} + \beta u^{*}}{mu^{*}}, 1)^{T},$$

$$l = [\frac{i\omega - d_{2}n^{2} - d - pu^{*} + \beta u^{*}}{mu^{*}} + \frac{(\beta - p)v^{*}}{-i\omega + d_{2}n^{2} + d + pu^{*} - \beta u^{*}}]^{-1}.$$
(4.2)

Suppose the solution of Eq. (4.1) is

$$U(x,t) = U(x,T_0,T_1,T_2,\cdots) = \sum_{k=1}^{+\infty} \varepsilon^k U_k(x,T_0,T_1,T_2\cdots),$$
(4.3)

Mathematical Biosciences and Engineering

where

$$U(x, T_0, T_1, T_2, \cdots) = (u(x, T_0, T_1, T_2, \cdots), v(x, T_0, T_1, T_2, \cdots))^T,$$

$$U_k(x, T_0, T_1, T_2, \cdots) = (u_k(x, T_0, T_1, T_2, \cdots), v_k(x, T_0, T_1, T_2, \cdots))^T$$

The derivation with respect to *t* is

$$\frac{\mathrm{d}}{\mathrm{d}t} = \frac{\partial}{\partial T_0} + \varepsilon \frac{\partial}{\partial T_1} + \varepsilon^2 \frac{\partial}{\partial T_2} + \dots = D_0 + \varepsilon D_1 + \varepsilon^2 D_2 + \dots,$$

where the differential operator $D_i = \frac{\partial}{\partial T_i}$, $i = 0, 1, 2, \cdots$.

Denote

$$u_j = u_j(x, T_0, T_1, T_2, \cdots), \ u_{j,1} = u_j(x, T_0 - 1, T_1, T_2, \cdots),$$
$$v_j = v_j(x, T_0, T_1, T_2, \cdots), \ v_{j,1} = v_j(x, T_0 - 1, T_1, T_2, \cdots), \quad j = 1, 2, 3, \cdots.$$

We obtain

$$\frac{dU(x,t)}{dt} = \varepsilon D_0 U_1 + \varepsilon^2 D_0 U_2 + \varepsilon^2 D_1 U_1 + \varepsilon^3 D_0 U_3 + \varepsilon^3 D_1 U_2 + \varepsilon^3 D_2 U_1 + \cdots,$$

$$\Delta U(x,t) = \varepsilon \Delta U_1(x,t) + \varepsilon^2 \Delta U_2(x,t) + \varepsilon^3 \Delta U_3(x,t) + \cdots.$$
(4.4)

We take perturbations as $\tau = \tau_c + \varepsilon \mu$, to deal with the delayed terms, we expend u(x, t-1), v(x, t-1) at $u(x, T_0 - 1, T_1, T_2, \cdots)$ and $v(x, T_0 - 1, T_1, T_2, \cdots)$, respectively, that is,

$$u(x,t-1) = \varepsilon u_{1,1} + \varepsilon^2 u_{2,1} - \varepsilon^2 D_1 u_{1,1} + \varepsilon^3 u_{3,1} - \varepsilon^3 D_1 u_{2,1} - \varepsilon^3 D_2 u_{1,1} + \cdots,$$

$$v(x,t-1) = \varepsilon v_{1,1} + \varepsilon^2 v_{2,1} - \varepsilon^2 D_1 v_{1,1} + \varepsilon^3 v_{3,1} - \varepsilon^3 D_1 v_{2,1} - \varepsilon^3 D_2 v_{1,1} + \cdots,$$
(4.5)

where $u_{j,1} = u_j(x, T_0 - 1, T_1, T_2, \cdots)$, $v_{j,1} = v_j(x, T_0 - 1, T_1, T_2, \cdots)$, $j = 1, 2, 3, \cdots$. Substituting Eqs. (4.3)-(4.5) into Eq. (4.1), for the ε -order terms, we obtain

$$\begin{cases} D_0 u_1 - \tau_c d_1 \Delta u_1 + \frac{r u^*}{\alpha} \tau_c u_{1,1} + \frac{r u^*}{\alpha} \tau_c u_1 + r \tau_c u_1 + m \tau_c (u^* v_1 + v^* u_1) = 0, \\ D_0 v_1 - \tau_c d_2 \Delta v_1 - \tau_c (\beta - p) (v^* u_1 + u^* v_1) + d \tau_c v_1 = 0. \end{cases}$$
(4.6)

Since $\pm i\omega\tau$ are the eigenvalues of the linear part of Eq. (4.1), the solution of Eq. (4.6) can be expressed in the following form

$$U_1(x, T_0, T_1, T_2, \cdots) = G(T_1, T_2, \cdots) e^{i\omega\tau_c T_0} h\cos(nx) + c.c.,$$
(4.7)

where c.c. means the complex conjugate of the preceding terms, and h is given in Eq. (4.2).

For the ε^2 -order terms, we obtain

$$\begin{cases} D_{0}u_{2} - \tau_{c}d_{1}\Delta u_{2} + \frac{ru^{*}}{\alpha}\tau_{c}u_{2,1} + \frac{ru^{*}}{\alpha}\tau_{c}u_{2} + r\tau_{c}u_{2} + m\tau_{c}(u^{*}v_{2} + v^{*}u_{2}) \\ = -D_{1}u_{1} + \mu d_{1}\Delta u_{1} + \frac{r}{\alpha}(\tau_{c}u^{*}D_{1}u_{1,1} - \tau_{c}u_{1}u_{1,1} - \mu u^{*}u_{1,1} - \mu u^{*}u_{1}) - r\mu u_{1} - m(\tau_{c}u_{1}v_{1} + \mu u^{*}v_{1} + \mu v^{*}u_{1}), \\ D_{0}v_{2} - \tau_{c}d_{2}\Delta v_{2} - \tau_{c}(\beta - p)(v^{*}u_{2} + u^{*}v_{2}) + d\tau_{c}v_{2} \\ = -D_{1}v_{1} + \mu d_{2}\Delta v_{1} + (\beta - p)\mu(u^{*}v_{1} + u_{1}v^{*}) + \tau_{c}(\beta - p)u_{1}v_{1} - d\mu v_{1}. \end{cases}$$

$$(4.8)$$

Mathematical Biosciences and Engineering

Substituting Eq. (4.7) into the right side of Eq. (4.8), we obtain the coefficient vector of term $e^{i\omega\tau_c T_0}$, denoted as m_1 , by solvability conditions, that is, $\langle h^*, m_1 \rangle = 0$, we obtain

$$\frac{\partial G}{\partial T_1} = M\mu G,\tag{4.9}$$

where

$$M = \frac{\alpha \overline{h}_{21} h_{11} (v^* - n^2 d_1 - u^* e^{-i\omega\tau_c} - u^* - r) - m\alpha u^* \overline{h}_{21} h_{12} - \alpha \overline{h}_{22} [(\beta - p)(u^* h_{12} + v^* h_{11}) - h_{12} (d_2 n^2 + d)]}{\alpha (h_{11} \overline{h}_{21} + h_{12} \overline{h}_{22}) - ru^* \tau_c h_{11} \overline{h}_{21} e^{-i\omega\tau_c}}$$

Suppose the solution of Eq. (4.8) is

$$\begin{cases} u_2 = \sum_{k=0}^{+\infty} (\eta_{0k} G \overline{G} + \eta_{1k} G^2 e^{2i\omega\tau_c T_0} + \overline{\eta}_{1k} \overline{G}^2 e^{-2i\omega\tau_c T_0}) \cos(kx), \\ v_2 = \sum_{k=0}^{+\infty} (\zeta_{0k} G \overline{G} + \zeta_{1k} G^2 e^{2i\omega\tau_c T_0} + \overline{\zeta}_{1k} \overline{G}^2 e^{-2i\omega\tau_c T_0}) \cos(kx). \end{cases}$$
(4.10)

Denoted $c_k = \langle \cos(nx) \cos(nx), \cos(kx) \rangle = \int_0^{\pi} \cos(nx) \cos(nx) \cos(kx) dx$, thus

$$\begin{aligned} \eta_{1k} &= -\frac{c_k}{F_{1,k}} [(r e^{-i\omega\tau_c} h_{11}^2 + \alpha m h_{11} h_{12}) (d_2 k^2 + 2i\omega - \beta u^* + pu^* + d) + m\alpha(\beta - p) h_{11} h_{12}], \\ \zeta_{1k} &= \frac{c_k}{F_{1,k}} [(2i\alpha\omega + d_1 k^2 \alpha + ru^* e^{-2i\omega\tau_c} + ru^* + r\alpha + mv^*) (\beta - p) h_{11} h_{12} - (\beta - p) v^* (r e^{-i\omega\tau_c} h_{11}^2 + m\alpha h_{11} h_{12})], \\ \eta_{0k} &= -\frac{c_k}{F_{2,k}} \{ [1 + u^* (\beta - p)] \alpha m (h_{11} \overline{h_{12}} + h_{12} \overline{h_{11}}) + (d_2 k^2 - \beta u^* + pu^* + d) r h_{11} \overline{h_{11}} (e^{i\omega\tau_c} + e^{i\omega\tau_c}) \}, \\ \zeta_{0k} &= -\frac{c_k}{F_{2,k}} \{ (\beta - p) v^* r h_{11} \overline{h_{12}} (e^{i\omega\tau_c} + e^{i\omega\tau_c}) + (h_{11} \overline{h_{12}} + h_{12} \overline{h_{11}}) [m\alpha - (\beta - p) (d_1 k^2 \alpha + ru^* + r\alpha + v^* m\alpha)] \}, \end{aligned}$$

$$(4.11)$$

with

$$F_{1,k} = [(2i\alpha\omega + d_1k^2\alpha + ru^*e^{-2i\omega\tau_c} + ru^* + r\alpha + m\alpha v^*)(2i\omega + d_2k^2 - \beta u^* + pu^* + d) + \alpha(\beta - p)v^*m] \times \int_0^{\pi} \cos(kx)\cos(kx)dx,$$

$$F_{2,k} = (\int_0^{\pi} \cos(kx)\cos(kx)dx)[mu^*(\beta - p)v^*\alpha + (d_2k^2 - \beta u^* + pu^* + d)(d_1k^2\alpha + ru^* + r\alpha + v^*m\alpha)].$$

For the ε^3 -order terms, we obtain

$$\begin{cases} D_{0}u_{3} - \tau_{c}d_{1}\Delta u_{3} + \frac{ru^{*}}{\alpha}\tau_{c}u_{3,1} + \frac{ru^{*}}{\alpha}\tau_{c}u_{3} + r\tau_{c}u_{3} + m\tau_{c}(u^{*}v_{3} + v^{*}u_{3}) \\ = -D_{1}u_{2} - D_{2}u_{1} + \mu d_{1}\Delta u_{2} + \frac{r}{\alpha}\tau_{c}(u^{*}D_{1}u_{2,1} + u^{*}D_{2}u_{1,1} - u_{1}u_{2,1} + u_{1}D_{1}u_{1,1} - u_{2}u_{1,1}) \\ - \frac{r}{\alpha}\mu(u^{*}u_{2,1} - u^{*}D_{1}u_{1,1} + u_{1}u_{1,1}) - \frac{r}{\alpha}\mu u^{*}u_{2} - r\mu u_{2} - m\tau_{c}(u_{2}v_{1} + u_{1}v_{2}) - m\mu(u_{1}v_{1} + u^{*}v_{2} + v^{*}u_{2}), \\ D_{0}v_{3} - \tau_{c}d_{2}\Delta v_{3} - \tau_{c}(\beta - p)(v^{*}u_{3} + u^{*}v_{3}) + d\tau_{c}v_{3} \\ = -D_{1}v_{2} - D_{2}v_{1} + \mu d_{2}\Delta v_{2} + (\beta - p)\tau_{c}(u_{2}v_{1} + u_{1}v_{2}) + \mu(\beta - p)(u_{1}v_{1} + v^{*}u_{2} + u^{*}v_{2}) - d\mu v_{2}. \end{cases}$$

$$(4.12)$$

Mathematical Biosciences and Engineering

Substituting solutions Eq. (4.7) and Eq. (4.10) into the right side of Eq. (4.12), we obtain the coefficient vector of term $e^{i\omega\tau_c T_0}$, denoted as m_2 , by solvability conditions, let $< h^*$, $m_2 >= 0$, we obtain

$$\frac{\partial G}{\partial T_2} = \chi G^2 \overline{G},\tag{4.13}$$

where

$$\chi = \frac{1}{F_3} \left[-r\tau_c \overline{h_{21}} \sum_{k \ge 0} (c_k \eta_{0k} h_{11} + \overline{h_{11}} \eta_{0k} e^{-2i\omega\tau_c} + \eta_{0k} h_{11} e^{-i\omega\tau_c} + \eta_{1k} \overline{h_{11}} e^{i\omega\tau_c}) + \overline{h_{22}} (\beta - p) \tau_c \alpha \sum_{k \ge 0} c_k (\eta_{0k} h_{12} + \eta_{1k} \overline{h_{12}} + \zeta_{0k} h_{11} + \overline{h_{11}} \zeta_{1k}) \right],$$

with

$$F_3 = \left[\overline{h_{21}}(\alpha h_{11} - ru^*\tau_c h_{11}e^{-i\omega\tau_c}) + \alpha \overline{h_{22}}h_{12}\right] \int_0^\pi \cos(nx)\cos(nx)dx.$$

According to the above analysis, the normal form of Hopf bifurcation for system (2.1) reduced on the center manifold is

$$\frac{\partial G}{\partial T} = \varepsilon \frac{\partial G}{\partial T_1} + \varepsilon^2 \frac{\partial G}{\partial T_2} + \cdots, \qquad (4.14)$$

making $G \to G/\varepsilon$, thus, Eq. (4.14) becomes:

$$\dot{G} = M\mu G + \chi G^2 \overline{G}, \tag{4.15}$$

where *M* and χ are given by Eq. (4.9) and Eq. (4.13), respectively.

Let $G = re^{i\theta}$ and substitute it into Eq. (4.15), and we obtain the Hopf bifurcation normal form in polar coordinates:

$$\begin{cases} \dot{r} = \operatorname{Re}(M)\mu r + \operatorname{Re}(\chi)r^{3}, \\ \dot{\theta} = \operatorname{Im}(M)\mu + \operatorname{Im}(\chi)r^{2}. \end{cases}$$
(4.16)

According to the bifurcation normal form Eq. (4.16) in polar coordinates, we have following theorem.

Theorem 4.1. For system (4.16), if $\frac{\operatorname{Re}(M)\mu}{\operatorname{Re}(\chi)} < 0$ holds, system (2.1) exists periodic solutions near equilibrium $E = (u_0, v_0)$, here $(u_0, v_0) = (\alpha, 0)$ or (u^*, v^*) with $u^* = \frac{d}{\beta - p}$, $v^* = \frac{r}{m}(1 - \frac{d}{\alpha(\beta - p)})$.

(1) If $\operatorname{Re}(M)\mu < 0$, the bifurcating periodic solutions are unstable.

(2) If $\operatorname{Re}(M)\mu > 0$, the bifurcating periodic solutions are locally asymptotically stable.

5. Numerical simulations

According to Ref. [12] we choose the following parameter (see Table 2) for numerical simulation. Obviously, (H_1) holds, and system (2.1) has two boundary equilibria $E_0 = (0,0)$ and $E_1 = (\alpha,0) = (5 \times 10^6, 0)$, and one nontrivial equilibrium $E^* = (u^*, v^*) = (2.5948 \times 10^5, 7.8643 \times 10^5)$. Actually, we only care about the stability of the nontrivial equilibrium E^* . In this section, we plot numerical simulations by using MATLAB software.

When $\tau_1 = 0$, $\tau_2 = 0$, according to Theorem 3.1, $E_0 = (0,0)$ is always unstable. This means that if system (2.1) without delays, the immune system cannot restrain the growth of tumor cells, and tumor cells proliferate quickly in the body. For equilibrium E^* , assumption (H_2) and the conditions

Table 2. The value of parameters.		
Parameter	Value	
d_1	$0.02 \ day^{-1}$	
d_2	$0.2 \ day^{-1}$	
r	$0.18 \ day^{-1}$	
α	5.0×10^6 viable cells ⁻¹	
т	$1.101 \times 10^{-7} day^{-1}$ viable cells ⁻¹	
р	$3.422 \times 10^{-10} day^{-1}$ viable cells ⁻¹	
d	$0.00152 \ day^{-1}$	
eta	$6.2 \times 10^{-9} day^{-1}$ viable cells ⁻¹	

Table 2. The value of parameters.

in Theorem 3.1 (3-2) hold, thus equilibrium $E^* = (u^*, v^*)$ is locally asymptotically stable. We choose the initial value (2.5947 × 10⁶, 7.8642 × 10⁵), $E^* = (u^*, v^*)$ is locally asymptotically stable (see Figure 2). Although tumor cells and immune cells can coexist at this moment, the immune system can suppress the growth of tumor cells effectively.



Figure 2. Simulated solution of system (2.1) for $\tau_1 = \tau_2 = 0$, showing a locally asymptotically stable equilibrium $E^* = (u^*, v^*)$.

By a simple calculation, (H_5) holds, from Eq. (3.7), we obtain $\tau_{12}^{(0)} = 14.8403$, we choose $\tau_1 = 3 \in (0, \tau_{12}^{(0)})$ and choose the initial value $(2.5947 \times 10^6, 7.8642 \times 10^5)$, according to Theorem 3.2, $E^* = (u^*, v^*)$ is locally asymptotically stable (see Figure 3). From Figure 3, we conclude that if tumor cells need a long time to proliferate, the tumor cells cannot diffuse, and it is easy to cure the tumor at this time. Thus, the diffusion of tumor cells can be controlled by making the size of the delay small. We reduce the diffusion of tumor cells by controlling the time of the proliferation of tumor cells.



Figure 3. Simulated solution of system (2.1) for $\tau_1 = 3$, $\tau_2 = 0$, showing a locally asymptotically stable equilibrium $E^* = (u^*, v^*)$.

1166

We choose $\tau_1 = 15$ and the initial value $(2.5947 \times 10^6, 7.8642 \times 10^5)$, from (4.15), we obtain $\operatorname{Re}(M) > 0$, $\operatorname{Re}(\chi) < 0$, thus, according to Theorem 4.1, system (2.1) exhibits stable periodic solutions near E^* , and the direction of Hopf bifurcation is forward. This means that, when τ is close to τ_c , the number of tumor cells can be controlled, and the number of tumor cells and immune cells varies periodically (see Figure 4).



Figure 4. Simulated solution of system (2.1) for $\tau_1 = 15$, $\tau_2 = 0$, showing a stable periodic solution near equilibrium $E^* = (u^*, v^*)$.

When $\tau_2 \neq 0, \tau_1 \neq 0$, the stable interval mentioned in Theorem 3.2 $I = {\tau_1 | 0 < \tau_1 < \tau_{12}^{(0)} = 14.8403}$, on which the equilibrium E^* is locally asymptotically stable for $\tau_2 = 0$, and (H_2) holds, from (3.8), we obtain $\tau_{21}^{(0)} = 4.3239$. We choose $\tau_1 = 1 \in I$, $\tau_2 = 1$, according to Theorem 3.3, we choose the initial value $(2.5947 \times 10^6, 7.8642 \times 10^5)$, $E^* = (u^*, v^*)$ is locally asymptotically stable (see Figure 5). The time of tumor growth is fixed within a certain range, and the tumor cells cannot escape the control of the immune system.



Figure 5. Simulated solution of system (2.1) for $\tau_1 = 1$, $\tau_2 = 1$, showing a locally asymptotically stable equilibrium $E^* = (u^*, v^*)$.

Remark: Comparing the above figures, we have the following results. When the time delay is less than the critical value, the equilibrium is locally asymptotically stable, that is, the immune system can restrain the growth of tumor cells effectively at this moment, and tumor cells may not proliferate insanely in the body. Tumor cells originate from the malignant growth of normal cells. It is shown that the immune system can restrain tumor growth, although immune cells and tumor cells could coexist, and the immune system has higher efficiency (see Figures 1, 2 and 5). As the time delay of immune cells identifying tumor cells increased, immune system is able to identify itself from nonself substances, and immunotherapy can work effectively. Then, the numbers of tumor cells and immune cells identifying tumor cells of signer 3). When the time delay of immune cells identifying tumor cells of tumor cells are larger, that is, the immune system loses

function, and the body may be damaged. On the other hand, the growing speed of tumor cells depends on the mood and diet of the patient, and it is very important to maintain an optimistic mood and a regular diet.

6. Conclusion

We constructed a tumor-immune system with two delays and diffusion, and investigated how the delays affect the dynamics of the system. We analyzed the existence and stability of equilibria, and studied the dynamic properties of Hopf bifurcation. The nontrivial equilibrium is locally asymptotically stable under suitable parameters for the system with two delays, and the system occurred stable periodical solutions when the delay of tumor cell proliferation passed through the critical value.

Conflict of interest

The authors declare that they have no competing interests.

Funding

This study was funded by Fundamental Research Funds for the Central Universities of China (Grant No. 2572021DJ01) and the Heilongjiang Provincial Natural Science Foundation of China (Grant No. LH2019A001).

Availability of data and materials

The datasets used or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

The idea of this research was introduced by Yuting Ding. All authors contributed to the main results and numerical simulations.

References

- 1. N. Bidmon, S. Kind, M. J. P. Welters, D. Joseph-Pietras, K. Laske, D. Maurer, et al., Development of an RNA-based kit for easy generation of TCR-engineered lymphocytes to control T-cell assay performance, *J. Immunol. Methods*, **458** (2018), 74–82. doi: 10.1016/j.jim.2018.04.007.
- L. Chen, D. Qiao, J. Wang, G. Tian, M. Wang, Cancer immunotherapy with lymphocytes genetically engineered with T cell receptors for solid cancers, *Immunol. Lett.*, **216** (2019), 51–62. doi: 10.1016/j.imlet.2019.10.002.
- 3. M. Yu, G. Huang, Y. Dong, Y. Takeuchi, Complicated dynamics of tumor-immune system interaction model with distributed time delay, *Discrete Cont. Dyn-B*, **7** (2020), 2391–2406. doi: 10.3934/dcdsb.2020015.

- 4. L. Han, C. He, Y. Kuang, Dynamics of a model of tumor-immune interaction with time delay and noise, *Discrete Cont. Dyn-S*, **9** (2020), 2347–2363. doi: 3934/dcdss.2020140.
- P. Bi, S. Ruan, Bifurcations in delay differential equations and applications to tumor and immune system interaction models, *SIAM J. Appl. Dyn. Syst.*, **12** (2013), 1847–1888. doi: 10.1137/120887898.
- 6. R. Yafia, Hopf bifurcation in differential equations with delay for tumor-immune system competition model, *SIAM J. Appl. Math.*, **6** (2007), 1693–1703. doi: 10.1137/060657947.
- L. Pang, S. Liu, X. Zhang, T. Tian, Mathematical modeling and dynamic analysis of antitumor-immune response, *J. Appl. Math. Comput.*, 62 (2020), 473–488. doi: 10.1007/s12190-019-01292-9.
- 8. Y. Jia, Bifurcation and pattern formation of a tumor-immune model with time-delay and diffusion, *Math. Comput. Simulat.*, **178** (2020), 92–108. doi: 10.1016/j.matcom.2020.06.011.
- 9. S. Banerjee, R. P. Sarkar. Delay-induced model for tumor-immune interaction and control of malignant tumor growth, *Biosystems*, **91** (2008), 268–288. doi: 10.1016/j.biosystems.2007.10.002.
- 10. S. Khajanchi, S. Banerjee, Stability and bifurcation analysis of delay induced tumor immune interaction model, *Appl. Math. Comput.*, **248** (2014), 652–671. doi: 10.1016/j.amc.2014.10.009.
- 11. L. R. Dickman, Y. Kuang, Analysis of tumor-immune dynamics in a delayed dendritic cell therapy model, *Chaos*, **11** (2020), 113108. doi: 10.1063/5.0006567.
- A. Kaddar, H. T. Alaoui, Global existence of periodic solution in a delayed tumor-immune model, *Math. Model Nat. Pheno.*, 5 (2010), 29–34. doi: 10.1051/mmnp/20105705.
- 13. M. Yu, Y. Dong, Y. Takeuchi, Dual role of delay effects in a tumour-immune system, *J. Biol. Dynam.*, **11** (2017), 334–347. doi: 10.1080/17513758.2016.1231347.
- F. A. Rihan, S. Lakshmanan, H. Maurer, Optimal control of tumour-immune model with time-delay and immuno-chemotherapy, *Appl. Math. Comput.*, 353 (2019), 147–165. doi: 10.1016/j.amc.2019.02.002.
- 15. F. A. Rihan, G. Velmurugan, for tumor-immune system, 10.1016/j.chaos.2019.109592.
 Dynamics of fractional-order delay differential model *Chaos Solitons Fractals*, **132** (2020), 109592. doi:
- F. A. Rihan, D. H. Abdel Rahman, S. Lakshmanan, A. S. Alkhajeh, A time delay model of tumour-immune system interactions: Global dynamics, parameter estimation, sensitivity analysis, *Appl. Math. Comput.*, 232 (2014), 606–623. doi: 10.1016/j.amc.2014.01.111.
- 17. P. Das, P. Das, S. Mukherjee, Stochastic dynamics of Michaelis-Menten kinetics based tumorimmune interactions, *Physica* A, **541** (2020), 123603. doi: 10.1016/j.physa.2019.123603.
- P. Das, P. Das, S. Das, An investigation on Monod-Haldane immune response based tumoreffector-interleukin-2 interactions with treatments, *Appl. Math. Comput.*, **361** (2019), 536–551. doi: 10.1016/j.amc.2019.05.032.
- 19. P. Das, R. K. Upadhyay, P. Das, D. Ghosh, Exploring dynamical complexity in a time-delayed tumor-immune model, *Chaos*, **30** (2020), 123118. doi: 10.1063/5.0025510.

- P. Das, S. Mukherjee, P. Das, An investigation on Michaelis-Menten kinetics based complex dynamics of tumor-immune interaction, *Chaos Solitons Fractals*, **128** (2019), 297–305. doi: 10.1016/j.chaos.2019.08.006.
- P. Das, S. Mukherjee, P. Das, S. Banerjee, Characterizing chaos and multifractality in noise-assisted tumor-immune interplay, *Nonlinear Dynam.*, **101** (2020), 675–685. doi: 10.1007/s11071-020-05781-6.
- P. Das, S. Das, P. Das, F. A. Rihan, M. Uzuntarla, D. Ghosh, Optimal control strategy for cancer remission using combinatorial therapy: A mathematical model-based approach, *Chaos Solitons Fractals*, 145 (2021), 110789. doi: 10.1016/j.chaos.2021.110789.
- P. Das, S. Das, R. K. Upadhyay, P. Das, Optimal treatment strategies for delayed cancer-immune system with multiple therapeutic approach, *Chaos Solitons Fractals*, **136** (2020), 109806. doi: 10.1016/j.chaos.2020.109806.
- 24. K. E. de Visser, A. Eichten, L. M. Coussens, Paradoxical roles of the immune system during cancer development, *Nat. Rev. Cancer*, **6** (2006), 24–37. doi: 10.1038/nrc1782.
- 25. J. Xie, T. Zhao, F. Hao, F. He, The effect of time delay on tumor-immune system during tumor growth, *J. Med. Biomech.*, **32** (2017), 319–324. doi: 10.16156/j.1004-7220.2017.04.004
- 26. T. Igakura, J. C. Stinchcomeb, P. K. C. Goon, G. P. Taylor, J. N. Weber, G. M. Griffiths, et al. Spread of HTLV-I between lymphocytes by virus-induced polarization of the cytoskeleton, *Sciences*, **5613** (2003), 1713–1716. doi: 10.1126/science.1080115.
- 27. S. Ruan, Nonlinear dynamics in tumor-immune system interaction models with delays, *Discrete Cont. Dyn-B*, **26** (2021), 541–602. doi: 10.3934/dcdsb.2020282.

Appendix A

In this section, we show the code of MATLAB software for simulating partial differential equations (that is, Figure 2).

```
function pdex43
m = 0; x = 0 : 0.1 : 1 * pi; t = 0 : 0.25 : 500;
sol = pdepe(m, @pdex44pde, @pdex44ic, @pdex44bc, x, t);
u1 = sol(:, :, 1);
u^2 = sol(:, :, 2);
figure
surf(x, t, u1); title('u(x, t)'); xlabel('x'); ylabel('t')
figure
surf(x, t, u2); title('v(x, t)'); xlabel('x'); ylabel('t')
function[c, f, s] = pdex44pde(x, t, u, DuDx)
d1 = 0.02; d2 = 0.2; r = 0.18; alpha = 5.0 * 10^{6}; m = 1.101 * 10^{-7};
beta = 6.2 * 10^{-9}; d = 0.0152; s = 3.422 * 10^{-10};
c = [1; 1];
f = [d1; d2]. * DuDx;
s = [r * u(1) * (1 - u(1)/alpha) - m * u(1) * u(2); beta * u(1) * u(2) - d * u(2) - s * u(1) * u(2)];
function u0 = pdex44ic(x);
u0 = [300000; 800000];
function [pl, ql, pr, qr] = pdex44bc(xl, ul, xr, ur, t)
pl = [0; 0]; ql = [1; 1]; pr = [0; 0]; qr = [1; 1];
```

Appendix B

In this section, we show the code of MATLAB software for simulating partial functional differential equations (that is, Figures 3-5).

function tumor – delay $d1 = 0.02; d2 = 0.2; r = 0.18; alpha = 5.0 * 10^{6}; m = 1.101 * 10^{-7};$ *beta* = 6.2×10^{-9} ; *d* = 0.0152; *s* = 3.422×10^{-10} ; q = 3;l = 1 * pi;N = 10; h = q/N; K = 6000; T = h * K; p = K + N + 1;M = 100; dx = l/M; u = zeros(M + 1, p); v = zeros(M + 1, p);for i = 1 : M + 1for j = 1 : N + 1u(i, j) = (2594700); v(i, j) = (786420);end end f = zeros(M + 1, p); g = zeros(M + 1, p);for j = N + 1 : p - 1for i = 2 : Mf(i, j) = r * u(i, j) * (1 - u(i, j - N)/alpha) - m * u(i, j) * v(i, j);g(i, j) = beta * u(i, j) * v(i, j) - d * v(i, j) - s * u(i, j) * v(i, j);u(i, j + 1) = u(i, j) + h * d1 * (u(i + 1, j) - 2 * u(i, j) + u(i - 1, j))/dx/dx + h * f(i, j);v(i, j+1) = v(i, j) + h * d2 * (v(i+1, j) - 2 * v(i, j) + v(i-1, j))/dx/dx + h * g(i, j);end u(1, j + 1) = u(2, j + 1); u(M + 1, j + 1) = u(M, j + 1);v(1, j + 1) = v(2, j + 1); v(M + 1, j + 1) = v(M, j + 1);end z1 = zeros(p, M + 1); z2 = zeros(p, M + 1);for i = 1 : M + 1for j = 1 : pz1(j,i) = u(i,j); z2(j,i) = v(i,j);end end x = 0: dx: 1 * pi; t = -q: h: T;[xx, tt] = meshgrid(x, t);figure; mesh(xx, tt, z1); title('u(x, t)'); xlabel('x'); ylabel('t'); figure; mesh(xx, tt, z2); title('v(x, t)'); xlabel('x'); ylabel('t');



© 2022 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)