



Research article

Mathematical analysis of an age-structured HIV-1 infection model with CTL immune response

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Abstract: In this paper, an age-structured HIV-1 infection model with CTL immune response is investigated. In the model, we consider the infection age (i.e. the time that has elapsed since an HIV virion has penetrated the cell) of infected T cells. The asymptotic smoothness of the semi-flow generated by the system is established. By calculation, the immune-inactivated reproduction rate \mathcal{R}_0 and the immune-activated reproduction rate \mathcal{R}_1 are obtained. By analyzing the corresponding characteristic equations, the local stability of an infection-free steady state and a CTL-inactivated infection steady state of the model is established. By using the persistence theory for infinite dimensional system, the uniform persistence of the system is established when $\mathcal{R}_1 > 1$. By means of suitable Lyapunov functionals and LaSalle's invariance principle, it is shown that if $\mathcal{R}_0 < 1$, the infection-free steady state is globally asymptotically stable; if $\mathcal{R}_1 < 1 < \mathcal{R}_0$, sufficient conditions are derived for the global stability of the CTL-inactivated infection steady state; if $\mathcal{R}_1 > 1$, sufficient conditions are obtained for the global attractivity of the CTL-activated infection steady state. Numerical simulations are carried out to illustrate the feasibility of the theoretical results.

Keywords: infection age; CTL immune response; persistence; Lyapunov functional; stability

1. Introduction

Human immunodeficiency virus (HIV) is a pathogen that infects T-helper cells of the immune system and can cause Acquired Immune Deficiency Syndrome (AIDS) [1]. These are white blood cells that move around the body, detecting faults and anomalies in cells as well as infections. When HIV targets and infiltrates these cells, it reduces the body's ability to combat other diseases. This increases the risk and impact of opportunistic infections and cancers. In past decades, many works have been developed for HIV-1 infection using simple differential equation models (see, for

example, [2–8]). Let x, y and v denote the concentrations of uninfected target cells (i.e. cells susceptible to HIV-1 infection), productively infected cells, and free virions, respectively. A basic mathematical model describing HIV-1 infection dynamics has been studied in [4, 8]:

$$\begin{aligned}\dot{x}(t) &= \Lambda - dx(t) - \beta x(t)v(t), \\ \dot{y}(t) &= \beta x(t)v(t) - ay(t), \\ \dot{v}(t) &= ky(t) - uv(t),\end{aligned}\tag{1.1}$$

where uninfected cells are produced at a rate Λ and die at rate dx per target cell, and become infected at rate βxv , where β is the rate constant describing the infection process; infected cells are produced at rate βxv and die at rate ay ; free virions are produced from infected cells at rate ky and are removed at rate uv .

We note that in system (1.1), the death rate and virus production rate of infected cells are both assumed to be constant. In reality, as argued by Nelson et al. [9], the production of new virus particles (virions) by an infected cell does not occur at a constant rate, but rather ramps up as viral proteins and unspliced viral RNA accumulate within the cytoplasm of an infected cell. In [9], in order to describe this phenomenon, by considering the variations in the death rate of productively infected T cells and the productions rate of viral particles as a function of the length of time a T cells has been infected, Nelson et al. developed and analyzed the following age-structured within-host HIV-1 infection model:

$$\begin{aligned}\dot{T}(t) &= s - dT(t) - \beta T(t)V(t), \\ \frac{\partial T^*(a, t)}{\partial t} + \frac{\partial T^*(a, t)}{\partial a} &= -\mu(a)T^*(a, t), \\ \dot{V}(t) &= \int_0^\infty p(a)T^*(a, t)da - uV(t),\end{aligned}\tag{1.2}$$

with boundary condition $T^*(0, t) = \beta T(t)V(t)$. In system (1.2), $T^*(a, t)$ denotes the density of infected T cells of infection age a (i.e. the time that has elapsed since an HIV virion has penetrated the cell) at time t , $\mu(a)$ is the age-dependent per capita death rate of infected cells, $p(a)$ is the viral production rate of an infected cell with age a . In [9], Nelson et al. discussed the local stability of the nontrivial equilibrium solution and provided a general stability condition for models with age structure. In [10], by constructing suitable Lyapunov functions, Huang et al. established the global dynamical properties for Nelson's age-structured model without (or with) drug treatment. In [11], Rong et al. considered two models with age-of-infection and combination therapies involving reverse transcriptase, protease, and entry/fusion inhibitors. In [12], considering the infection rate of microparasitic infections is an increasing function of the parasite dose, Xu et al. further investigated a within-host HIV-1 infection model with saturation incidence and age-since-infection structure for infected cells. Recently, great attention has been paid by many researchers to age-structured model of HIV infection due to their greater flexibility in exploring fundamental issues of viral production and death, and allow coupling of biological processes happening on different time scales (see, for example, [10–17]).

It is worth noting that the effect of immune response is ignored in the models above. However, in most virus infections, cytotoxic T lymphocytes (CTLs) play a critical role in cell-mediated immunity by regulating the functions of other immune cells (such as the B cells and macrophages) and attacking diseased cells and tumors [18]. For HIV-1 infection, the main clinical indicators of that HIV-1 positive patient are in the follow up both the viral load and the $CD4^+$ T cells count in blood plasma,

therapy is started, make a portion to the immune cells to be toxic thereby introducing toxicity in the immune system of the individual [19]. In [20], Cao et al. have shown that CTL immune response is often associated with better virus control and slower disease progression during the early stage of HIV infection. In [4], in order to discuss the effect of the population dynamics of viral infection with CTL immune response, Nowak et al. proposed the basic HIV-1 infection model with immune response. At present, there are few works on global dynamics in the age-structured within-host HIV-1 infection model with CTL immune response [21, 22].

Motivated by the works of Nelson et al. [9], Nowak et al. [4] and Regoes et al. [23], in the present paper, we are concerned with the effects of age-structured, CTL immune response and saturation incidence on the dynamics of HIV-1 infection. To this end, we consider the following HIV-1 infection model:

$$\begin{aligned} \dot{x}(t) &= \Lambda - dx(t) - \frac{\beta x(t)v(t)}{1 + \alpha v(t)}, \\ \frac{\partial y(a, t)}{\partial t} + \frac{\partial y(a, t)}{\partial a} &= -\mu(a)y(a, t) - p(a)y(a, t)z(t), \quad a > 0, \\ \dot{v}(t) &= \int_0^\infty k(a)y(a, t)da - uv(t), \\ \dot{z}(t) &= z(t) \int_0^\infty c(a)y(a, t)da - bz(t), \end{aligned} \tag{1.3}$$

with boundary condition

$$y(0, t) = \frac{\beta x(t)v(t)}{1 + \alpha v(t)}, \tag{1.4}$$

and initial condition

$$X_0 := (x(0), y(\cdot, 0), v(0), z(0)) = (x^0, y_0(\cdot), v^0, z^0) \in \mathcal{X}, \tag{1.5}$$

where $\mathcal{X} = \mathbb{R}^+ \times L_+^1(0, \infty) \times \mathbb{R}^+ \times \mathbb{R}^+$, $L_+^1(0, \infty)$ is the set of all integrable functions from $(0, \infty)$ into $\mathbb{R}^+ = [0, \infty)$. In system (1.3), $x(t)$ represents the concentration of uninfected target T cells at time t , $y(a, t)$ denotes the density of infected T cells of infection age a at time t , $v(t)$ denotes the concentration of infectious free virion at time t , and $z(t)$ denotes the concentration of CTLs at time t . The definitions of all parameters in system (1.3) are listed in Table 1.

In the sequel, we further make the following assumptions.

- (H1) $c, k, p, \mu \in L_+^\infty(0, \infty)$ with essential upper bounds $\bar{c}, \bar{k}, \bar{p}$ and $\bar{\mu}$, respectively.
- (H2) There are positive constants μ_0, μ_1, \bar{b} and satisfying $\mu_0 \leq d$, $\bar{b} = b - \bar{c} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\}$ and $\mu_1 = \min\{\mu_0, u, \bar{b}\}$. $\mu(a)$ is a bounded function on \mathbb{R}^+ satisfying $\mu(a) \geq \mu_0$ for $a \geq 0$.
- (H3) For any $a > 0$, there exist $a_k, a_c > a$ such that $k(a)$ is positive in a neighbourhood of a_k and $c(a)$ is positive in a neighbourhood of a_c .

Using the theory of age-structured dynamical systems introduced in [24, 25], one can show that system (1.3) has a unique solution $(x(t), y(\cdot, t), v(t), z(t))$ satisfying the boundary condition (1.4) and the initial condition (1.5). Moreover, it is easy to show that all solutions of system (1.3) with the boundary condition (1.4) and the initial condition (1.5) are defined on $[0, +\infty)$ and remain positive for all $t \geq 0$. Furthermore, \mathcal{X} is positively invariant and system (1.3) exhibits a continuous semi-flow

$\Phi : \mathbb{R}^+ \times \mathcal{X} \rightarrow \mathcal{X}$, namely,

$$\Phi_t(X_0) = \Phi(t, X_0) := (x(t), y(\cdot, t), v(t), z(t)), \quad t \geq 0, \quad X_0 \in \mathcal{X}. \quad (1.6)$$

Given a point $(x, \varphi, v, z) \in \mathcal{X}$, we have the norm $\|(x, \varphi, v, z)\|_{\mathcal{X}} := x + \int_0^\infty \varphi(a) da + v + z$.

Table 1. The definitions of the parameters in system (1.1).

| Parameter | Description |
|-----------|--|
| Λ | The recruitment rate of healthy T cells |
| d | The per capita death rate of uninfected cells |
| u | Clearance rate of virions |
| α | Saturation constant |
| β | The rate at which an uninfected cell becomes infected by an infectious virus |
| a | Age of infection, that is, the time since an HIV virion penetrated cell |
| b | The death rate of CTL cells |
| $\mu(a)$ | The age-dependent per capita death rate of infected cells with age a |
| $k(a)$ | The viral production rate of infected cells with age a |
| $p(a)$ | The killing rate of infected cells with age a |
| $c(a)$ | The proliferate rate of virus-specific CTL cells with age a |

The organization of this paper is as follows. In the next section, we establish the asymptotic smoothness of the semi-flow generated by system (1.3). In Section 3, we calculate the immune-inactivated reproduction rate and the immune-activated reproduction rate and discuss the existence of feasible steady states of system (1.3) with the boundary condition (1.4). In Section 4, by analyzing the corresponding characteristic equations, we study the local asymptotic stability of an infection-free steady state and a CTL-inactivated infection steady state of system (1.3), respectively. In Section 5, we show that if the immune-activated reproduction rate is greater than unity, system (1.3) is uniformly persistent. In Section 6, we are concerned with the global stability (attractivity) of each of feasible steady states of system (1.3) by means of Lyapunov functionals and LaSalle's invariance principle. In Section 7, numerical examples are carried out to illustrate the feasibility of theoretical results. A brief conclusion is given in Section 8 to end this work.

2. Boundedness and Asymptotic smoothness

In order to discuss the global dynamics of system (1.3) with the boundary condition (1.4), in this section, we are concerned with the boundedness of solutions of system (1.3) and the asymptotic smoothness of the semi-flow $\{\Phi(t)\}_{t \geq 0}$ generated by system (1.3).

2.1. Boundedness of solutions

In this subsection, we prove the boundedness of semi-flow $\{\Phi(t)\}_{t \geq 0}$. Denote

$$\|X_0\|_{\mathcal{X}} = x^0 + \int_0^\infty y_0(a) da + v^0 + z^0$$

and

$$N(t) = \|\Phi(t, X_0)\|_{\mathcal{X}} = x(t) + \int_0^\infty y(a, t) da + v(t) + z(t).$$

Proposition 2.1. *Let Φ_t be defined as in (1.6). Then the following statements hold.*

- (i) $\frac{d}{dt}\|\Phi_t(X_0)\|_{\mathcal{X}} \leq \Lambda + \bar{k} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\} - \mu_1 N(t)$ for all $t \geq 0$;
- (ii) $\|\Phi_t(X_0)\|_{\mathcal{X}} \leq \max\left\{\frac{1}{\mu_1} \left[\Lambda + \bar{k} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\}\right], \|X_0\|_{\mathcal{X}}\right\}$ for all $t \geq 0$;
- (iii) $\limsup_{t \rightarrow +\infty} \|\Phi_t(X_0)\|_{\mathcal{X}} \leq \frac{1}{\mu_1} \left[\Lambda + \bar{k} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\}\right]$;
- (iv) Φ_t is point dissipative: there is a bounded set that attracts all points in \mathcal{X} .

Proof. Let $\Phi_t(X_0) = \Phi(t, X_0) := (x(t), y(\cdot, t), v(t), z(t))$ be any nonnegative solution of system (1.3) with the boundary condition (1.4) and the initial condition (1.5). We derive from system (1.3) that

$$\begin{aligned} \frac{d}{dt} \left(x(t) + \int_0^\infty y(a, t) da \right) &= \Lambda - dx(t) - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} - y(a, t)|_0^\infty \\ &\quad - \int_0^\infty (\mu(a) + p(a)z(t))y(a, t) da \\ &\leq \Lambda - \mu_0 \left(x(t) + \int_0^\infty y(a, t) da \right). \end{aligned} \quad (2.1)$$

The variation of constants formula implies

$$x(t) + \int_0^\infty y(a, t) da \leq \frac{\Lambda}{\mu_0} - e^{-\mu_0 t} \left(\frac{\Lambda}{\mu_0} - \|X_0\|_{\mathcal{X}} \right),$$

which yields

$$x(t) + \int_0^\infty y(a, t) da \leq \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\} \quad (2.2)$$

for all $t \geq 0$. We derive from Eq (2.2) and the third and fourth equations of system (1.3) that

$$\begin{aligned} \frac{dv(t)}{dt} &\leq \bar{k} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\} - uv(t), \\ \frac{dz(t)}{dt} &\leq \bar{c} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\} z(t) - bz(t). \end{aligned} \quad (2.3)$$

It follows from Eqs (2.1) and (2.3) that

$$\begin{aligned} \frac{d}{dt} N(t) &\leq \Lambda - \mu_0 \left(x(t) + \int_0^\infty y(a, t) da \right) + \bar{k} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\} - uv(t) - \bar{b}z(t) \\ &\leq \Lambda + \bar{k} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\} - \mu_1 N(t). \end{aligned} \quad (2.4)$$

Again, using variation of constants formula we have from Eq (2.4) that

$$N(t) \leq \max\left\{\frac{1}{\mu_1} \left[\Lambda + \bar{k} \max\left\{\frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}}\right\}\right], \|X_0\|_{\mathcal{X}}\right\}$$

for all $t \geq 0$. This completes the proof.

The following results are direct consequences of Proposition 2.1.

Proposition 2.2. *If $X_0 \in \mathcal{X}$ and $\|X_0\|_{\mathcal{X}} \leq K$ for some $K \geq \frac{1}{\mu_1} \left[\Lambda + \bar{k} \max \left\{ \frac{\Lambda}{\mu_0}, \|X_0\|_{\mathcal{X}} \right\} \right]$, then*

$$x(t) \leq K, \int_0^\infty y(a, t) da \leq K, v(t) \leq K, z(t) \leq K \quad (2.5)$$

for all $t \geq 0$.

Proposition 2.3. *Let $C \in \mathcal{X}$ be bounded. Then*

- (1) $\Phi_t(C)$ is bounded;
- (2) Φ_t is eventually bounded on C .

2.2. Asymptotic smoothness

In this subsection, we show the asymptotic smoothness of the semi-flow $\{\Phi(t)\}_{t \geq 0}$.

Denote

$$\pi(a) = e^{-\int_0^a (\mu(s) + p(s)z) ds}, \quad a \in \mathbb{R}^+. \quad (2.6)$$

It follows from (H1) and (H2) that $0 < e^{-(\bar{\mu} + \bar{p}K)a} \leq \pi(a) \leq e^{-\mu_0 a}$ for all $a \geq 0$. Clearly, $\pi(a)$ is a decreasing function.

Let $(x(t), y(\cdot, t), v(t), z(t))$ be a solution of system (1.3) with the boundary condition (1.4) and the initial condition (1.5). Integrating the second equation of system (1.3) along the characteristic line $t - a = \text{const.}$, we have

$$y(a, t) = \begin{cases} L(t - a)\pi(a), & 0 \leq a < t, \\ y_0(a - t) \frac{\pi(a)}{\pi(a - t)}, & 0 \leq t \leq a, \end{cases} \quad (2.7)$$

where $L(t) := y(0, t) = \frac{\beta x(t)v(t)}{1 + \alpha v(t)}$.

Using a similar argument as that in [12], it's easy to verify the following result.

Proposition 2.4. *The function $L(t)$ is Lipschitz continuous on \mathbb{R}^+ .*

Before giving our main results, we need the following Lemmas.

Lemma 2.1. [26] *The semi-flow $\Phi : \mathbb{R}^+ \times \mathcal{X}_+ \rightarrow \mathcal{X}_+$ is asymptotically smooth if there are maps $\Theta, \Psi : \mathbb{R}^+ \times \mathcal{X}_+ \rightarrow \mathcal{X}_+$ such that $\Phi(t, x) = \Theta(t, x) + \Psi(t, x)$ and the following hold for any bounded closed set $C \subset \mathcal{X}_+$ that is forward invariant under Φ :*

- (1) $\lim_{t \rightarrow +\infty} \text{diam} \Theta(t, C) = 0$;
- (2) there exists $t_C \geq 0$ such that $\Psi(t, C)$ has compact closure for each $t \geq t_C$.

Lemma 2.2. [26] *Let C be a subset of $L^1(\mathbb{R}^+)$. Then C has compact closure if and only if the following assumptions hold:*

- (i) $\sup_{f \in C} \int_0^\infty |f(a)| da < \infty$;
- (ii) $\lim_{r \rightarrow \infty} \int_r^\infty |f(a)| da = 0$ uniformly in $f \in C$;

- (iii) $\lim_{h \rightarrow 0^+} \int_0^\infty |f(a+h) - f(a)| da = 0$ uniformly in $f \in C$;
 (iv) $\lim_{h \rightarrow 0^+} \int_0^h |f(a)| da = 0$ uniformly in $f \in C$.

By applying Lemmas 2.1 and 2.2, we now prove the asymptotic smoothness of the semiflow Φ generated by system (1.3).

Theorem 2.1. *The semi-flow Φ generated by system (1.3) is asymptotically smooth.*

Proof. We first decompose the semi-flow Φ into two parts: for $t \geq 0$, let $\Psi(t, X_0) := (x(t), \tilde{y}(\cdot, t), v(t), z(t))$, $\Theta(t, X_0) := (0, \tilde{\phi}_y(\cdot, t), 0, 0)$, where

$$\begin{aligned} \tilde{y}(a, t) &= \begin{cases} L(t-a)\pi(a), & 0 \leq a \leq t, \\ 0, & 0 \leq t < a, \end{cases} \\ \tilde{\phi}_y(a, t) &= \begin{cases} 0, & 0 \leq a \leq t, \\ y_0(a-t) \frac{\pi(a)}{\pi(a-t)}, & 0 \leq t < a. \end{cases} \end{aligned} \quad (2.8)$$

Clearly, we have $\Phi = \Theta + \Psi$ for $t \geq 0$.

Let C be a bounded subset of \mathcal{X} and $K > [\Lambda + \bar{k} \max\{\frac{\Delta}{\mu_0}, \|X_0\|_{\mathcal{X}}\}]/\mu_1$ the bound for C . Let $\Phi(t, X_0) = (x(t), y(\cdot, t), v(t), z(t))$, where $X_0 = (x^0, y_0(\cdot), v^0, z^0) \in C$. Then

$$\begin{aligned} \|\Theta(t, X_0)\| &= \|\tilde{\phi}_y(\cdot, t)\|_{L^1} = \int_0^\infty |\tilde{\phi}_y(a, t)| da \\ &= \int_t^\infty y_0(a-t) \frac{\pi(a)}{\pi(a-t)} da. \end{aligned} \quad (2.9)$$

Letting $a - t = \sigma$, it follows from (2.9) that

$$\begin{aligned} \|\tilde{\phi}_y(\cdot, t)\|_{L^1} &= \int_0^\infty y_0(\sigma) e^{-\int_\sigma^{\sigma+t} (\mu(s)+p(s)z) ds} d\sigma \\ &\leq K e^{-\mu_0 t}, \end{aligned} \quad (2.10)$$

yielding $\lim_{t \rightarrow +\infty} \|\Theta(t, X_0)\| = 0$, hence, $\lim_{t \rightarrow +\infty} \text{diam } \Theta(t, C) = 0$ and the assumption (1) in Lemma 2.1 holds.

In the following we show that $\Psi(t, C)$ has compact closure for each $t \geq t_C$ by verifying the assumptions (i)–(iv) of Lemma 2.2. From Proposition 2.2 we see that $x(t)$, $v(t)$ and $z(t)$ remain in the compact set $[0, K]$. Next, we show that $\tilde{y}(a, t)$ remain in a pre-compact subset of L^1_+ independent of X_0 . It is easy to show that $\tilde{y}(a, t) \leq \bar{L} e^{-\mu_0 a}$, where $\bar{L} = \beta K^2 / (1 + \alpha K)$. Therefore, the assumptions (i),(ii) and (iv) of Lemma 2.2 follow directly. We need only to verify that (iii) of Lemma 2.2 holds.

Since we are concerned with the limit as $h \rightarrow 0$, we assume that $h \in (0, t)$. In this case, we have

$$\begin{aligned}
 \int_0^\infty |\tilde{y}(a+h, t) - \tilde{y}(a, t)| da &= \int_0^{t-h} |L(t-a-h)\pi(a+h) - L(t-a)\pi(a)| da \\
 &\quad + \int_{t-h}^t L(t-a)\pi(a) da \\
 &\leq \int_0^{t-h} L(t-a-h) |\pi(a+h) - \pi(a)| da \\
 &\quad + \int_0^{t-h} |L(a-t-h) - L(a-t)| \pi(a) da \\
 &\quad + \int_{t-h}^t L(t-a)\pi(a) da.
 \end{aligned} \tag{2.11}$$

By Proposition 2.2., there is positive constant M_L such that

$$|L(a+h) - L(a)| \leq M_L h. \tag{2.12}$$

It follows from (2.11) and (2.12) that

$$\begin{aligned}
 \int_0^\infty |\tilde{y}(a+h, t) - \tilde{y}(a, t)| da &\leq \bar{L} \int_0^{t-h} \pi(a) \int_a^{a+h} (\mu(s) + p(s)z) ds da + M_L h + \bar{L} h \\
 &\leq [(\bar{\mu} + \bar{p}K)\bar{L} + M_L + \bar{L}] h.
 \end{aligned} \tag{2.13}$$

Hence, the condition (iii) of Lemma 2.2 holds. By Lemma 2.1, the asymptotic smoothness of the semi-flow Φ generated by system (1.3) follows. This completes the proof.

The following result is immediate from Theorem 2.33 in [26] and Theorem 2.1.

Theorem 2.2. *There exists a global attractor \mathcal{A} of bounded sets in \mathcal{X} .*

3. Steady states and basic reproduction number

In this section, we are concerned with the local stability of each of feasible steady states of system (1.3) with the boundary condition (1.4).

Clearly, system (1.3) always has an infection-free steady state $E_0(\Lambda/d, 0, 0, 0)$. If system (1.3) has a CTL-inactivated infection steady state $E_1(x_1, y_1(a), v_1, 0)$, then it must satisfy the following equations:

$$\begin{aligned}
 \Lambda - dx_1 - \frac{\beta x_1 v_1}{1 + \alpha v_1} &= 0, \\
 \frac{d}{da} y_1(a) &= -\mu(a) y_1(a), \\
 \int_0^\infty k(a) y_1(a) da &= uv_1, \\
 y_1(0) &= \frac{\beta x_1 v_1}{1 + \alpha v_1}.
 \end{aligned} \tag{3.1}$$

We derive from the first equation of (3.1) that

$$x_1 = \frac{\Lambda(1 + \alpha v_1)}{d + (\alpha d + \beta)v_1}. \quad (3.2)$$

It follows from the second equation of (3.1) that

$$y_1(a) = y_1(0)\phi_1(a), \quad (3.3)$$

where $\phi_1(a) = e^{-\int_0^a \mu(s)ds}$. We obtain from the third equation of (3.1) and (3.3) that

$$v_1 = \frac{y_1(0) \int_0^\infty k(a)\phi_1(a)da}{u}. \quad (3.4)$$

On substituting (3.2)–(3.4) into the fourth equation of (3.1), we have

$$y_1(0) = \frac{du}{(\alpha d + \beta) \int_0^\infty k(a)\phi_1(a)da} (\mathcal{R}_0 - 1), \quad (3.5)$$

where

$$\mathcal{R}_0 = \frac{\Lambda\beta \int_0^\infty k(a)\phi_1(a)da}{du}. \quad (3.6)$$

Here, \mathcal{R}_0 is called the immune-inactivated reproduction rate of system (1.3), which represents the number of newly infected cells produced by one infected cell during its lifespan. Hence, if $\mathcal{R}_0 > 1$, in addition to the infection-free steady state E_0 , system (1.3) admits a CTL-inactivated infection steady state $E_1(x_1, y_1(a), v_1, 0)$, where

$$x_1 = \frac{\Lambda(1 + \alpha v_1)}{d + (\alpha d + \beta)v_1}, \quad y_1(a) = \frac{du\phi_1(a)}{(\alpha d + \beta) \int_0^\infty k(a)\phi_1(a)da} (\mathcal{R}_0 - 1), \quad v_1 = \frac{d}{\alpha d + \beta} (\mathcal{R}_0 - 1).$$

Further, if system (1.3) has a CTL-activated infection steady state $E^*(x^*, y^*(a), v^*, z^*)$, then it must satisfy the following equations:

$$\begin{aligned} \Lambda - dx^* - \frac{\beta x^* v^*}{1 + \alpha v^*} &= 0, \\ \frac{d}{da} y^*(a) &= -\mu(a)y^*(a) - p(a)y^*(a)z^*, \\ \int_0^\infty k(a)y^*(a)da &= uv^*, \\ z^* \left(\int_0^\infty c(a)y^*(a)da - b \right) &= 0, \\ y^*(0) &= \frac{\beta x^* v^*}{1 + \alpha v^*}. \end{aligned} \quad (3.7)$$

It follows from the first equation of (3.7) that

$$x^* = \frac{\Lambda(1 + \alpha v^*)}{d + (\alpha d + \beta)v^*}. \quad (3.8)$$

We obtain from the second equation of (3.7) that

$$y^*(a) = y^*(0)\phi_1(a)\phi_2(a, z^*), \quad (3.9)$$

where $\phi_2(a, z^*) = e^{-\int_0^a p(s)z^* ds}$. When $z^* \neq 0$, it follows from (3.9) and the fourth equation of (3.7) that

$$y^*(0) = \frac{b}{\int_0^\infty c(a)\phi_1(a)\phi_2(a, z^*)da}. \quad (3.10)$$

We derive from the third equation of (3.7) and (3.10) that

$$v^* = \frac{b \int_0^\infty k(a)\phi_1(a)\phi_2(a, z^*)da}{u \int_0^\infty c(a)\phi_1(a)\phi_2(a, z^*)da}. \quad (3.11)$$

On substituting (3.8), (3.10) and (3.11) into the first equation of (3.7), we have that

$$\Lambda\beta G_1(z^*)G_2(z^*) = duG_2(z^*) + b(\alpha d + \beta)G_1(z^*), \quad (3.12)$$

where

$$G_1(z^*) = \int_0^\infty k(a)\phi_1(a)\phi_2(a, z^*)da, \quad G_2(z^*) = \int_0^\infty c(a)\phi_1(a)\phi_2(a, z^*)da. \quad (3.13)$$

From (3.13), it is easy to show that

$$0 < G_i(z^*) \leq G_i(0) \quad \text{and} \quad G'_i(z^*) \leq 0, \quad i = 1, 2. \quad (3.14)$$

Denote

$$\begin{aligned} \Phi(z) &= \Lambda\beta G_1(z)G_2(z) - duG_2(z) - b(\alpha d + \beta)G_1(z) \\ &= [duG_2(z) + b(\alpha d + \beta)G_1(z)] \left(\frac{\Lambda\beta G_1(z)G_2(z)}{duG_2(z) + b(\alpha d + \beta)G_1(z)} - 1 \right). \end{aligned} \quad (3.15)$$

Clearly

$$\Phi(0) = [duG_2(0) + b(\alpha d + \beta)G_1(0)](\mathcal{R}_1 - 1), \quad (3.16)$$

where

$$\mathcal{R}_1 = \frac{\Lambda\beta \int_0^\infty k(a)\phi_1(a)da \int_0^\infty c(a)\phi_1(a)da}{du \int_0^\infty c(a)\phi_1(a)da + b(\alpha d + \beta) \int_0^\infty k(a)\phi_1(a)da}.$$

Here, \mathcal{R}_1 is called the immune-activated reproduction rate which expresses the CTL load during the lifespan of a CTL cell. Clearly, if $\mathcal{R}_1 > 1$, it therefore follows from (3.14) and (3.16) that $\Phi(0) > 0$. Further, for $z > 0$ sufficiently large, we note that

$$\frac{\Lambda\beta G_1(z)G_2(z)}{duG_2(z) + b(\alpha d + \beta)G_1(z)} \rightarrow 0,$$

Then by (3.15), for $z > 0$ sufficiently large, there exists a $z_0^* > 0$ such that $\Phi(z_0^*) < 0$. Therefore, if $\mathcal{R}_1 > 1$, there exists a $z^* \in (0, z_0^*)$ such that $\Phi(z^*) = 0$. Hence, if $\mathcal{R}_1 > 1$, in addition to the infection-free steady state E_0 and the CTL-inactivated infection steady state E_1 , system (1.3) exists a unique infection steady state $E^*(x^*, y^*(a), v^*, z^*)$.

4. Local stability

In this section, we are concerned with the local stability of the infection-free steady state E_0 and the CTL-inactivated infection steady state E_1 of system (1.3), respectively.

We first consider the local stability of the infection-free steady state $E_0(\Lambda/d, 0, 0, 0)$.

Let $x(t) = x_0(t) + \Lambda/d, y(a, t) = y_0(a, t), v(t) = v_0(t), z(t) = z_0(t)$. Linearizing system (1.3) at the steady state E_0 , it follows that

$$\begin{aligned} \dot{x}_0(t) &= -dx_0(t) - \frac{\Lambda\beta}{d}v_0(t), \\ \frac{\partial y_0(a, t)}{\partial t} + \frac{\partial y_0(a, t)}{\partial a} &= -\mu(a)y_0(a, t), \\ \dot{v}_0(t) &= \int_0^\infty k(a)y_0(a, t)da - uv_0(t), \\ \dot{z}_0(t) &= -bz_0(t), \\ y_0(0, t) &= \frac{\Lambda\beta}{d}v_0(t). \end{aligned} \quad (4.1)$$

Looking for solutions of system (4.1) of the form $x_0(t) = x_{01}e^{\lambda t}, y_0(a, t) = y_{01}(a)e^{\lambda t}, v_0(t) = v_{01}e^{\lambda t}, z_0(t) = z_{01}e^{\lambda t}$, where $x_{01}, y_{01}(a), v_{01}$ and z_{01} will be determined later, one obtains the characteristic equation of system (1.3) at the steady state E_0 of the form:

$$(\lambda + b)(1 - f_1(\lambda)) = 0, \quad (4.2)$$

where

$$f_1(\lambda) = \frac{\Lambda\beta}{d(\lambda + u)} \int_0^\infty k(a)e^{-(\lambda + \mu(s))ds} da.$$

Clearly, Eq (4.2) always has one negative real root $\lambda = -b$, other roots of (4.2) are determined by equation

$$f_1(\lambda) = 1. \quad (4.3)$$

Clearly, we have $f_1(0) = \mathcal{R}_0$. It is easy to show that $f_1'(\lambda) < 0$ and $\lim_{\lambda \rightarrow +\infty} f_1(\lambda) = 0$. Hence, $f_1(\lambda)$ is a decreasing function. Therefore, if $\mathcal{R}_0 > 1$, then $f_1(\lambda) = 1$ has a unique positive root. Hence, if $\mathcal{R}_0 > 1$, the steady state E_0 is unstable.

Now, we claim that all roots of Eq (4.3) have negative real parts if $\mathcal{R}_0 < 1$. If not, there exists a root $\lambda_1 = a_1 + ib_1$ with $a_1 \geq 0$. In this case, substituting λ_1 into (4.3), we obtain

$$|f_1(\lambda_1)| \leq \frac{\Lambda\beta \int_0^\infty k(a)\phi_1(a)da}{du} = \mathcal{R}_0 < 1,$$

a contradiction. Hence, if $\mathcal{R}_0 < 1$, all roots of equation (4.2) have negative real parts. Accordingly, the steady state $E_0(\Lambda/d, 0, 0, 0)$ is locally asymptotically stable if $\mathcal{R}_0 < 1$.

Now, we consider the local stability of the CTL-inactivated infection steady state $E_1(x_1, y_1(a), v_1, 0)$.

Let $x(t) = x_1(t) + x_1, y(a, t) = y_1(a, t) + y_1(a), v(t) = v_1(t) + v_1, z(t) = z_1(t)$. Linearizing system (1.3) at the steady state E_1 , we obtain that

$$\begin{aligned} \dot{x}_1(t) &= -\left(d + \frac{\beta v_1}{1 + \alpha v_1}\right)x_1(t) - \frac{\beta x_1}{(1 + \alpha v_1)^2}v_1(t), \\ \frac{\partial y_1(a, t)}{\partial t} + \frac{\partial y_1(a, t)}{\partial a} &= -\mu(a)y_1(a, t) - p(a)y_1(a)z_1(t), \\ \dot{v}_1(t) &= \int_0^\infty k(a)y_1(a, t)da - uv_1(t), \\ \dot{z}_1(t) &= \left(\int_0^\infty c(a)y_1(a)da - b\right)z_1(t), \\ y_1(0, t) &= \frac{\beta v_1}{1 + \alpha v_1}x_1(t) + \frac{\beta x_1}{(1 + \alpha v_1)^2}v_1(t). \end{aligned} \quad (4.4)$$

Looking for solutions of system (4.4) of the form $x_1(t) = x_{11}e^{\lambda t}, y_1(a, t) = y_{11}(a)e^{\lambda t}, v_1(t) = v_{11}e^{\lambda t}, z_1(t) = z_{11}e^{\lambda t}$, where $x_{11}, y_{11}(a), v_{11}$ and z_{11} will be determined later, we obtain the following linear eigenvalue problem:

$$\begin{aligned} \left(\lambda + d + \frac{\beta v_1}{1 + \alpha v_1}\right)x_{11} &= -\frac{\beta x_1}{(1 + \alpha v_1)^2}v_{11}, \\ y'_{11}(a) &= -(\lambda + \mu(a))y_{11}(a) - p(a)y_{11}(a)z_{11}, \\ (\lambda + u)v_{11} &= \int_0^\infty k(a)y_{11}(a)da, \\ \lambda &= \int_0^\infty c(a)y_{11}(a)da - b, \\ y_{11}(0) &= \frac{\beta v_1}{1 + \alpha v_1}x_{11} + \frac{\beta x_1}{(1 + \alpha v_1)^2}v_{11}. \end{aligned} \quad (4.5)$$

We derive from the fourth equation of (4.5) that

$$\lambda = \left(\frac{du \int_0^\infty c(a)\phi_1(a)da}{(\alpha d + \beta) \int_0^\infty k(a)\phi_1(a)da} + b\right)(\mathcal{R}_1 - 1). \quad (4.6)$$

Clearly, If $\mathcal{R}_1 > 1, \lambda > 0$, in this case, E_1 is unstable. If $\mathcal{R}_1 < 1 < \mathcal{R}_0$, it follows from the fourth equation of (4.4) that $z_1 \rightarrow 0$, hence, in the following discussion, we only consider the simplified system

$$\begin{aligned} \left(\lambda + d + \frac{\beta v_1}{1 + \alpha v_1}\right)x_{11} &= -\frac{\beta x_1}{(1 + \alpha v_1)^2}v_{11}, \\ y'_{11}(a) &= -(\lambda + \mu(a))y_{11}(a), \\ (\lambda + u)v_{11} &= \int_0^\infty k(a)y_{11}(a)da, \\ y_{11}(0) &= \frac{\beta v_1}{1 + \alpha v_1}x_{11} + \frac{\beta x_1}{(1 + \alpha v_1)^2}v_{11}. \end{aligned} \quad (4.7)$$

It follows from the first and the second equations of system (4.7) that

$$x_{11} = -\frac{\beta x_1}{(1 + \alpha v_1)[(\lambda + d)(1 + \alpha v_1) + \beta v_1]}v_{11}, \quad (4.8)$$

and

$$y_{11}(a) = y_{11}(0)e^{-\int_0^a (\lambda + \mu(s))ds}. \quad (4.9)$$

We derive from the third equation of system (4.5) that

$$v_{11} = \frac{y_{11}(0) \int_0^\infty k(a)e^{-\int_0^a (\lambda + \mu(s))ds} da}{\lambda + u}. \quad (4.10)$$

On substituting (4.8)–(4.10) into the fifth equation of system (4.7), one obtains that

$$f_2(\lambda) = 1, \quad (4.11)$$

where

$$f_2(\lambda) = \frac{\beta x_1}{(1 + \alpha v_1)^2} \frac{(\lambda + d) \int_0^\infty k(a)e^{-\int_0^a (\lambda + \mu(s))ds} da}{(\lambda + u) \left(\lambda + d + \frac{\beta v_1}{1 + \alpha v_1} \right)}.$$

We claim that all roots of Eq (4.11) have negative real parts. Otherwise, Eq (4.11) has at least one root $\lambda_2 = a_2 + ib_2$ satisfying $a_2 \geq 0$. In this case, we have

$$|f_2(\lambda_2)| \leq \frac{1}{1 + \frac{\alpha d}{(\alpha d + \beta)}(\mathcal{R}_0 - 1)} \frac{u}{|a_2 + u + ib_2|}. \quad (4.12)$$

Clearly, if $\mathcal{R}_0 > 1$, then $1 + \frac{\alpha d}{(\alpha d + \beta)}(\mathcal{R}_0 - 1) > 1$, which mean that $|f_2(\lambda_2)| < 1$, a contradiction. Therefore, if $\mathcal{R}_1 < 1 < \mathcal{R}_0$, the CTL-inactivated infection steady state E_1 is locally asymptotically stable.

In conclusion, we have the following result.

Theorem 4.1. *For system (1.3) with the boundary condition (1.4), if $\mathcal{R}_0 < 1$, the infection-free steady state $E_0(\Lambda/d, 0, 0, 0)$ is locally asymptotically stable; if $\mathcal{R}_1 < 1 < \mathcal{R}_0$, E_0 is unstable and the CTL-inactivated infection steady state $E_1(x_1, y_1(a), v_1, 0)$ exists and is locally asymptotically stable.*

5. Uniform persistence

In this section, we investigate the uniform persistence of the semi-flow $\{\Phi(t)\}_{t \geq 0}$ generated by system (1.3) when the immune-activated reproduction rate $\mathcal{R}_1 > 1$.

Define

$$\bar{a}_1 = \inf \left\{ a : \int_a^\infty k(u)du = 0 \right\}, \quad \bar{a}_2 = \inf \left\{ a : \int_a^\infty c(u)du = 0 \right\}.$$

Noting that $k(\cdot), c(\cdot) \in L_+^\infty(0, \infty)$, we have $\bar{a}_1 > 0, \bar{a}_2 > 0$.

Denote

$$\begin{aligned} \mathcal{X} &= L_+^1(0, +\infty) \times \mathbb{R}^+ \times \mathbb{R}^+, \quad \bar{a} = \max\{\bar{a}_1, \bar{a}_2\} \\ \tilde{\mathcal{Y}} &= \left\{ (y(\cdot, t), v(t), z(t))^\top \in \mathcal{X} : \int_0^{\bar{a}} y(a, t)da + v(t) + z(t) > 0 \right\}, \end{aligned}$$

and

$$\mathcal{Y} = \mathbb{R}^+ \times \tilde{\mathcal{Y}}, \quad \partial \mathcal{Y} = \mathcal{X} \setminus \mathcal{Y}, \quad \partial \tilde{\mathcal{Y}} = \mathcal{X} \setminus \tilde{\mathcal{Y}}.$$

By [36] and using a similar argument as in the proof of Theorem 5.1 in [38], we have the following result.

Proposition 5.1. *The subsets \mathcal{Y} and $\partial\mathcal{Y}$ are both positively invariant under the semi-flow $\{\Phi(t)\}_{t \geq 0}$, namely, $\Phi(t, \mathcal{Y}) \subset \mathcal{Y}$ and $\Phi(t, \partial\mathcal{Y}) \subset \partial\mathcal{Y}$ for $t \geq 0$.*

The following result is helpful to the proof of uniform persistence of the semi-flow $\{\Phi(t)\}_{t \geq 0}$ generated by system (1.3).

Theorem 5.1. *The infection-free steady state $E_0(A/\mu, 0, 0, 0)$ is globally asymptotically stable for the semi-flow $\{\Phi(t)\}_{t \geq 0}$ restricted to $\partial\mathcal{Y}$.*

Proof. Let $(x^0, y_0(\cdot), v^0, z^0) \in \partial\mathcal{Y}$. Then $(y_0(\cdot), v^0, z^0) \in \partial\tilde{\mathcal{Y}}$. We consider the following system

$$\begin{aligned} \frac{\partial y(a, t)}{\partial t} + \frac{\partial y(a, t)}{\partial a} &= -\mu(a)y(a, t) - p(a)y(a, t)z(t), \\ \dot{v}(t) &= \int_0^\infty k(a)y(a, t)da - uv(t), \\ \dot{z}(t) &= z(t) \int_0^\infty c(a)y(a, t)da - bz(t), \\ y(0, t) &= \frac{\beta x(t)v(t)}{1 + \alpha v(t)}, \\ y(a, 0) &= y_0(a), \quad v(0) = 0, \quad z(0) = 0. \end{aligned} \tag{5.1}$$

Since $\limsup_{t \rightarrow +\infty} x(t) \leq \Lambda/d$, by comparison principle, we have

$$y(a, t) \leq \hat{y}(a, t), \quad v(t) \leq \hat{v}(t), \quad z(t) \leq \hat{z}(t), \tag{5.2}$$

where $\hat{y}(a, t)$, $\hat{v}(t)$ and $\hat{z}(t)$ satisfy the following auxiliary system

$$\begin{aligned} \frac{\partial \hat{y}(a, t)}{\partial t} + \frac{\partial \hat{y}(a, t)}{\partial a} &= -\mu(a)\hat{y}(a, t) - p(a)\hat{y}(a, t)\hat{z}(t), \\ \dot{\hat{v}}(t) &= \int_0^\infty k(a)\hat{y}(a, t)da - u\hat{v}(t), \\ \dot{\hat{z}}(t) &= \hat{z}(t) \int_0^\infty c(a)\hat{y}(a, t)da - b\hat{z}(t), \\ \hat{y}(0, t) &= \frac{\Lambda\beta}{d} \frac{\hat{v}(t)}{1 + \alpha\hat{v}(t)}, \\ \hat{y}(a, 0) &= y_0(a), \quad \hat{v}(0) = 0, \quad \hat{z}(0) = 0. \end{aligned} \tag{5.3}$$

Solving the first equation of system (5.3), we have

$$\hat{y}(a, t) = \begin{cases} \hat{L}(t-a)\pi(a), & 0 \leq a < t, \\ y_0(a-t) \frac{\pi(a)}{\pi(a-t)}, & 0 \leq t \leq a, \end{cases} \tag{5.4}$$

where

$$\hat{L}(t) := \hat{y}(0, t) = \frac{\Lambda\beta}{d} \frac{\hat{v}(t)}{1 + \alpha\hat{v}(t)}. \tag{5.5}$$

On substituting (5.5) into the second and the third equations of (5.3), it follows that

$$\begin{aligned}\hat{v}(t) &= \int_0^t k(a)\hat{L}(t-a)\pi(a)da - u\hat{v}(t) + L_1(t), \\ \hat{z}(t) &= \hat{z}(t) \int_0^t c(a)\hat{L}(t-a)\pi(a)da - b\hat{z}(t) + \hat{z}(t)L_2(t), \\ \hat{v}(0) &= 0, \quad \hat{z}(0) = 0.\end{aligned}\tag{5.6}$$

where

$$\begin{aligned}L_1(t) &= \int_t^\infty k(a)y_0(a-t)\frac{\pi(a)}{\pi(a-t)}da, \\ L_2(t) &= \int_t^\infty c(a)y_0(a-t)\frac{\pi(a)}{\pi(a-t)}da,\end{aligned}$$

Since $(y_0(\cdot), v^0, z^0) \in \partial\tilde{\mathcal{Y}}$, we have $L_i(t) \equiv 0$ ($i = 1, 2$) for all $t \geq 0$. It therefore follows from (5.6) that

$$\begin{aligned}\hat{v}(t) &= \int_0^t k(a)\hat{L}(t-a)\pi(a)da - u\hat{v}(t), \\ \hat{z}(t) &= \hat{z}(t) \int_0^t c(a)\hat{L}(t-a)\pi(a)da - b\hat{z}(t), \\ \hat{L}(t) &:= \hat{y}(0, t) = \frac{\Lambda\beta}{d} \frac{\hat{v}(t)}{1 + \alpha\hat{v}(t)}, \\ \hat{v}(0) &= 0, \quad \hat{z}(0) = 0.\end{aligned}\tag{5.7}$$

It is easy to show that system (5.7) has a unique solution $\hat{v}(t) = 0, \hat{z}(t) = 0, \hat{L}(t) = 0$.

We obtain from (5.4) that $\hat{y}(a, t) = 0$ for $0 \leq a < t$. For $a \geq t$, we have

$$\|\hat{y}(a, t)\|_{L^1} = \left\| y_0(a-t)\frac{\pi(a)}{\pi(a-t)} \right\|_{L^1} \leq e^{-\mu_0 t} \|y_0\|_{L^1},$$

which yields $\lim_{t \rightarrow +\infty} \hat{y}(a, t) = 0$. By comparison principle, it follows that $\lim_{t \rightarrow +\infty} y(a, t) = 0$ and $v(t) = 0, z(t) = 0$ as t tends to infinity. We obtain from the first equation of system (1.3) that $\lim_{t \rightarrow +\infty} x(t) = \Lambda/d$. This completes the proof.

Using a similar argument as that in the proof of Theorem 5.1, we have the following result.

Theorem 5.2. *The CTL-inactivated infection steady state $E_1(x_1, y_1(a), v_1, 0)$ is globally asymptotically stable for the semi-flow $\{\Phi(t)\}_{t \geq 0}$ restricted to $\partial\mathcal{Y}$.*

Theorem 5.3. *If $\mathcal{R}_1 > 1$, then the semi-flow $\{\Phi(t)\}_{t \geq 0}$ generated by system (1.3) is uniformly persistent with respect to the pair $(\mathcal{Y}, \partial\mathcal{Y})$; that is, there exists an $\varepsilon > 0$ such that $\liminf_{t \rightarrow +\infty} \|\Phi(t, x)\|_{\mathcal{X}} \geq \varepsilon$ for $x \in \mathcal{Y}$. Furthermore, there is a compact subset $\mathcal{A}_0 \subset \mathcal{Y}$ which is a global attractor for $\{\Phi(t)\}_{t \geq 0}$ in \mathcal{Y} .*

Proof. By Theorems 5.1 and 5.2, we see that the infection-free steady state $E_0(A/\mu, 0, 0, 0)$ and the CTL-inactivated infection steady state $E_1(x_1, y_1(a), v_1, 0)$ are globally asymptotically stable in $\partial\mathcal{Y}$. Hence, applying Theorem 4.2 in [35], in the following, we verify that

$$W^s(E_i) \cap \mathcal{Y} = \emptyset \quad (i = 0, 1),$$

where

$$W^s(E_0) = \{x \in \mathcal{Y} : \lim_{t \rightarrow +\infty} \Phi(t, x) = E_0\}, \quad W^s(E_1) = \{x \in \mathcal{Y} : \lim_{t \rightarrow +\infty} \Phi(t, x) = E_1\}.$$

Here, we only show $W^s(E_1) \cap \mathcal{Y} = \emptyset$ holds since the proof of $W^s(E_0) \cap \mathcal{Y} = \emptyset$ is simple. Assume $W^s(E_1) \cap \mathcal{Y} \neq \emptyset$. Then there exists a solution $w \in \mathcal{Y}$ such that $\Phi(t, w) \rightarrow E_1$ as $t \rightarrow \infty$. In this case, one can find a sequence $\{w_n\} \subset \mathcal{Y}$ such that

$$\|\Phi(t, w_n) - \bar{w}\|_{\mathcal{X}} < \frac{1}{n}, \quad t \geq 0,$$

where $\bar{w} = (x_1, y_1(a), v_1, 0)$.

Denote $\Phi(t, w_n) = (x_n(t), y_n(\cdot, t), v_n(t), z_n(t))$ and $w_n = (x_n(0), y_n(\cdot, 0), v_n(0), z_n(0))$. Since $\mathcal{R}_1 > 1$, we can choose n sufficiently large satisfying $x_1 - \frac{1}{n} > 0$ and

$$\left(\frac{du \int_0^\infty c(a)\phi_1(a)da}{(\alpha d + \beta) \int_0^\infty k(a)\phi_1(a)da} + b \right) (\mathcal{R}_1 - 1) > \frac{1}{n} \int_0^\infty c(a)da \quad (5.8)$$

For such an $n > 0$, there exists a $T_1 > 0$ such that for $t > T_1$,

$$\begin{aligned} x_1 - \frac{1}{n} < x_n(t) < x_1 + \frac{1}{n}, \quad y_1(a) - \frac{1}{n} \leq y_n(\cdot, t) \leq y_1(a) + \frac{1}{n}, \\ v_1 - \frac{1}{n} \leq v_n(t) \leq v_1 + \frac{1}{n}, \quad 0 \leq z_n(t) \leq \frac{1}{n}. \end{aligned} \quad (5.9)$$

Consider the following auxiliary system

$$\begin{aligned} \frac{\partial \tilde{y}(a, t)}{\partial t} + \frac{\partial \tilde{y}(a, t)}{\partial a} &= -\mu(a)\tilde{y}(a, t) - p(a)\tilde{y}(a, t)\frac{1}{n}, \\ \dot{\tilde{v}}(t) &= \int_0^\infty k(a)\tilde{y}(a, t)da - u\tilde{v}(t), \\ \dot{\tilde{z}}(t) &= \tilde{z}(t) \left(\int_0^\infty c(a) \left(y_1(a) - \frac{1}{n} \right) da - b \right), \\ \tilde{y}(0, t) &= \frac{\beta \left(x_1 - \frac{1}{n} \right) \tilde{v}(t)}{1 + \frac{\alpha}{n}}. \end{aligned} \quad (5.10)$$

It is easy to show that if $\mathcal{R}_1 > 1$, system (5.10) has a unique steady state $E_0(0, 0, 0)$.

Looking for solutions of system (5.10) of the form $\tilde{y}(a, t) = \tilde{y}_1(a)e^{\lambda t}$, $\tilde{v}(t) = \tilde{v}_1 e^{\lambda t}$, $\tilde{z}(t) = \tilde{z}_1 e^{\lambda t}$, where the function $\tilde{y}_1(a)$ and the constants \tilde{v}_1, \tilde{z}_1 will be determined later, we obtain the following linear eigenvalue problem:

$$\begin{aligned} \tilde{y}_1'(a) &= -(\lambda + \mu(a))\tilde{y}_1(a) - p(a)\tilde{y}_1(a)\frac{1}{n}, \\ \int_0^\infty k(a)\tilde{y}_1(a)da &= (\lambda + \mu)\tilde{v}_1, \\ \lambda &= \int_0^\infty c(a) \left(y_1(a) - \frac{1}{n} \right) da - b, \\ \tilde{y}_1(0) &= \frac{\beta \left(x_0 - \frac{1}{n} \right) \tilde{v}_1}{1 + \frac{\alpha}{n}}. \end{aligned} \quad (5.11)$$

We derive from the third equation of (5.11) that

$$\lambda = \left(\frac{du \int_0^\infty c(a)\phi_1(a)da}{(\alpha d + \beta) \int_0^\infty k(a)\phi_1(a)da} + b \right) (\mathcal{R}_1 - 1) - \frac{1}{n} \int_0^\infty c(a)da. \quad (5.12)$$

Clearly, if $\mathcal{R}_1 > 1$, Eq (5.11) has at least one positive root λ_0 , which yields the solution $(\tilde{y}(\cdot, t), \tilde{v}(t), \tilde{z}(t))$ of system (5.10) is unbounded. By comparison principle, the solution $\Phi(t, y_n)$ of system (1.3) is unbounded, which contradicts Proposition 5.1. Therefore, the semi-flow $\{\Phi(t)\}_{t \geq 0}$ generated by system (1.3) is uniformly persistent. Furthermore, there is a compact subset $\mathcal{A}_0 \subset \mathcal{Y}$ which is a global attractor for $\{\Phi(t)\}_{t \geq 0}$ in \mathcal{Y} . This completes the proof.

6. Global stability

In this section, we discuss the global stability of each of feasible steady states of system (1.3). The strategy of proofs is to use suitable Lyapunov functionals and LaSalle's invariance principle. Further, we employ a Volterra type functional defined by $G(x) = x - 1 - \ln x$ in [37], which is positive and attains minimum value 0 at $x = 1$.

We first give a result on the global stability of the infection-free steady state $E_0(\Lambda/d, 0, 0, 0)$ of system (1.3).

Theorem 6.1. *If $\mathcal{R}_0 < 1$, the infection-free steady state $E_0(\Lambda/d, 0, 0, 0)$ of system (1.3) is globally asymptotically stable.*

Proof. Let $(x(t), y(a, t), v(t), z(t))$ be any positive solution of system (1.3) with the boundary condition (1.4). Denote $x_0 = \Lambda/d$.

Define

$$V_1(t) = x(t) - x_0 - x_0 \ln \frac{x(t)}{x_0} + \int_0^\infty F_1(a)y(a, t)da + k_1 v(t), \quad (6.1)$$

where the positive constant k_1 and the nonnegative kernel function $F_1(a)$ will be determined later.

Calculating the derivative of $V_1(t)$ along positive solutions of system (1.3), it follows that

$$\begin{aligned} \frac{d}{dt} V_1(t) &= \left(1 - \frac{x_0}{x(t)} \right) \left[\Lambda - dx(t) - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} \right] \\ &\quad + \int_0^\infty F_1(a) \frac{\partial y(a, t)}{\partial t} da + k_1 \left[\int_0^\infty k(a)y(a, t)da - uv(t) \right]. \end{aligned} \quad (6.2)$$

On substituting $\Lambda = dx_0$, $\frac{\partial y(a,t)}{\partial t} = -(\mu(a) + p(a)z(t))y(a,t) - \frac{\partial y(a,t)}{\partial a}$ into Eq (6.2), one obtains

$$\begin{aligned} \frac{d}{dt}V_1(t) &= \left(1 - \frac{x_0}{x(t)}\right)[-d(x(t) - x_0)] \\ &\quad - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} + \frac{\beta x_0 v(t)}{1 + \alpha v(t)} - \int_0^\infty F_1(a) \frac{\partial y(a,t)}{\partial a} da \\ &\quad - \int_0^\infty F_1(a)\mu(a)y(a,t)da + k_1 \int_0^\infty k(a)y(a,t)da - k_1 uv(t) \\ &= \left(1 - \frac{x_0}{x(t)}\right)[-d(x(t) - x_0)] \\ &\quad - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} + \frac{\beta x_0 v(t)}{1 + \alpha v(t)} - F_1(a)y(a,t)|_0^\infty + \int_0^\infty F_1'(a)y(a,t)da \\ &\quad - \int_0^\infty F_1(a)\mu(a)y(a,t)da + k_1 \int_0^\infty k(a)y(a,t)da - k_1 uv(t). \end{aligned} \quad (6.3)$$

Choose

$$k_1 = \frac{\beta x_0}{u}, \quad F_1(a) = k_1 \int_a^\infty k(u)e^{-\int_a^u \mu(s)ds} du,$$

Then, we have

$$\begin{aligned} F_1(0) &= \frac{\beta x_0}{u} \int_0^\infty k(a)e^{-\int_0^a \mu(s)ds} da = \mathcal{R}_0, \\ F_1'(a) &= -\frac{\beta x_0}{u}k(a) + \mu(a)F_1(a), \quad \lim_{a \rightarrow +\infty} f_1(a) = 0. \end{aligned} \quad (6.4)$$

We therefore obtain from (6.3)–(6.4) that

$$\begin{aligned} \frac{d}{dt}V_1(t) &= \left(1 - \frac{x_0}{x(t)}\right)[-d(x(t) - x_0)] + (\mathcal{R}_0 - 1) \frac{\beta x(t)v(t)}{1 + \alpha v(t)} - \frac{\alpha \beta x_0 v(t)^2}{1 + \alpha v(t)} \\ &\quad - z(t) \int_0^\infty F_1(a)p(a)y(a,t)da. \end{aligned} \quad (6.5)$$

Clearly, if $\mathcal{R}_0 < 1$, we obtain from (6.5) that $V_1'(t) \leq 0$ and $V_1'(t) = 0$ implies that $x = x_0$, $y(a,t) = 0$ and $v = 0$. Hence, the largest invariant subset of $\{V_1'(t) = 0\}$ is the singleton $(x_0, 0, 0)$. Further, for $\varepsilon > 0$ sufficiently small satisfying $\int_0^\infty c(a)\varepsilon da - b < 0$, there is a $T > 0$, such that if $t > T$, $y(a,t) < \varepsilon$. It therefore follows from the fourth equation of system (1.3) that for $t > T$,

$$\dot{z}(t) \leq \left(\int_0^\infty c(a)\varepsilon da - b \right) z(t).$$

By comparison, we derive that

$$\lim_{t \rightarrow +\infty} z(t) = 0.$$

From Section 4, we see that if $\mathcal{R}_0 < 1$, E_0 is locally asymptotically stable. Accordingly, the global asymptotic stability of E_0 of system (1.3) follows from LaSalle's invariance principle. This completes the proof.

In the following, we establish the global asymptotic stability of the CTL-inactivated infection steady state $E_1(x_1, y_1(a), v_1, 0)$ and the global attractivity of the CTL-activated infection steady state $E^*(x^*, y^*(a), v^*, z^*)$ of system (1.3), respectively.

Denote

$$D_0 = \left\{ (x^0, y_0, v^0, z^0) \in \mathcal{X} \mid \int_0^\infty k(a)y_0(a)da > 0, \int_0^\infty c(a)y_0(a)da > 0 \right\}.$$

In order to guarantee the Lyapunov functional in proving the global stability of E_1 and E^* is well-defined in infinite dimension, we make the following assumption:

$$(H4) \quad x^0 > 0, v^0 > 0, z^0 > 0, \int_0^\infty |\ln y_0(a)| da < +\infty.$$

We now define a positive function

$$F_2(a) = \frac{\beta x_1}{u(1 + \alpha v_1)} \int_a^\infty k(u) e^{-\int_a^u \mu(s) ds} du. \quad (6.6)$$

Then, we have

$$F_2(0) = \frac{\beta x_1}{u(1 + \alpha v_1)} \int_0^\infty k(a) e^{-\int_0^a \mu(s) ds} da = 1, \quad (6.7)$$

and

$$\lim_{a \rightarrow \infty} F_2(a) = 0, \quad F_2'(a) = -\frac{\beta x_1}{u(1 + \alpha v_1)} k(a) + \mu(a) F_2(a). \quad (6.8)$$

Theorem 6.2. *Assume there exists a positive constant k_2 satisfying $F_2(a)p(a) = k_2c(a)$. If (H4) holds, then the CTL-inactivated infection steady state $E_1(x_1, y_1(a), v_1, 0)$ of system (1.3) is globally asymptotically stable if $\mathcal{R}_1 < 1 < \mathcal{R}_0$.*

Proof. Let $(x(t), y(a, t), v(t), z(t))$ be any positive solution of system (1.3) with the boundary condition (1.4).

Define

$$V_2(t) = x_1 G\left(\frac{x(t)}{x_1}\right) + \int_0^\infty F_2(a)y_1(a)G\left(\frac{y(a, t)}{y_1(a)}\right) da + \frac{\beta x_1}{u(1 + \alpha v_1)} v_1 G\left(\frac{v(t)}{v_1}\right) + k_2 z(t). \quad (6.9)$$

Using a similar argument as that in the proof of Lemmas 7.1 and 7.2 in [27], one can show that all integrals involved in $V_2(t)$ are finite.

Calculating the derivative of $V_2(t)$ along positive solutions of system (1.3), it follows that

$$\begin{aligned}
 \frac{d}{dt}V_2(t) &= \left(1 - \frac{x_1}{x(t)}\right) \left[\Lambda - dx(t) - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} \right] \\
 &\quad + \int_0^\infty F_2(a)y_1(a) \frac{\partial}{\partial t} G\left(\frac{y(a,t)}{y_1(a)}\right) da \\
 &\quad + \frac{\beta x_1}{u(1 + \alpha v_1)} v_1 \left(1 - \frac{v_1}{v(t)}\right) \left[\int_0^\infty k(a)y(a,t) da - uv(t) \right] \\
 &\quad + k_2 \left[\int_0^\infty c(a)y(a,t) da - b \right] z(t) \\
 &= \left(1 - \frac{x_1}{x(t)}\right) \left[\Lambda - dx(t) - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} \right] \\
 &\quad + \int_0^\infty F_2(a) \left(1 - \frac{y_1(a)}{y(a,t)}\right) \frac{\partial y(a,t)}{\partial t} da \\
 &\quad + \frac{\beta x_1}{u(1 + \alpha v_1)} \left(1 - \frac{v_1}{v(t)}\right) \left[\int_0^\infty k(a)y(a,t) da - uv(t) \right] \\
 &\quad + k_2 \left[\int_0^\infty c(a)y(a,t) da - b \right] z(t).
 \end{aligned} \tag{6.10}$$

On substituting $\Lambda = dx_1 + \beta x_1 v_1 / (1 + \alpha v_1)$ and $\frac{\partial y(a,t)}{\partial t} = -(\mu(a) + p(a)z(t))y(a,t) - \frac{\partial y(a,t)}{\partial a}$ into Eq (6.10), one obtains

$$\begin{aligned}
 \frac{d}{dt}V_2(t) &= \left(1 - \frac{x_1}{x(t)}\right) \left[-d(x(t) - x_1) + \frac{\beta x_1 v_1}{1 + \alpha v_1} \right] - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} + \frac{\beta x_1 v(t)}{1 + \alpha v(t)} \\
 &\quad - \int_0^\infty F_2(a) \left(1 - \frac{y_1(a)}{y(a,t)}\right) \left[\frac{\partial y(a,t)}{\partial a} + (\mu(a) + p(a)z(t))y(a,t) \right] da \\
 &\quad + \frac{\beta x_1}{u(1 + \alpha v_1)} \left[\int_0^\infty k(a)y(a,t) da - uv(t) - \frac{v_1}{v(t)} \int_0^\infty k(a)y(a,t) da + uv_1 \right] \\
 &\quad + k_2 \left[\int_0^\infty c(a)y(a,t) da - b \right] z(t).
 \end{aligned} \tag{6.11}$$

A direct calculation shows that

$$y_1(a) \frac{\partial}{\partial a} G\left(\frac{y(a,t)}{y_1(a)}\right) = \left(1 - \frac{y_1(a)}{y(a,t)}\right) \left(\frac{\partial y(a,t)}{\partial a} + \mu(a)y(a,t) \right). \tag{6.12}$$

On substituting Eq (6.12) into Eq (6.11), we have

$$\begin{aligned}
 \frac{d}{dt}V_2(t) &= \left(1 - \frac{x_1}{x(t)}\right) \left[-d(x(t) - x_1) + \frac{\beta x_1 v_1}{1 + \alpha v_1} \right] - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} + \frac{\beta x_1 v(t)}{1 + \alpha v(t)} \\
 &\quad - \int_0^\infty F_2(a)y_1(a) \frac{\partial}{\partial a} G\left(\frac{y(a,t)}{y_1(a)}\right) da - \int_0^\infty F_2(a) \left(1 - \frac{y_1(a)}{y(a,t)}\right) p(a)y(a,t)z(t) da \\
 &\quad + \frac{\beta x_1}{u(1 + \alpha v_1)} \left[\int_0^\infty k(a)y(a,t) da - uv(t) - \frac{v_1}{v(t)} \int_0^\infty k(a)y(a,t) da + uv_1 \right] \\
 &\quad + k_2 \left[\int_0^\infty c(a)y(a,t) da - b \right] z(t).
 \end{aligned} \tag{6.13}$$

Using integration by parts, it follows from Eq (6.13) that

$$\begin{aligned} \frac{d}{dt}V_2(t) &= \left(1 - \frac{x_1}{x(t)}\right) \left[-d(x(t) - x_1) + \frac{\beta x_1 v_1}{1 + \alpha v_1}\right] - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} + \frac{\beta x_1 v(t)}{1 + \alpha v(t)} \\ &\quad - F_2(a)y_1(a)G\left(\frac{y(a,t)}{y_1(a)}\right)\Big|_0^\infty + \int_0^\infty G\left(\frac{y(a,t)}{y_1(a)}\right)[F_2'(a)y_1(a) + F_2(a)y_1'(a)]da \\ &\quad - \int_0^\infty F_2(a)\left(1 - \frac{y_1(a)}{y(a,t)}\right)p(a)y(a,t)z(t)da \\ &\quad + \frac{\beta x_1}{u(1 + \alpha v_1)} \left[\int_0^\infty k(a)y(a,t)da - uv(t) - \frac{v_1}{v(t)} \int_0^\infty k(a)y(a,t)da + uv_1 \right] \\ &\quad + k_2 z(t) \left[\int_0^\infty c(a)y(a,t)da - b \right]. \end{aligned} \quad (6.14)$$

On substituting Eqs (6.7)–(6.8) into Eq (6.14), and noting that $y_1'(a) = -\mu(a)y_1(a)$, $y_1(0) = \beta x_1 v_1 / (1 + \alpha v_1)$ and $y(0, t) = \beta x(t)v(t) / (1 + \alpha v(t))$, we obtain from Eq (6.14) that

$$\begin{aligned} \frac{d}{dt}V_2(t) &= -d \frac{(x(t) - x_1)^2}{x(t)} - \frac{\beta x_1 v_1}{1 + \alpha v_1} \left(\frac{x_1}{x(t)} - 1 \right) + \frac{\beta x_1 v(t)}{1 + \alpha v(t)} \\ &\quad - \frac{\beta x_1 v_1}{1 + \alpha v_1} - \frac{\beta x_1 v_1}{1 + \alpha v_1} \ln \frac{x(t)v(t)(1 + \alpha v_1)}{x_1 v_1 (1 + \alpha v(t))} \\ &\quad - \frac{\beta x_1}{u(1 + \alpha v_1)} \int_0^\infty k(a)y_1(a)G\left(\frac{y(a,t)}{y_1(a)}\right)da \\ &\quad + \frac{\beta x_1}{u(1 + \alpha v_1)} \left[\int_0^\infty k(a)y(a,t)da - uv(t) - \frac{v_1}{v(t)} \int_0^\infty k(a)y(a,t)da + uv_1 \right] \\ &\quad - \int_0^\infty F_2(a)\left(1 - \frac{y_1(a)}{y(a,t)}\right)p(a)y(a,t)z(t)da + k_2 z(t) \left[\int_0^\infty c(a)y(a,t)da - b \right]. \end{aligned} \quad (6.15)$$

Noting that $\frac{\beta x_1}{u(1 + \alpha v_1)} \int_0^\infty k(a)y_1(a)da = \frac{\beta x_1}{u(1 + \alpha v_1)} uv_1 = \frac{\beta x_1 v_1}{1 + \alpha v_1}$, we have from Eq (6.15) that

$$\begin{aligned} \frac{d}{dt}V_2(t) &= -d \frac{(x(t) - x_1)^2}{x(t)} - \frac{\alpha \beta x_1 (v(t) - v_1)^2}{(1 + \alpha v(t))(1 + \alpha v_1)^2} \\ &\quad - \frac{\beta x_1 v_1}{1 + \alpha v_1} G\left(\frac{x_1}{x(t)}\right) - \frac{\beta x_1 v_1}{1 + \alpha v_1} G\left(\frac{1 + \alpha v(t)}{1 + \alpha v_1}\right) \\ &\quad - \frac{\beta x_1}{u(1 + \alpha v_1)} \int_0^\infty k(a)y_1(a)G\left(\frac{v_1 y(a,t)}{v(t)y_1(a)}\right)da \\ &\quad + k_2 \left(\frac{du \int_0^\infty c(a)\phi_1(a)da}{(\alpha d + \beta) \int_0^\infty k(a)\phi_1(a)da} + b \right) (\mathcal{R}_1 - 1)z(t). \end{aligned} \quad (6.16)$$

Since the function $G(x) = x - 1 - \ln x \geq 0$ for all $x > 0$ and $G(x) = 0$ holds iff $x = 1$. Hence, $V_2'(t) \leq 0$ holds if $\mathcal{R}_1 < 1$. It is readily seen from (6.16) that $V_2'(t) = 0$ if and only if

$$x(t) = x_1, \quad \frac{y(a,t)v_1}{y_1(a)v(t)} = 1, \quad \frac{1 + \alpha v(t)}{1 + \alpha v_1} = 1, \quad (6.17)$$

for all $a \geq 0$. It is easy to verify that the largest invariant subset of $\{V_2'(t) = 0\}$ is the singleton E_1 . By Theorem 4.2, we see that if $\mathcal{R}_1 < 1 < \mathcal{R}_0$, E_1 is locally asymptotically stable. Therefore, using LaSalle's invariance principle, we see that if $\mathcal{R}_1 < 1 < \mathcal{R}_0$ and (H4) hold, the global asymptotic stability of E_1 follows. This completes the proof.

In the following, we define a positive function

$$F_3(a) = \frac{\beta x^*}{u(1 + \alpha v^*)} \int_a^\infty k(u) e^{-\int_a^u (\mu(s) + p(s)z^*) ds} du. \quad (6.18)$$

It is easy to show that

$$F_3(0) = \frac{\beta x^*}{u(1 + \alpha v^*)} \int_0^\infty k(a) e^{-\int_0^a (\mu(s) + p(s)z^*) ds} da = 1, \quad (6.19)$$

and

$$\lim_{a \rightarrow \infty} F_3(a) = 0, \quad F_3'(a) = -\frac{\beta x^*}{u(1 + \alpha v^*)} k(a) + (\mu(a) + p(a)z^*) F_3(a). \quad (6.20)$$

Theorem 6.3. *Assume there exists a positive constant k_3 satisfying $F_3(a)p(a) = k_3c(a)$. If (H4) holds, then the CTL-activated infection steady state $E^*(x^*, y^*(a), v^*, z^*)$ of system (1.3) is globally attractive if $\mathcal{R}_1 > 1$.*

Proof. Let $(x(t), y(a, t), v(t), z(t))$ be any positive solution of system (1.3) with the boundary condition (1.4).

Define

$$\begin{aligned} V_3(t) = & x^* G\left(\frac{x(t)}{x^*}\right) + \int_0^\infty F_3(a) y^*(a) G\left(\frac{y(a, t)}{y^*(a)}\right) da \\ & + \frac{\beta x^*}{u(1 + \alpha v^*)} v^* G\left(\frac{v(t)}{v^*}\right) + k_3 z^* G\left(\frac{z(t)}{z^*}\right). \end{aligned} \quad (6.21)$$

Using a similar argument as that in the proof of Lemmas 7.1 and 7.2 in [27], one can show that all integrals involved in $V_3(t)$ are finite.

Calculating the derivative of $V_3(t)$ along positive solutions of system (1.3), it follows that

$$\begin{aligned} \frac{d}{dt} V_3(t) = & \left(1 - \frac{x^*}{x(t)}\right) \left[\Lambda - dx(t) - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} \right] \\ & + \int_0^\infty F_3(a) \left(1 - \frac{y^*(a)}{y(a, t)}\right) \frac{\partial y(a, t)}{\partial t} da \\ & + \frac{\beta x^*}{u(1 + \alpha v^*)} \left(1 - \frac{v^*}{v(t)}\right) \left[\int_0^\infty k(a) y(a, t) da - uv(t) \right] \\ & + k_3 \left(1 - \frac{z^*}{z(t)}\right) \left[\int_0^\infty c(a) y(a, t) da - b \right] z(t). \end{aligned} \quad (6.22)$$

On substituting $\Lambda = dx^* + \beta x^* v^* / (1 + \alpha v^*)$ and $\frac{\partial y(a, t)}{\partial t} = -(\mu(a) + p(a)z(t))y(a, t) - \frac{\partial y(a, t)}{\partial a}$ into Eq (6.22),

one obtains

$$\begin{aligned} \frac{d}{dt} V_3(t) = & \left(1 - \frac{x^*}{x(t)}\right) \left[-d(x(t) - x^*) + \frac{\beta x^* v^*}{1 + \alpha v^*} \right] - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} + \frac{\beta x^* v(t)}{1 + \alpha v(t)} \\ & - \int_0^\infty F_3(a) \left(1 - \frac{y^*(a)}{y(a,t)}\right) \left[\frac{\partial y(a,t)}{\partial a} + (\mu(a) + p(a)z(t))y(a,t) \right] da \\ & + \frac{\beta x^*}{u(1 + \alpha v^*)} \left[\int_0^\infty k(a)y(a,t)da - uv(t) - \frac{v^*}{v(t)} \int_0^\infty k(a)y(a,t)da + uv^* \right] \\ & + k_3 \left[z(t) \int_0^\infty c(a)y(a,t)da - bz(t) - z^* \int_0^\infty c(a)y(a,t)da + bz^* \right]. \end{aligned} \quad (6.23)$$

A direct calculation shows that

$$y^*(a) \frac{\partial}{\partial a} G \left(\frac{y(a,t)}{y^*(a)} \right) = \left(1 - \frac{y^*(a)}{y(a,t)}\right) \left(\frac{\partial y(a,t)}{\partial a} + (\mu(a) + p(a)z^*)y(a,t) \right). \quad (6.24)$$

On substituting Eq (6.24) into Eq (6.23), we get

$$\begin{aligned} \frac{d}{dt} V_3(t) = & \left(1 - \frac{x^*}{x(t)}\right) \left[-d(x(t) - x^*) + \frac{\beta x^* v^*}{1 + \alpha v^*} \right] - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} + \frac{\beta x^* v(t)}{1 + \alpha v(t)} \\ & - \int_0^\infty F_3(a) y^*(a) \frac{\partial}{\partial a} G \left(\frac{y(a,t)}{y^*(a)} \right) da \\ & - \int_0^\infty F_3(a) \left(1 - \frac{y^*(a)}{y(a,t)}\right) p(a)y(a,t)(z(t) - z^*) da \\ & + \frac{\beta x^*}{u(1 + \alpha v^*)} \left[\int_0^\infty k(a)y(a,t)da - uv(t) - \frac{v^*}{v(t)} \int_0^\infty k(a)y(a,t)da + uv^* \right] \\ & + k_3 \left[z(t) \int_0^\infty c(a)y(a,t)da - bz(t) - z^* \int_0^\infty c(a)y(a,t)da + bz^* \right]. \end{aligned} \quad (6.25)$$

Using integration by parts, it follows from Eq (6.25) that

$$\begin{aligned} \frac{d}{dt} V_3(t) = & -\frac{d}{x(t)} (x(t) - x^*)^2 - \frac{\beta x^* v^*}{1 + \alpha v^*} \left(\frac{x^*}{x(t)} - 1 - \ln \frac{x^*}{x(t)} \right) \\ & - \frac{\beta x^* v^*}{1 + \alpha v^*} \ln \frac{x^*}{x(t)} - \frac{\beta x(t)v(t)}{1 + \alpha v(t)} + \frac{\beta x^* v(t)}{1 + \alpha v(t)} \\ & - F_3(a) y^*(a) G \left(\frac{y(a,t)}{y^*(a)} \right) \Big|_0^\infty + \int_0^\infty G \left(\frac{y(a,t)}{y^*(a)} \right) [F_3'(a) y^*(a) + F_3(a) y^{*'}(a)] da \\ & - \int_0^\infty F_3(a) \left(1 - \frac{y^*(a)}{y(a,t)}\right) p(a)y(a,t)(z(t) - z^*) da \\ & + \frac{\beta x^*}{u(1 + \alpha v^*)} \left[\int_0^\infty k(a)y(a,t)da - uv(t) - \frac{v^*}{v(t)} \int_0^\infty k(a)y(a,t)da + uv^* \right] \\ & + k_3 \left[z(t) \int_0^\infty c(a)y(a,t)da - bz(t) - z^* \int_0^\infty c(a)y(a,t)da + bz^* \right]. \end{aligned} \quad (6.26)$$

On substituting Eqs (6.19)–(6.20) into Eq (6.26), and noting that

$$y^*(a) = -(\mu(a) + p(a)z^*)y^*(a), y^*(0) = \beta x^* v^* / (1 + \alpha v^*)$$

and

$$\frac{\beta x^*}{u(1 + \alpha v^*)} \int_0^\infty k(a)y^*(a)da = \frac{\beta x^*}{u(1 + \alpha v^*)} uv^* = \frac{\beta x^* v^*}{1 + \alpha v^*},$$

we obtain from Eq (6.26) that

$$\begin{aligned} \frac{d}{dt} V_3(t) = & -d \frac{(x(t) - x^*)^2}{x(t)} - \frac{\alpha \beta x^* (v(t) - v^*)^2}{(1 + \alpha v^*)^2 (1 + \alpha v(t))} \\ & - \frac{\beta x^* v^*}{1 + \alpha v^*} G\left(\frac{x^*}{x(t)}\right) - \frac{\beta x^* v^*}{1 + \alpha v^*} G\left(\frac{1 + \alpha v(t)}{1 + \alpha v^*}\right) \\ & - k_3 \int_0^\infty k(a)y^*(a)G\left(\frac{y(a,t)v^*}{y^*(a)v(t)}\right) da. \end{aligned} \quad (6.27)$$

Since the function $G(x) = x - 1 - \ln x \geq 0$ for all $x > 0$ and $G(x) = 0$ holds iff $x = 1$. Hence, $V_3'(t) \leq 0$ holds if $\mathcal{R}_1 > 1$. It is readily seen from (6.27) that $V_3'(t) = 0$ if and only if

$$x(t) = x^*, v(t) = v^*, \frac{y(a,t)v^*}{y^*(a)v(t)} = 1, \frac{1 + \alpha v(t)}{1 + \alpha v^*} = 1, \quad (6.28)$$

for all $a \geq 0$. We now look for the invariant subset \mathcal{M} within the set

$$\mathcal{M} = \{(x, y, v) : x(t) = x^*, y(a, t) = y^*(a), v(t) = v^*\}.$$

Because $x(t) = x^*, y(a, t) = y^*(a)$ and $v(t) = v^*$ on \mathcal{M} and consequently, it follows from the second equation of system (1.3) that

$$\frac{d}{da} y^*(a) = -\mu(a)y^*(a) - p(a)y^*(a)z(t),$$

which yields $z(t) = z^*$. It is easy to verify that the largest invariant subset of $\{V_3'(t) = 0\}$ is the singleton E^* . Therefore, using LaSalle's invariance principle, we see that if $\mathcal{R}_1 > 1$ and (H4) hold, the global attractivity of E^* follows. This completes the proof.

7. Numerical simulations

In this section, we give some numerical examples for system (1.3) to illustrate the theoretical results in Sections 3 and 4. Based on the works of [28–33], parameter values of system (1.3) are summarized in Table 2. In the following, we will use the finite difference method [34] for all numerical simulations. Further, to ensure the precision of numerical simulations, time- and age-steps are both set as 0.05.

Table 2. Parameter values for the age-structured HIV-1 model (1.3).

| Parameter | Symbol | Case 1 | Case2 | Case 3 | Source |
|--|---|----------------------|----------------------|----------------------|---------|
| Recruitment rate of healthy T cells | Λ ($\text{ml}^{-1}\text{day}^{-1}$) | 0.8×10^6 | 0.98×10^6 | 1.8×10^6 | Assumed |
| Death rate of uninfected cells | d (day^{-1}) | 0.01 | 0.01 | 0.01 | [29] |
| Infection rate | β (ml day^{-1}) | 1.3×10^{-8} | 1.3×10^{-8} | 1.3×10^{-8} | [28] |
| Saturation constant | α | 0.00015 | 0.00015 | 0.00015 | [33] |
| Death rate of infected cells | μ_m (day^{-1}) | 0.7 | 0.7 | 0.7 | [30] |
| Clearance rate of virions | u (day^{-1}) | 23 | 23 | 23 | [31] |
| Death rate of CTL cells | b (day^{-1}) | 0.5 | 0.5 | 0.5 | [32] |
| Killing rate of infected cells | p_m (day^{-1}) | 0.00094 | 0.00094 | 0.00094 | [32] |
| Viral production rate of infected cells | k_m (day^{-1}) | 11.349 | 11.349 | 11.349 | [32] |
| Proliferate rate of virus-specific CTL cells | c_m (day^{-1}) | 0.003 | 0.003 | 0.003 | Assumed |

As argued by Markowitz et al. [30], the faster rate of loss of virus-producing cells shows that the generation time for HIV-1 in vivo is correspondingly shorter, ~ 2.0 days, which is obtained by summing up some factors, such as the eclipse time of ~ 1.0 day. This value indicates that HIV-1 typically undergoes 180 generations per year in an infected person. Thus, the death rate of infected cells μ_m in Table 2 is set as 0.7 day^{-1} .

7.1. Dynamical behaviors of system (1.3)

When the viruses invade through cytomembrane, infected cells cannot die immediately, due to that it takes some time for viruses to replicate, transcribe and translate. For this reason, we assume that the death rate of infected cells increases from 0 to a peak value μ_m with the infection age. The age-dependent per capita death rate is set as

$$\mu(a) = \begin{cases} \mu_m \sin(0.1\pi a), & a < a_0, \\ \mu_m, & a \geq a_0, \end{cases} \quad (7.1)$$

where a_0 denotes the mean value of the time for viruses to replicate, transcribe and translate (in this section, a_0 is set as 5 day). Further, the maturing rate of new T cells and the kill ratio by T cells are selected as follows:

$$c(a) = \begin{cases} c_m \sin(0.1\pi a), & a < a_0, \\ c_m, & a \geq a_0, \end{cases} \quad p(a) = \begin{cases} p_m \sin(0.1\pi a), & a < a_0, \\ p_m, & a \geq a_0, \end{cases} \quad (7.2)$$

where c_m and p_m are the peak levels of $c(a)$ and $p(a)$, respectively. As for the viral production rate of infected cells, it keeps at 0 for a short time a_1 , and then increases from 0 to a peak value k_m . Based on the works of [9, 11], the specific function is set as follows

$$k(a) = \begin{cases} 0, & a < a_1, \\ k_m(1 - e^{-\theta(a-a_1)}), & a \geq a_1, \end{cases} \quad (7.3)$$

where θ determines how quickly $k(a)$ reaches the saturation level k_m . For simplicity, we assume that $\theta = 1$ and $a_1 = 0.5$ day in the following numerical simulations.

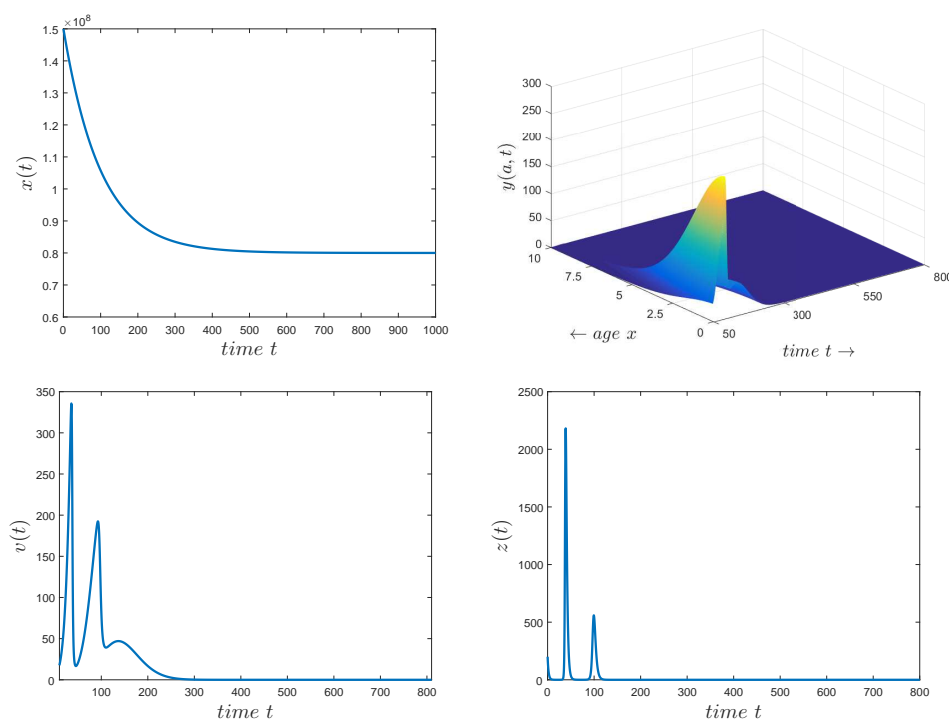


Figure 1. The temporal solution found by numerical integration of system (1.3) with the boundary condition (1.4) and the initial condition $x(0) = 1.5 \times 10^8$, $y(0, 0) = 100$, $v(0) = 200$, $z(0) = 200$, where $\mathcal{R}_0 = 0.9131 < 1$.

We first choose parameter values as in Case 1 of Table 2. Then we have the basic reproduction number $\mathcal{R}_0 = 0.8227 < 1$. By Theorem 4.1, we see that the infection-free steady state $E_0(80003167.69, 0, 0, 0)$ is locally asymptotically stable. Numerical simulation illustrates this fact (see Figure 1).

Next, we choose parameter values as in Case 2 of Table 2. By direct calculation, we get the basic reproduction number $\mathcal{R}_0 = 1.0079 > 1$ and the immune response reproduction number $\mathcal{R}_1 = 0.9939 < 1$. By Theorem 4.1, we see that in addition to the infection-free steady state $E_0(80003167.69, 0, 0, 0)$, system (1.3) has a CTL-inactivated infection steady state $E_1(97994001.72, 59.69\phi_1(a), 47.19, 0)$ which is locally asymptotically stable. Numerical simulation illustrates this fact (see Figure 2).

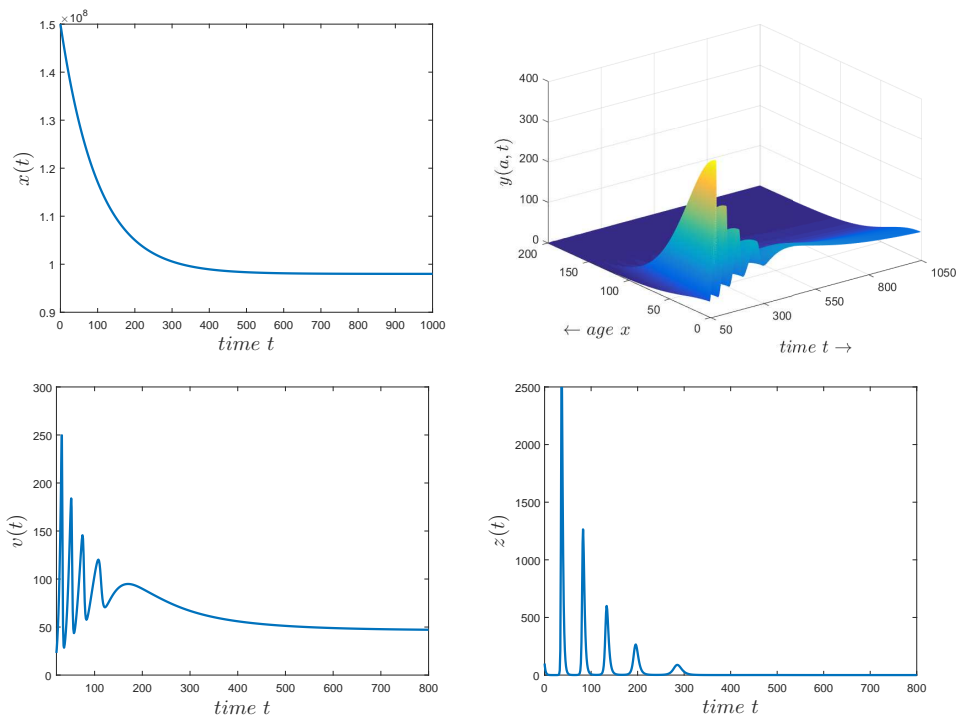


Figure 2. The temporal solutions found by numerical integration of system (1.3) with the boundary condition (1.4) and the initial condition $x(0) = 1.5 \times 10^8$, $y(0, 0) = 100$, $v(0) = 200$, $z(0) = 100$, where $\mathcal{R}_1 = 0.9939 < 1 < \mathcal{R}_0 = 1.0079$.

Remark 7.1. For system (1.3), a direct calculation shows that the characteristic equation of system (1.3) at the CTL-activated infection steady state E^* is of the form

$$\frac{\beta x^*}{(1 + \alpha v^*)^2} \frac{\int_0^\infty k(a) \phi_1(a) \phi_2(a, z^*) e^{-\lambda a} da}{\lambda + u} = 1 + \frac{\beta v^*}{1 + \alpha v^*} \frac{1}{\lambda + d} + \frac{\beta x^*}{(1 + \alpha v^*)^2} \frac{1}{\lambda + u} f(\lambda), \quad (7.4)$$

where

$$f(\lambda) = \frac{z^* \int_0^\infty \int_0^a k(a) p(s) y^*(s) \phi_1(a-s) \phi_2(a-s, z^*) e^{-\lambda(a-s)} ds da \int_0^\infty c(a) \phi_1(a) \phi_2(a, z^*) e^{-\lambda a} da}{\lambda + z^* \int_0^\infty \int_0^a c(a) p(s) y^*(s) \phi_1(a-s) \phi_2(a-s, z^*) e^{-\lambda(a-s)} ds da}.$$

We failed in studying the local asymptotic stability of E^* due to the complexity of Eq (7.4). In particular, we choose parameter values as in Case 3 of Table 2. By calculation, we have the immune-activated reproduction rate $\mathcal{R}_1 = 1.8256 > 1$. As can be seen from the discussion in Section 3, in addition to the infection-free steady state E_0 and the CTL-inactivated infection steady state E_1 , system (1.3) has a unique CTL-activated infection steady state $E^*(179981502.93, 216.65\phi_1(a) \phi_2(a, 638.84), 93.90, 638.84)$. Numerical simulation indicates that if $\mathcal{R}_1 > 1$, the CTL-activated infection steady state E^* is locally asymptotically stable in some special cases (see Figure 3).

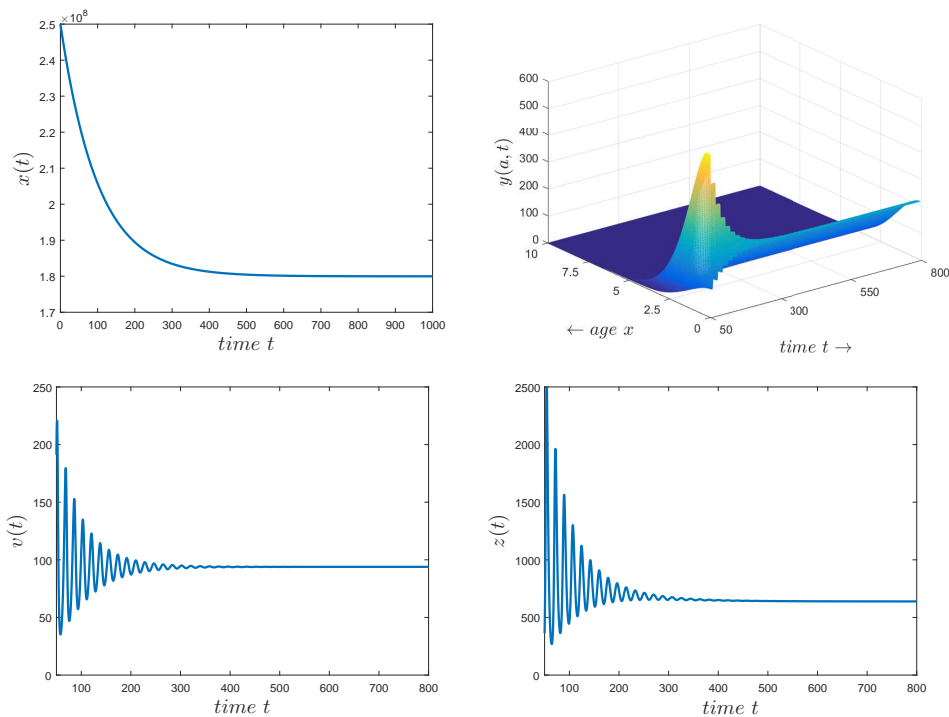


Figure 3. The temporal solutions found by numerical integration of system (1.3) with the boundary condition (1.4) and the initial condition $x(0) = 2.5 \times 10^8$, $y(0, 0) = 100$, $v(0) = 200$, $z(0) = 800$, where $\mathcal{R}_1 = 1.8256 > 1$.

7.2. The effects of CTL response

In order to investigate the effects of CTL immune response, we carry out the following numerical simulations. For convenience, parameter values are chosen as in Table 2. From Figure 4, it is clear that the concentrations of infected cells and free virions with CTL immune response is obviously lower than those without CTL immune response, which indicates that CTL immune response indeed has an important impact on infected cells and free virions and can help our body to eliminate the virions.

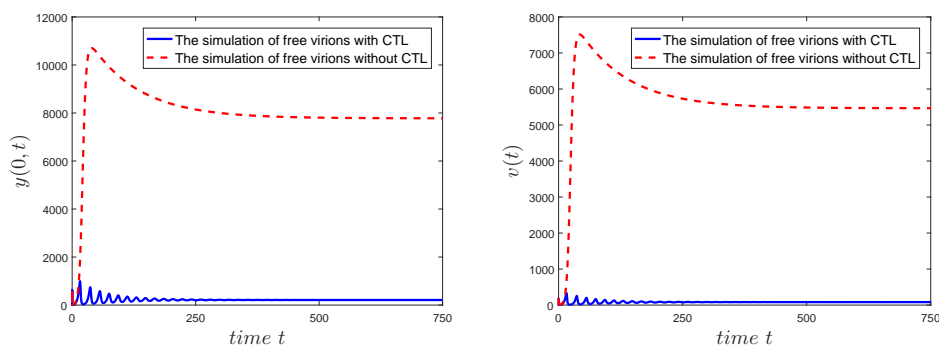


Figure 4. The temporal solutions of infected cells and free virions with and without CTL immune response found by numerical integration of system (1.3) with the boundary condition (1.4) and the similar initial condition to Figure 3.

From Figure 5, we further observe that when the proliferate rate of virus-specific CTL cells c_m increase from 0.002 to 0.004 (day^{-1}), both infected cells and free virions decreases to lower levels. This implies that CTL response can effectively reduce the quantity of infected cells and the serum viral load.

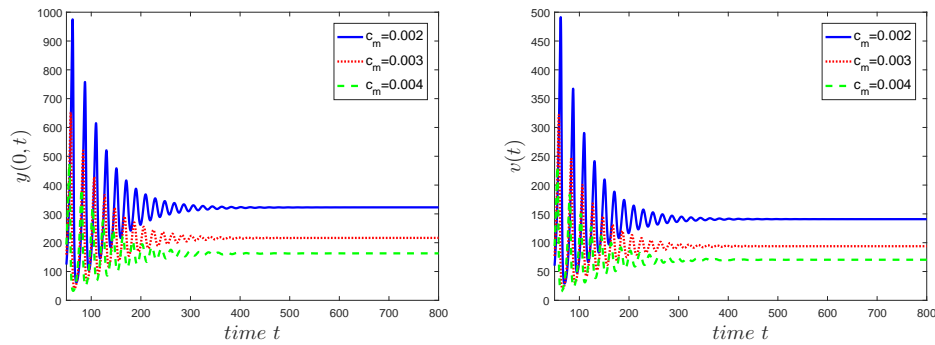


Figure 5. The temporal solutions of infected cells and free virions with different values of c_m found by numerical integration of system (1.3) with the boundary condition (1.4) and the similar initial condition to Figure 3.

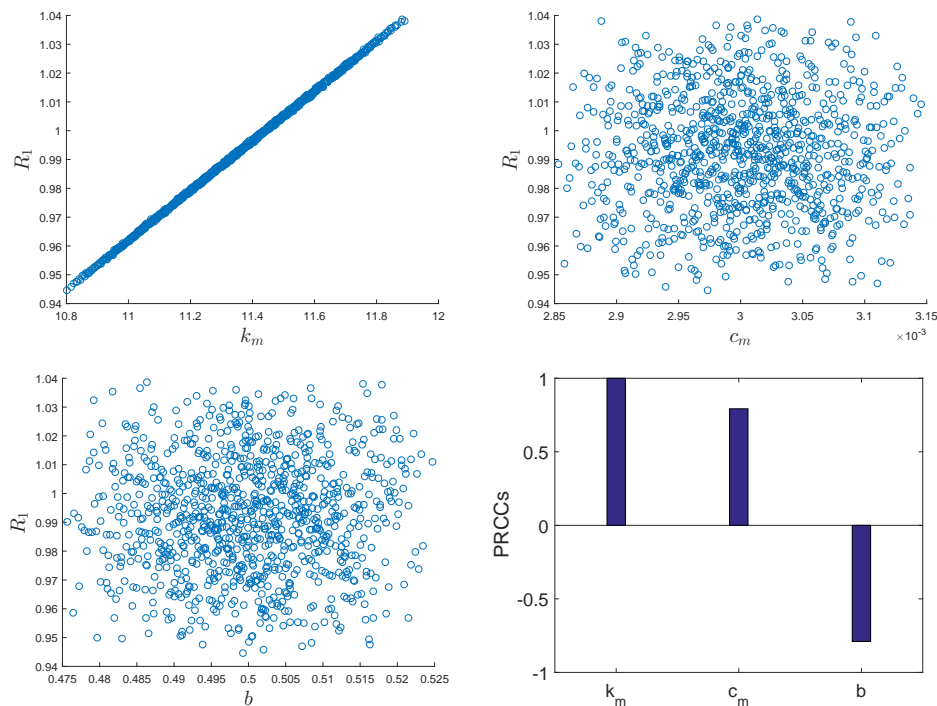


Figure 6. Scatter plots of \mathcal{R}_1 with respect to β , η , ϕ , σ , ξ and γ (first three figures). Tornado plot of partial rank correlation coefficients in respect to \mathcal{R}_1 (last figure).

Remark 7.2. *In our model, the death rate and the viral production rate of infected cells, the killing rate of infected cells by CTL and the proliferate rate of virus-specific CTL cells are assumed to vary according to the time a cell has been infected. Compared with the standard CTL response models without age structure, age-structure has more realistic representations of the biology of HIV-1 infection.*

We now carry out the sensitivity analysis of \mathcal{R}_1 . Through analysis of the sample derived from Latin hypercube sampling, we can obtain large efficient data in respect to different parameters of \mathcal{R}_1 . The first three figures in Figure 6 shows the scatter plots of \mathcal{R}_1 in respect to k_m , c_m and b , respectively, which implies that k_m and c_m are both positive correlative variables with \mathcal{R}_1 ; b is negative correlative variable with \mathcal{R}_1 . It is worth mentioning that k_m contributes more to \mathcal{R}_1 compared to c_m , namely, k_m is a more important factor in \mathcal{R}_1 . The last figure in Figure 6 shows a tornado plot of partial rank correlation coefficients with respect to \mathcal{R}_1 , indicating the importance of each parameter's uncertainty in contributing to \mathcal{R}_1 in the time to eradicate infection, which has the similar results to the first three figures in Figure 6.

In the following, we carry out corresponding numerical simulations about the relation between the immune-activated reproduction rate \mathcal{R}_1 and the proliferate rate of virus-specific CTL cells c_m . As shown in Figure 7, we find that as the proliferate rate c_m decreases, the value of \mathcal{R}_1 changes from greater than one to less than one.

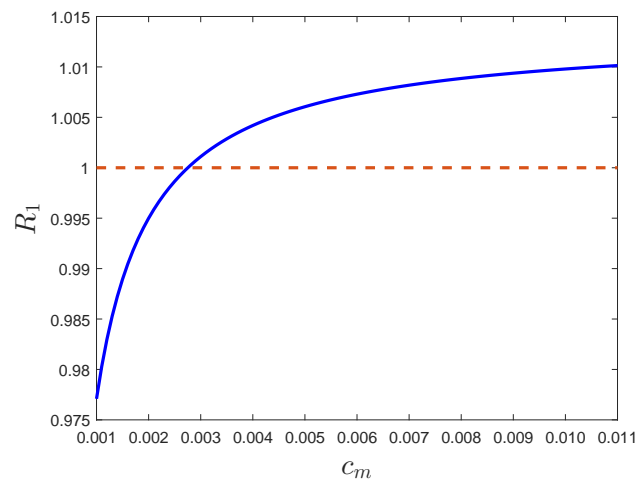


Figure 7. The curve of \mathcal{R}_1 with respect to the proliferate rate of virus-specific CTL cells c_m .

8. Conclusion

In this work, we have investigated an age-structured HIV-1 infection model with CTL immune response. The model allows the production rate of viral particles, the death rate of productively infected cells, the removed rate of infected cells and the proliferate rate of virus-specific CTLs to vary and depend on the infection age. By constructing suitable Lyapunov functionals and using LaSalle's invariance principle, we have investigated the global dynamics of each of feasible steady state of system (1.3). By Theorem 6.1, we see that if the immune-inactivated reproduction rate \mathcal{R}_0 is less than

unity, the infection-free steady state is globally asymptotically stable. In this case, the virus is finally cleared up. By Theorem 6.2, we know that if the immune-activated reproduction rate \mathcal{R}_1 satisfies $\mathcal{R}_1 < 1 < \mathcal{R}_0$, sufficient conditions are derived for the global stability of the CTL-inactivated infection steady state. In this case, the infection becomes chronic but without CTL immune response. If $\mathcal{R}_1 > 1$, by Theorem 6.3, sufficient conditions are obtained for the global attractivity of the CTL-activated infection steady state. In this case, the infection turns to chronic with CTL immune response. We would like to point out here that Theorems 6.2 and 6.3 have room for improvement, we leave this for future work.

Acknowledgments

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Conflict of interest

The authors declare that they have no competing interests.

References

1. H. D. Kwon, Optimal treatment strategies derived from a HIV model with drug-resistant mutants, *Appl. Math. Comput.*, **188** (2007), 1193–1204.
2. S. Bonhoeffer, R. M. May, G. M. Shaw, et al., Virus dynamics and drug therapy, *Proc. Natl. Acad. Sci. USA*, **94** (1997), 6971–6976.
3. D. D. Ho, A. U. Neumann, A. S. Perelson, et al., Rapid turnover of plasma virions and CD4⁺ lymphocytes in HIV-1 infection, *Nature*, **373** (1995), 123–126.
4. M. A. Nowak, R. M. Anderson, M. C. Boerlijst, et al., HIV-1 evolution and disease progression, *Science*, **274** (1996), 1008–1011.
5. M. A. Nowak, S. Bonhoeffer, G. M. Shaw, et al., Anti-viral drug treatment: Dynamics of resistance in free virus and infected cell populations, *J. Theor. Biol.*, **184** (1997), 203–217.
6. A. S. Perelson and P. W. Nelson, Mathematical analysis of HIV-1 dynamics in vivo, *SIAM Rev.*, **41** (1999), 3–44.
7. A. S. Perelson, D. E. Kirschner and R. De Boer, Dynamics of HIV infection of CD4⁺T cells, *Math. Biosci.*, **114** (1993), 81–125.
8. A. S. Perelson, A. U. Neumann, M. Markowitz, et al., HIV-1 dynamics in vivo: Virion clearance rate, infected cell life-span, and viral generation time, *Science*, **271** (1996), 1582–1586.
9. P. W. Nelson, M. A. Gilchrist, D. Coombs, et al., An age-structured model of HIV infection that allows for variations in the production rate of viral particles and the death rate of productively infected cells, *Math. Biosci. Eng.*, **1** (2004), 267–288.
10. G. Huang, X. Liu and Y. Takeuchi, Lyapunov functions and global stability for age-structured HIV infection model, *SIAM J. Appl. Math.*, **72** (2012), 25–38.

11. L. Rong, Z. Feng and A. S. Perelson, Mathematical analysis of age-structured HIV-1 dynamics with combination antiretroviral therapy, *SIAM J. Appl. Math.*, **67** (2007), 731–756.
12. R. Xu, X. Tian and S. Zhang, An age-structured within-host HIV-1 infection model with virus-to-cell and cell-to-cell transmissions, *J. Biol. Dyn.*, **12** (2017), 89–117.
13. G. W. Suryawanshi and A. Hoffmann, A multi-scale mathematical modeling framework to investigate anti-viral therapeutic opportunities in targeting HIV-1 accessory proteins, *J. Theor. Biol.*, **386** (2015), 89–104.
14. J. Xu, Y. Geng and Y. Zhou, Global dynamics for an age-structured HIV virus infection model with cellular infection and antiretroviral therapy, *Appl. Math. Comput.*, **305** (2017), 62–83.
15. J. Wang, J. Lang and X. Zou, Analysis of an age structured HIV infection model with virus-to-cell infection and cell-to-cell transmission, *Nonlinear Anal. RWA*, **34** (2017), 75–96.
16. J. Wang, R. Zhang and T. Kuniya, Global dynamics for a class of age-infection HIV models with nonlinear infection rate, *J. Math. Anal. Appl.*, **432** (2015), 289–313.
17. A. Alshorman, C. Samarasinghe, W. Lu, et al., An HIV model with age-structured latently infected cells, *J. Biol. Dyn.*, **11** (2017), 192–215.
18. Z. Liu and Z. Li, Molecular imaging in tracking tumor-specific cytotoxic T lymphocytes (CTLs), *Theranostics*, **4** (2014), 990–1001.
19. P. K. Roy and A. N. Chatterjee, T-cell proliferation in a mathematical model of CTL activity through HIV-1 infection, *Proc. World Congr. Eng.*, **1** (2010), 1–6.
20. J. Cao, J. McNevin, S. Holte, et al., Comprehensive analysis of human immunodeficiency virus type 1 (HIV-1)-specific gamma interferon-secreting CD8⁺T cells in primary HIV-1 infection, *J. Virol.*, **77** (2003), 6867–6878.
21. C. Browne, Immune response in virus model structured by cell infection-age, *Math. Biosci. Eng.*, **13** (2016), 887–909.
22. J. Pang, J. Chen, Z. Liu, et al., Local and global stabilities of a viral dynamics model with infection-age and immune response, *J. Dyn. Differ. Equ.*, **31** (2019), 793–813.
23. R. R. Regoes, D. Ebert and S. Bonhoeffer, Dose-dependent infection rates of parasites produce the Allee effect in epidemiology, *Proc. R. Soc. London B*, **269** (2002), 271–279.
24. M. Iannelli, Mathematical Theory of Age-Structured Population Dynamics, *Applied Mathematics Monographs*, **7**, Consiglio Nazionale delle Ricerche (C.N.R), Giardini Pisa, 1995, comitato nazionale per le scienze matematiche.
25. G. Webb, Theory of Nonlinear Age-Dependent Population Dynamics, *Marcel Dekker, New York*, 1985.
26. H. L. Smith and H. R. Thieme, Dynamical Systems and Population Persistence, *American Mathematical Society, Providence*, 2011.
27. J. Yang, R. Xu and X. Luo, Dynamical analysis of an age-structured multi-group SIVS epidemic model, *Math. Biosci. Eng.*, **16** (2019), 636–666.
28. D. Burg, L. Rong, A. U. Neumann, et al., Mathematical modeling of viral kinetics under immune control during primary HIV-1-infection, *J. Theor. Biol.*, **259** (2009), 751–759.

29. N. M. Dixit and A. S. Perelson, Complex patterns of viral load decay under antiretroviral therapy: Influence of pharmacokinetics and intracellular delay, *J. Theor. Biol.*, **226** (2004), 95–109.
30. M. Markowitz, M. Louie, A. Hurley, et al., A novel antiviral intervention results in more accurate assessment of human immunodeficiency virus type 1 replication dynamics and t-cell decay in vivo, *J. Virol.*, **77** (2003), 5037–5038.
31. B. Ramratnam, S. Bonhoeffer, J. Binley, et al., Rapid production and clearance of HIV-1 and hepatitis C virus assessed by large volume plasma apheresis, *Lancet*, **354** (1999), 1782–1785.
32. J. Wang, M. Guo, X Liu, et al., Threshold dynamics of HIV-1 virus model with cell-to-cell transmission, cell-mediated immune responses and distributed delay, *Appl. Math. Comput.*, **291** (2016), 149–161.
33. X. Zhou and J. Cui, Global stability of the viral dynamics with Crowley-Martin functional response, *Bull. Korean Math. Soc.*, **48** (2011), 555–574.
34. M. Iannelli and F. Milner, *The Basic Approach to Age-Structured Population Dynamics: Models, Methods and Numerics*, Springer, 2017.
35. J. K. Hale and P. Waltman, Persistence in infinite-dimensional systems, *SIAM J. Math. Anal.*, **20** (1989), 388–395.
36. P. Magal, Compact attractors for time periodic age-structured population models, *Electron. J. Differ. Equ.*, **65** (2001), 1–35.
37. P. Magal, C. C. McCluskey and G. F. Webb, Lyapunov functional and global asymptotic stability for an infection-age model, *Appl. Anal.*, **89** (2010), 1109–1140.
38. R. Xu, X. Tian and F. Zhang, Global dynamics of a tuberculosis transmission model with age of infection and incomplete treatment, *Adv. Differ. Equ.*, **242** (2017), 1–34.



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