



Research article

Qualitative behaviour of a stochastic hepatitis C epidemic model in cellular level

Dwi Lestari^{1,2}, Noorma Yulia Megawati^{1,*}, Nanang Susyanto¹ and Fajar Adi-Kusumo¹

¹ Department of Mathematics, Universitas Gadjah Mada, Yogyakarta, Indonesia

² Department of Mathematics Education, Universitas Negeri Yogyakarta, Yogyakarta, Indonesia

* **Correspondence:** Email: noorma_yulia@ugm.ac.id.

Abstract: In this paper, a mathematical model describing the dynamical of the spread of hepatitis C virus (HCV) at a cellular level with a stochastic noise in the transmission rate is developed from the deterministic model. The unique time-global solution for any positive initial value is served. The Ito's Formula, the suitable Lyapunov function, and other stochastic analysis techniques are used to analyze the model dynamics. The numerical simulations are carried out to describe the analytical results. These results highlight the impact of the noise intensity accelerating the extinction of the disease.

Keywords: hepatitis C virus; stochastic noise; transmission rate; extinction; persistence in mean; numerical simulation

1. Introduction

Hepatitis C is a liver inflammatory disease caused by a viral infection. Hepatitis C virus (HCV) is an RNA-type virus from the Flaviviridae family (genus Hepacivirus) which has a high replication rate. Hepatitis can develop into liver cancer [1] and can cause serious complications even death [2, 3]. Around 60 – 85% of acute hepatitis C may develop into a chronic condition, 10–15% may develop cirrhosis, and 25% may develop liver cancer [1, 2, 4]. It is estimated that 58 million people have a chronic hepatitis C, with 1.5 million new infections occurring per year over the world [5]. There is no vaccine available for HCV [3, 4, 6, 7], thus it is important to gain a better insight into the nature of this disease.

One of the accurate tools to understand the dynamics of hepatitis C is using a mathematical model. There are several mathematical models developed for HCV for example in [8, 9] studying the HCV model in population level. The HCV model incorporating treatment, therapy, or isolation were investigated in [10–14]. In reference [15], the optimal control strategies for HCV epidemics considering the uncertainty of the model was discussed.

The spread of HCV at the cellular level was also investigated by several researchers e.g., [16–20]. In 1998, Neumann et al. studied the dynamic of HCV and the effect of antiviral. They have found that HCV is highly dynamic. An existing deterministic model cannot be applied to describe randomness in many biological factors, for instance, the random occurrence of cell infection, mutation, and apoptosis [21,22]. Furthermore, during the spread of HCV, different cells and infectious virus particles reacting in the same environment can give different effects. In other words, some uncertain factors influence the spread of HCV in body cells such as lifestyle (alcohol consumption or smoking), patient compliance level, lipid metabolism, metabolic syndrome, and body weight [23]. Motivated by this phenomenon, we are interested in extending the mathematical model given in [24] by considering uncertainty factors. A stochastic model will be derived from the deterministic model. This model provides better insight into the uncertainty and variability of the disease dynamic. Moreover, the solution of the stochastic model is in the form of distribution [25], while the solution of the deterministic model only produces one predictive value [26]. Several papers also discussed the stochastic HCV model, e.g., [27–32]. In contrast, in this paper, the noise parameter is added in the transmission rate representing the characteristic of variability and the treatment in the model is considered.

The outline of this paper is organized as follows. In Section 2, the HCV model involving stochastic disturbances in the transmission rate are developed. In addition, the existence, uniqueness, and boundedness of the solution are established. In Section 3, we derive the extinction and persistence in mean condition. In Section 4, the numerical simulations are carried out to illustrate the analytical results. The conclusions are presented in the last section.

2. Materials and methods

2.1. Mathematical model

Recall a deterministic mathematical model of HCV given in [24] as

$$\begin{aligned}\frac{dT}{dt} &= \Lambda - \delta_1 T - (1 - \eta)\beta VT \\ \frac{dI}{dt} &= (1 - \eta)\beta VT - \delta_2 I \\ \frac{dV}{dt} &= (1 - \epsilon)kI - cV.\end{aligned}\tag{2.1}$$

where T is the number of uninfected cells, I is the number of infected cells, and V is the number of free viruses. The uninfected cells are produced at rate Λ and die naturally at a constant rate δ_1 . Cells become infected when they are interacting with a virus with a constant rate β , and once infected, they will die at a constant rate δ_2 . HCV is produced by infected cells at a constant rate k and cleared at a constant rate c .

For the deterministic model (2.1), the disease-free equilibrium point is $E_0 = \left(\frac{\Lambda}{\delta_1}, 0, 0\right)$ and the endemic equilibrium point is $E_1 = \left(\frac{\delta_2 c}{(1-\eta)(1-\epsilon)k\beta}, \frac{\Lambda}{\delta_2} - \frac{\delta_1 c}{(1-\eta)(1-\epsilon)k\beta}, \frac{(1-\epsilon)k\Lambda}{\delta_2 c} - \frac{\delta_1}{(1-\eta)\beta}\right)$. Using the next-generation matrix method [33], the basic reproduction number of the system (2.1) is $R_0^d = \frac{(1-\epsilon)(1-\eta)\beta k \Lambda}{c \delta_1 \delta_2}$.

Table 1. Variables and parameters of the model.

Variable/Parameter	Description	Unit	Range
$T(t)$	concentration of uninfected liver cells	cell/mL	≥ 0
$I(t)$	concentration of infected liver cells	cell/mL	≥ 0
$V(t)$	concentration of free viruses	virion/mL	≥ 0
Λ	rate of production	cell/day×mL	≥ 0
η	the effectiveness of drug in stopping infection	–	[0,1]
k	rate of free virus production	virion/cell×day	≥ 0
ϵ	the effectiveness of drug in blocking virus	–	[0,1]
β	transmission rate	mL/virion × day	≥ 0
δ_1	death rate of uninfected liver cells	1/day	≥ 0
δ_2	death rate of infected liver cells	1/day	≥ 0
c	virion clearance	1/day	≥ 0

In this paper, we generalize system (2.1) by incorporating a stochastic noise parameter in the transmission rate as

$$\begin{cases} dT = (\Lambda - \delta_1 T - (1 - \eta)\beta VT) dt - \sigma(1 - \eta)VT dB(t) \\ dI = ((1 - \eta)\beta VT - \delta_2 I) dt + \sigma(1 - \eta)VT dB(t) \\ dV = ((1 - \epsilon)kI - cV) dt \end{cases} \quad (2.2)$$

where $B(t)$ is a standard Brownian motion and σ is a real constant which is known as the intensity of noise. The value σ is the standard deviation of transmission rate data that represents the variability of the transmission rate. The other description of parameters model (2.2) is given in Table 1. Clearly, the system (2.1) is the special case of the system (2.2), where $\sigma = 0$.

2.2. Existence and uniqueness of the solution

In this section, the theorem of the existence, uniqueness, and boundedness of the solution system (2.2) are established. Let (Ω, \mathcal{F}, P) be a complete probability space with a filtration $\{\mathcal{F}_t\}_{t \geq 0}$ satisfying the conditions that it is right continuous and \mathcal{F}_0 contains all P-null sets.

Definition 2.1. [34]

Let $T > 0$, $F(.,.) : [0, T] \times \mathbb{R}^n \rightarrow \mathbb{R}^n$, $G(.,.) : [0, T] \times \mathbb{R}^n \rightarrow \mathbb{R}^{n \times m}$ be measurable function, and $X(t)$ satisfy

$$dX(t) = F(X(t))dt + G(X(t))dB(t), \quad (2.3)$$

where $F(.,.)$ and $G(.,.)$ are the coefficients of (2.3). Then the coefficients (2.3) are locally Lipschitz, if

$$|F(t, X) - F(t, \bar{X})| + |G(t, X) - G(t, \bar{X})| \leq C_1 |X - \bar{X}|; X, \bar{X} \in \mathbb{R}^n, t \in [0, T], \quad (2.4)$$

for some constant $C_1 \geq 0$.

Definition 2.2. [34]

The coefficients of (2.3) satisfy linear growth condition, if

$$|F(t, X)| + |G(t, X)| \leq C_2(1 + |X|); X \in \mathbb{R}^n, t \in [0, T], \quad (2.5)$$

for some constant C_2 .

Lemma 2.3. *The system (2.2) with the initial condition in \mathbb{R}_+^3 is uniformly ultimately bounded and belongs to the following closed and bounded positively invariant set*

$$\Gamma = \left\{ (T, I, V) \in \mathbb{R}_+^3 \mid 0 < T + I \leq \frac{\Lambda}{\mu}, 0 < V \leq (1 - \epsilon) \left(\frac{k\Lambda}{c\mu} \right) \right\} \subset \mathbb{R}_+^3, \quad (2.6)$$

for every $t \geq 0$.

Proof. Let $a(t)$ be a function such that $a(t) \leq \Lambda$. We know that $N(t) = T(t) + I(t)$ is the concentration of uninfected and infected cells at time t . Take $\mu = \min\{\delta_1, \delta_2\}$, then

$$\begin{aligned} dN &= dT + dI, \\ &= (\Lambda - \delta_1 T - \delta_2 I)dt, \\ dN &\leq (\Lambda - \mu N)dt. \end{aligned} \quad (2.7)$$

Let $dN = (a(t) - \mu N)dt$, it follows that

$$\begin{aligned} dN &= (a(t) - \mu N)dt, \\ \frac{dN}{dt} + \mu N &= a(t), \end{aligned}$$

thus,

$$N(t) = e^{-\mu t} \left(\int_0^t e^{\mu \tau} a(\tau) d\tau + N_0 \right). \quad (2.8)$$

Since $a(t) \leq \Lambda$, we obtain

$$\begin{aligned} N &\leq e^{-\mu t} \left(\int_0^t e^{\mu \tau} \Lambda d\tau + N_0 \right), \\ &= \frac{\Lambda}{\mu} + \left(N_0 - \frac{\Lambda}{\mu} \right) e^{-\mu t}. \end{aligned} \quad (2.9)$$

By taking a limit $t \rightarrow \infty$ of Eq (2.9), we get

$$N(t) \leq \frac{\Lambda}{\mu}.$$

Analog with the technique for solving Eq (2.7), for $V(0) = V_0$ and $I(t) < N(t) \leq \frac{\Lambda}{\mu}$, we get

$$\begin{aligned} dV(t) &= [(1 - \epsilon)kI - cV]dt, \\ &\leq [(1 - \epsilon) \left(\frac{k\Lambda}{\mu} \right) - cV]dt, \end{aligned}$$

then

$$V(t) \leq (1 - \epsilon) \left(\frac{k\Lambda}{c\mu} \right) + \left(V_0 - (1 - \epsilon) \left(\frac{k\Lambda}{\mu} \right) \right) e^{-ct}. \quad (2.10)$$

If we take $\lim_{t \rightarrow \infty} V(t)$, then

$$V(t) \leq (1 - \epsilon) \left(\frac{k\Lambda}{c\mu} \right).$$

Since the right-hand side of Eq (2.2) satisfies the Lipschitz condition, the solution exists and unique on $[0, b)$ for some $b > 0$. Assume that there exists $t_1 \in (0, b)$ such that $V(t_1) = 0$ and all other variables are positive on $t_1 \in (0, b)$. Therefore, for all $t \in [0, t_1]$

$$\begin{aligned} dV &= ((1 - \epsilon)kI - cV)dt, \\ &\geq -cVdt, \end{aligned}$$

thus

$$V(t_1) \geq C_3 e^{-ct_1} > 0, \text{ where } C_3 \text{ positive constant.} \tag{2.11}$$

This is contradiction with $V(t_1) = 0$. Then $V(t) > 0$. Analoguely, we get $I(t) > 0$ and $T(t) > 0$. Therefore, we obtain that $(T(t), I(t)) \in (0, \frac{\Lambda}{\mu})$ for all $t \in [0, T]$ and $0 < V(t) \leq (1 - \epsilon) \left(\frac{k\Lambda}{c\mu} \right)$. \square

Next, we define the necessary condition that guarantees the existence and uniqueness of time-global solution of Eq (2.2).

Theorem 2.4. *Let the coefficients of the system (2.2) satisfy the Lipschitz condition. Then for any initial value $(T(0), I(0), V(0)) \in \Gamma$, there exists a unique time-global solution $(T(t), I(t), V(t)) \in \Gamma \subset \mathbb{R}_+^3, t \geq 0$ with probability 1.*

Proof. According to Definition 2.1, Definition 2.2, and Theorem 5.2.1 in [34], there exists a unique global solution. However, we have that the coefficients of the system (2.2) only satisfy the Lipschitz condition [35], thus system (2.2) has a unique local solution on $t \in [0, \tau_e)$ for any initial value $(T(0), I(0), V(0)) \in \Gamma$ where τ_e is the explosion time (i.e. the time when the solution tends to infinity). To guarantee the solution of system (2.2) is a unique global solution, it is necessary to show that $\tau_e = \infty$ [36, 37].

Let $k_0 > 0$ be sufficiently large such that every component of $(T(0), I(0), V(0))$ is in the interval $[\frac{1}{k_0}, k_0]$. For every $k \geq k_0$, we define the stopping time or the first passage time (i.e. first period when the stochastic process penetrates the barrier) as

$$\tau_k = \inf \left\{ t \in [0, \tau_e) \mid T(t) \notin \left(\frac{1}{k}, k \right), I(t) \notin \left(\frac{1}{k}, k \right) \text{ or } V(t) \notin \left(\frac{1}{k}, k \right) \right\}.$$

Throughout this paper, we set $\inf \emptyset = \infty$. It is known that the lower bound of \mathbb{R} is an empty set and the largest infimum of the empty set is infinity. Since τ_k is increasing as $k \rightarrow \infty$, and

$$\tau_0 \leq \tau_1 \leq \tau_2 \leq \dots \leq \tau_k \leq \tau_{k+1} \leq \dots,$$

then it follows that

$$\tau_\infty = \lim_{k \rightarrow \infty} \tau_k,$$

thus $\tau_\infty \leq \tau_e$ almost sure. Next, it is necessary to show

$$\lim_{k \rightarrow \infty} \tau_k = \infty.$$

We will prove by contradiction. Suppose that $P(\tau_\infty < \infty) < 1$. If $\tau_\infty < \infty$, then there exist $T^* > 0$ and $\varepsilon \in (0, 1)$ such that $P\{\tau_k \leq T^*\} > \varepsilon, \forall k \geq k_0$. In this case, the proof technique analogue is to [37–39]. Define a function $C^{2,1}, Q : [0, \infty) \times \mathbb{R}_+^3 \rightarrow \mathbb{R}_+$ where

$$Q(T, I, V) = (T - 1 - \ln T) + (I - 1 - \ln I) + (V - 1 - \ln V).$$

Since

$$y - 1 - \ln y \geq 0, \forall y > 0,$$

then Q is a non-negative function. By using Itô's formula, we get

$$\begin{aligned} dQ &= \frac{\partial Q}{\partial t} dt + \frac{\partial Q}{\partial T} dT + \frac{\partial Q}{\partial I} dI + \frac{\partial Q}{\partial V} dV + \left(\frac{1}{2}\right) \left(\frac{\partial^2 Q}{\partial T^2} (dT)^2\right) + \left(\frac{1}{2}\right) \left(\frac{\partial^2 Q}{\partial I^2} (dI)^2\right) \\ &\quad + \left(\frac{1}{2}\right) \left(\frac{\partial^2 Q}{\partial V^2} (dV)^2\right) + \frac{\partial^2 Q}{\partial T \partial I} dT dI + \frac{\partial^2 Q}{\partial T \partial V} dT dV + \frac{\partial^2 Q}{\partial I \partial V} dI dV, \\ &= \left(1 - \frac{1}{T}\right) \{[\Lambda - \delta_1 T - (1 - \eta)\beta VT] dt - \sigma(1 - \eta)VT dB(t)\} \\ &\quad + \left(1 - \frac{1}{I}\right) \{[(1 - \eta)\beta VT - \delta_2 I] dt + \sigma(1 - \eta)VT dB(t)\} \\ &\quad + \left(1 - \frac{1}{V}\right) \{[(1 - \epsilon)kI - cV] dt\} + \frac{1}{2}(\sigma VT)^2(1 - \eta)^2 \left(\frac{1}{T^2} + \frac{1}{I^2}\right) (dB(t))^2, \\ &\leq \left\{ \Lambda + \delta_1 + \delta_2 + (1 - \epsilon)kI + (1 - \eta)\beta V + c + \frac{1}{2}(\sigma VT)^2(1 - \eta)^2 \right. \\ &\quad \left. \left(\frac{1}{T^2} + \frac{1}{I^2}\right) \right\} dt + \left\{ \sigma(1 - \eta)VT \left(\frac{1}{T} - \frac{1}{I}\right) \right\} dB(t), \\ &\leq Mdt + \sigma(1 - \eta)VT \left(\frac{1}{T} - \frac{1}{I}\right) dB(t), \end{aligned} \tag{2.12}$$

where M is a positive constant. Integrating Eq (2.12) from 0 to $\tau_k \wedge T^*$ yields

$$\begin{aligned} Q(T(\tau_k \wedge T^*), I(\tau_k \wedge T^*), V(\tau_k \wedge T^*)) - Q(T(0), I(0), V(0)) \\ \leq Mt + \int_0^{\tau_k \wedge T^*} \sigma(1 - \eta)VT \left(\frac{1}{T} - \frac{1}{I}\right) dB(t). \end{aligned} \tag{2.13}$$

Taking the expectation of both sides in Eq (2.13) leads to

$$\begin{aligned} E(Q(T(\tau_k \wedge T^*), I(\tau_k \wedge T^*), V(\tau_k \wedge T^*))) &\leq E(Q(T(0), I(0), V(0))) + E(Mt) \\ &\quad + E \left[\int_0^{\tau_k \wedge T^*} \sigma(1 - \eta)VT \left(\frac{1}{T} - \frac{1}{I}\right) dB(t) \right], \end{aligned}$$

$$\leq Q(T(0), I(0), V(0)) + Mt. \quad (2.14)$$

Define $\Omega_k = \{\tau_k \leq T^*\}$ for any $k \geq k_0$. Then for every $w \in \Omega_k$ there are components of $Q(T(\tau_k \wedge T^*), I(\tau_k \wedge T^*), V(\tau_k \wedge T^*))$ which are equal to either k or $\frac{1}{k}$, thus

$$Q(T(\tau_k \wedge T^*), I(\tau_k \wedge T^*), V(\tau_k \wedge T^*)) \geq \min \left\{ k - 1 - \ln k, \frac{1}{k} - 1 - \ln \left(\frac{1}{k} \right) \right\}. \quad (2.15)$$

Combining Eqs (2.14) and (2.15) yields

$$\begin{aligned} Q(T(0), I(0), V(0)) + Mt &\geq E(Q(T(\tau_k \wedge T^*), I(\tau_k \wedge T^*), V(\tau_k \wedge T^*), w)), \\ &= E(1_{\Omega_k(w)} \cdot Q(T(\tau_k), I(\tau_k), V(\tau_k)), w), \\ &\geq P(\Omega_k) \min \left\{ k - 1 - \ln k, \frac{1}{k} - 1 - \ln \left(\frac{1}{k} \right) \right\}, \\ &\geq \varepsilon \min \left\{ k - 1 - \ln k, \frac{1}{k} - 1 - \ln \left(\frac{1}{k} \right) \right\}, \end{aligned} \quad (2.16)$$

where 1_{Ω_k} is the indicator function. By taking a limit $k \rightarrow \infty$ of Eq (2.16), we obtain

$$\infty = Q(T(0), I(0), V(0)) + Mt < \infty.$$

This is contradiction. We get $\tau_\infty = \infty$ or $P(\tau_\infty = \infty) = 1$. Therefore, there exists a unique time-global solution $(T(t), I(t), V(t)) \in \Gamma \subset \mathbb{R}_+^3, t \geq 0$ with probability 1. \square

3. The equilibria and their stabilities

In this section, the theorem of the extinction and persistence in the mean condition system (2.2) is derived.

3.1. Extinction

In this part, we investigate the almost surely exponential stability of the disease-free equilibrium point by using the suitable Lyapunov function and another technique of stochastic analysis.

Lemma 3.1. *If*

$$\sigma > \frac{\beta}{\sqrt{2(\delta_2 - (1 - \epsilon)k \wedge c)}}, \quad (3.1)$$

the disease-free equilibrium point $(T, I, V) = (\Lambda/\delta_1, 0, 0)$ is almost surely exponentially stable in Γ .

Proof. The technical proof of this theorem follows from [38, 40]. Define a function $Q = \ln(I + V)$. Using Itô's formula, we obtain

$$\begin{aligned} dQ &= \frac{\partial Q}{\partial t} dt + \frac{\partial Q}{\partial T} dT + \frac{\partial Q}{\partial I} dI + \frac{\partial Q}{\partial V} dV + \frac{1}{2} \left[\frac{\partial^2 Q}{\partial T^2} dT dT + \frac{\partial^2 Q}{\partial I^2} dI dI + \frac{\partial^2 Q}{\partial V^2} dV dV \right] \\ &\quad + \frac{\partial^2 Q}{\partial T \partial I} dT dI + \frac{\partial^2 Q}{\partial T \partial V} dT dV + \frac{\partial^2 Q}{\partial I \partial V} dI dV, \end{aligned}$$

$$\begin{aligned}
 &= \frac{1}{I+V} [((1-\eta)\beta VT - \delta_2 I)dt + \sigma(1-\eta)VTdB] + \frac{1}{I+V} [(1-\epsilon)kI - cV]dt \\
 &\quad + \frac{1}{2} \left[-\frac{1}{(I+V)^2} (\sigma^2(1-\eta)^2(VT)^2)dt \right], \\
 &= \left\{ (1-\eta)\beta Z - \frac{\delta_2 I}{I+V} - \frac{cV}{I+V} + \frac{(1-\epsilon)kI}{I+V} - \frac{1}{2}\sigma^2(1-\eta)^2 Z^2 \right\} dt + \sigma(1-\eta)ZdB,
 \end{aligned}$$

where $Z = \frac{VT}{I+V}$. Futhermore, we have

$$\begin{aligned}
 dQ &\leq -\frac{1}{2}\sigma^2(1-\eta)^2 \left[Z - \frac{\beta}{\delta^2(1-\eta)} \right]^2 dt + \frac{\beta^2}{2\sigma^2} dt - [\delta_2 - (1-\epsilon)k \wedge c]dt + \sigma(1-\eta)ZdB, \\
 &\leq \frac{\beta^2}{2\sigma^2} dt - [\delta_2 - (1-\epsilon)k \wedge c]dt + \sigma(1-\eta)ZdB.
 \end{aligned} \tag{3.2}$$

Taking the integral of (3.2) and dividing both sides by t and then computing the limit superior $t \rightarrow \infty$ yield

$$\begin{aligned}
 \limsup_{t \rightarrow \infty} \frac{1}{t} \ln(I(t) + V(t)) &\leq \limsup_{t \rightarrow \infty} \frac{1}{t} \{ \ln(I(0) + V(0)) + \frac{\beta^2}{2\sigma^2} - [\delta_2 - (1-\epsilon)k \wedge c] \} \\
 &\quad + \limsup_{t \rightarrow \infty} \int_0^t \frac{1}{t} \sigma(1-\eta)ZdB.
 \end{aligned} \tag{3.3}$$

By the strong law of large number for Martingales [41], we have

$$\limsup_{t \rightarrow \infty} \int_0^t \frac{1}{t} \sigma(1-\eta)ZdB = 0. \tag{3.4}$$

Thus, from Eqs (3.3) and (3.4), we obtain

$$\limsup_{t \rightarrow \infty} \frac{1}{t} \ln(I(t) + V(t)) \leq \limsup_{t \rightarrow \infty} \frac{1}{t} \{ \ln(I(0) + V(0)) + \frac{\beta^2}{2\sigma^2} - [\delta_2 - (1-\epsilon)k \wedge c] \} < 0.$$

□

By Lemma 3.1, if the noise is increasing and satisfies the condition (3.1) HCV will die out. However, in this study, we also get that a bounded variation of infection rate could also lead to extinction. This is presented in the following theorem.

Theorem 3.2. *Let $(T(0), I(0), V(0)) \in \Gamma$ be the initial value of system (2.2) and $(T(t), I(t), V(t)) \in \Gamma$ be the corresponding solution. If*

$$\begin{aligned}
 R_0^s &= \frac{(1-\epsilon)(1-\eta)\beta k \Lambda}{c\delta_1\delta_2} - \left[\frac{1}{4c\delta_1^2\delta_2^2} (1-\epsilon)^2(1-\eta)^2 k^2 \Lambda^2 \sigma^2 \right] \\
 &= R_0^d - \left[\frac{1}{4c\delta_1^2\delta_2^2} (1-\epsilon)^2(1-\eta)^2 k^2 \Lambda^2 \sigma^2 \right] < 1
 \end{aligned}$$

and $\sigma^2 < \frac{4\delta_1\delta_2\beta}{(1-\epsilon)(1-\eta)k\Lambda}$, then HCV will be extinct almost surely, i.e.,

$$\lim_{t \rightarrow \infty} T(t) = \frac{\Lambda}{\delta_1}, \lim_{t \rightarrow \infty} I(t) = 0, \lim_{t \rightarrow \infty} V(t) = 0.$$

Proof. The proof technique follows from [38, 40, 42, 43]. Consider the first equation of system (2.2), then we get

$$T(t) - T(0) = \Lambda t - \int_0^t (1 - \eta)\beta V(s)T(s)ds - \int_0^t \delta_1 T(s)ds - \int_0^t \sigma(1 - \eta)V(s)T(s)dB(s),$$

or

$$\frac{1}{t} \int_0^t T(s)ds = \frac{\Lambda}{\delta_1} - \frac{(T(t) - T(0))}{\delta_1 t} - \frac{1}{\delta_1 t} \int_0^t (1 - \eta)\beta V(s)T(s)ds - \frac{1}{\delta_1 t} \int_0^t \sigma(1 - \eta)V(s)T(s)dB(s).$$

For simplicity, we define an integrable function $\langle X \rangle = \frac{1}{t} \int_0^t X(s)ds$, $X(t) \in [0, +\infty)$, thus

$$\langle T \rangle = \frac{\Lambda}{\delta_1} - \frac{(T(t) - T(0))}{\delta_1 t} - \frac{1}{\delta_1} (1 - \eta)\beta \langle VT \rangle - \frac{1}{\delta_1 t} \int_0^t \sigma(1 - \eta)V(s)T(s)dB(s). \quad (3.5)$$

Taking the limit of Eq (3.5) as t tend to infinity, we obtain

$$\begin{aligned} \lim_{t \rightarrow \infty} \langle T \rangle &= \frac{1}{\delta_1} \lim_{t \rightarrow \infty} \left[\Lambda - \frac{(T(t) - T(0))}{t} - (1 - \eta)\beta \langle VT \rangle - \frac{1}{t} \int_0^t \sigma(1 - \eta)V(s)T(s)dB(s) \right], \\ &\leq \frac{1}{\delta_1} \lim_{t \rightarrow \infty} \left[\Lambda - \frac{1}{t} \int_0^t \sigma(1 - \eta)V(s)T(s)dB(s) \right], \\ &\leq \frac{\Lambda}{\delta_1}. \end{aligned}$$

Next, define a function $Q = \ln((1 - \epsilon)kI + \delta_2 V)$. Applying Ito's formula, we obtain

$$\begin{aligned} dQ &= \frac{\partial Q}{\partial t} dt + \frac{\partial Q}{\partial T} dT + \frac{\partial Q}{\partial I} dI + \frac{\partial Q}{\partial V} dV + \frac{1}{2} \left[\frac{\partial^2 Q}{\partial T^2} dT dT + \frac{\partial^2 Q}{\partial I^2} dI dI + \frac{\partial^2 Q}{\partial V^2} dV dV \right] \\ &\quad + \frac{\partial^2 Q}{\partial T \partial I} dT dI + \frac{\partial^2 Q}{\partial T \partial V} dT dV + \frac{\partial^2 Q}{\partial I \partial V} dI dV, \\ &= \frac{(1 - \epsilon)k}{(1 - \epsilon)kI + \delta_2 V} [(1 - \eta)\beta VT - \delta_2 I] dt + \sigma(1 - \eta)VT dB \\ &\quad + \frac{\delta_2}{(1 - \epsilon)kI + \delta_2 V} [(1 - \epsilon)kI - cV] dt \\ &\quad + \frac{1}{2} \left[-\frac{1}{((1 - \epsilon)kI + \delta_2 V)^2} (1 - \epsilon)^2 k^2 (\sigma^2 (1 - \eta)^2 (VT)^2) dt \right], \\ &\leq \left[\frac{(1 - \epsilon)k(1 - \eta)\beta T}{\delta_2} - c - \frac{1}{2\delta_2^2} (1 - \epsilon)^2 k^2 \sigma^2 (1 - \eta)^2 T^2 \right] dt \\ &\quad + \frac{(1 - \epsilon)k}{(1 - \epsilon)kI + \delta_2 V} \sigma(1 - \eta)VT dB. \end{aligned} \quad (3.6)$$

Integrating both sides of Eq (3.6) from 0 to t yields

$$\ln((1 - \epsilon)kI + \delta_2 V) \leq \ln((1 - \epsilon)kI(0) + \delta_2 V(0)) + \frac{(1 - \epsilon)k}{\delta_2} (1 - \eta)\beta \int_0^t T(s)ds$$

$$\begin{aligned}
 & - ct - \frac{1}{4\delta_2^2}(1 - \epsilon)^2 k^2 \sigma^2 (1 - \eta)^2 \int_0^t (T(s))^2 ds \\
 & + \int_0^t \frac{(1 - \epsilon)k}{(1 - \epsilon)kI(s) + \delta_2 V(s)} \sigma(1 - \eta)VT dB(s).
 \end{aligned} \tag{3.7}$$

Dividing both sides of Eq (3.7) by t and taking a limit superior for $t \rightarrow \infty$, we get

$$\begin{aligned}
 \limsup_{t \rightarrow \infty} \frac{1}{t} \ln((1 - \epsilon)kI + \delta_2 V) & \leq \limsup_{t \rightarrow \infty} \frac{1}{t} [\ln((1 - \epsilon)kI(0) + \delta_2 V(0)) + \frac{(1 - \epsilon)k}{\delta_2} (1 - \eta)\beta \int_0^t T(s)ds] \\
 & - c - \limsup_{t \rightarrow \infty} \frac{1}{4\delta_2^2}(1 - \epsilon)^2 k^2 \sigma^2 (1 - \eta)^2 \int_0^t (T(s))^2 ds \\
 & + \int_0^t \frac{(1 - \epsilon)k}{(1 - \epsilon)kI(s) + \delta_2 V(s)} \sigma(1 - \eta)VT dB(s).
 \end{aligned}$$

By the strong law of large number for Martingales [41],

$$\limsup_{t \rightarrow \infty} \frac{1}{t} \int_0^t \frac{(1 - \epsilon)k}{(1 - \epsilon)kI(s) + \delta_2 V(s)} \sigma(1 - \eta)VT dB(s) = 0,$$

we obtain

$$\begin{aligned}
 \limsup_{t \rightarrow \infty} \frac{1}{t} \ln((1 - \epsilon)kI + \delta_2 V) & \leq \frac{(1 - \epsilon)k}{\delta_1 \delta_2} (1 - \eta)\beta \Lambda - c - \frac{1}{4\delta_2^2}(1 - \epsilon)^2 k^2 \sigma^2 (1 - \eta)^2 \frac{(\Lambda)^2}{(\delta_1)^2}, \\
 & \leq c \left\{ \left[\frac{(1 - \epsilon)(1 - \eta)\beta k \Lambda}{c \delta_1 \delta_2} - \frac{1}{4c\delta_2^2 \delta_1^2} (1 - \epsilon)^2 (1 - \eta)^2 k^2 \Lambda^2 \sigma^2 \right] - 1 \right\}, \\
 & \leq c[R_0^s - 1] < 0.
 \end{aligned}$$

Thus, $\lim_{t \rightarrow \infty} I(t) = 0$ and $\lim_{t \rightarrow \infty} V(t) = 0$. In other words, HCV will be extinct. □

3.2. Persistence in mean

In this subsection, we give a sufficient condition to guarantee the persistence in the mean condition.

Definition 3.3. [31, 40, 44]

System (2.2) is said to be persistent in mean if

$$\liminf_{t \rightarrow \infty} \frac{1}{t} \int_0^t [I(s) + V(s)] ds > 0$$

almost surely.

Theorem 3.4. Let $(T(0), I(0), V(0)) \in \Gamma$ be the initial value of system (2.2) and $(T(t), I(t), V(t)) \in \Gamma$ be the corresponding solution. If

$$\begin{aligned}
 R_1^s & = \frac{(1 - \epsilon)(1 - \eta)\beta k \Lambda}{c \delta_1 \delta_2} - \left[\frac{1}{2c\delta_1^2 (1 + \delta_2)^2} (1 - \epsilon)^2 (1 - \eta)^2 k^2 \Lambda^2 \sigma^2 \right] \\
 & = R_0^d - \left[\frac{1}{2c\delta_1^2 (1 + \delta_2)^2} (1 - \epsilon)^2 (1 - \eta)^2 k^2 \Lambda^2 \sigma^2 \right] > 1,
 \end{aligned}$$

then HCV will be persistent in mean, i.e.,

$$\liminf_{t \rightarrow \infty} \frac{1}{t} \int_0^t \{I(s) + V(s)\} ds \geq \frac{\delta_2 / (\delta_2 \vee c)}{(1 - \eta)(1 - \epsilon)\beta\Lambda k} [\delta_1 c (R_1^s - 1)].$$

Proof. Analogue to the proof technique in [40] and [44]. Integrating the system (2.2) yields

$$\begin{aligned} T(t) - T(0) + I(t) - I(0) + V(t) - V(0) &= \Lambda t - \delta_1 \int_0^t T(s) ds - \delta_2 \int_0^t I(s) ds \\ &+ (1 - \epsilon)k \int_0^t I(s) ds - c \int_0^t V(s) ds \end{aligned} \quad (3.8)$$

Dividing both sides Eq (3.8) by t , we get

$$\begin{aligned} \frac{T(t) - T(0)}{t} + \frac{I(t) - I(0)}{t} + \frac{V(t) - V(0)}{t} &= \Lambda - \frac{\delta_1}{t} \int_0^t T(s) ds - \frac{\delta_2}{t} \int_0^t I(s) ds \\ &+ \frac{(1 - \epsilon)k}{t} \int_0^t I(s) ds - \frac{c}{t} \int_0^t V(s) ds \\ \frac{1}{t} \int_0^t T(s) ds &= \frac{\Lambda}{\delta_1} - \frac{\delta_2}{\delta_1 t} \int_0^t I(s) ds + \frac{(1 - \epsilon)k}{\delta_1 t} \int_0^t I(s) ds - \frac{c}{\delta_1 t} \int_0^t V(s) ds - \Phi \end{aligned} \quad (3.9)$$

where

$$\Phi(t) = \frac{T(t) - T(0)}{t} + \frac{I(t) - I(0)}{t} + \frac{V(t) - V(0)}{t}.$$

Define a function $Q = \ln[(1 - \epsilon)kI(t) + (1 + \delta_2)V(t)]$. Applying Itô's formula, we obtain

$$\begin{aligned} dQ &= \frac{(1 - \epsilon)k[(1 - \eta)\beta VT - \delta_2 I]dt + \sigma(1 - \eta)VT dB}{(1 - \epsilon)kI + (1 + \delta_2)V} \\ &+ \frac{(1 + \delta_2)((1 - \epsilon)kI - cV)dt}{(1 - \epsilon)kI + (1 + \delta_2)V} + \frac{1}{2} \left[- \frac{(1 - \eta)^2(1 - \epsilon)^2 k^2 \sigma^2 (VT)^2}{[(1 - \epsilon)kI + (1 + \delta_2)V]^2} dt \right], \\ &\geq \left(\frac{(1 - \epsilon)(1 - \eta)k\beta T}{\delta_2} - c - \frac{1}{2} \frac{(1 - \eta)^2(1 - \epsilon)^2 k^2 \sigma^2 T^2}{(1 + \delta_2)^2} \right) dt \\ &+ \frac{\sigma(1 - \eta)(1 - \epsilon)kVT dB}{(1 - \epsilon)kI + (1 + \delta_2)V}. \end{aligned} \quad (3.10)$$

Integrating Eq (3.10) from 0 to t and dividing both sides by t , we have

$$\begin{aligned} \frac{1}{t} \ln[(1 - \epsilon)kI(t) + (1 + \delta_2)V(t)] &\geq \frac{1}{t} \ln[(1 - \epsilon)kI(0) + (1 + \delta_2)V(0)] \\ &+ \frac{1}{t} \int_0^t \frac{(1 - \epsilon)(1 - \eta)k\beta T(s) ds}{\delta_2} - c \\ &- \frac{1}{t} \int_0^t \frac{1}{2} \frac{(1 - \eta)^2(1 - \epsilon)^2 k^2 \sigma^2 T(s)^2 ds}{(1 + \delta_2)^2} \\ &+ \frac{1}{t} \int_0^t \frac{\sigma(1 - \eta)(1 - \epsilon)kV(s)T(s)dB(s)}{(1 - \epsilon)kI + (1 + \delta_2)V}. \end{aligned} \quad (3.11)$$

According to Eqs (3.9) and (3.11), we get

$$\frac{\delta_2}{\delta_1 t} \int_0^t I(s) ds + \frac{c}{\delta_1 t} \int_0^t V(s) ds \geq \frac{\delta_2}{(1-\eta)(1-\epsilon)\beta\Lambda k} [c(R_1^s - 1) - \Phi(t) - \Psi(t) + M(t)] \quad (3.12)$$

where

$$\begin{aligned} R_1^s &= \frac{(1-\epsilon)(1-\eta)\beta k \Lambda}{c\delta_1\delta_2} - \left[\frac{1}{2c\delta_1^2(1+\delta_2)^2} (1-\epsilon)^2(1-\eta)^2 k^2 \Lambda^2 \sigma^2 \right], \\ \Psi(t) &= \frac{1}{t} \ln[(1-\epsilon)kI(0) + (1+\delta_2)V(0)], \\ M(t) &= \frac{1}{t} \int_0^t \frac{\sigma(1-\eta)(1-\epsilon)kV(s)T(s)dB(s)}{(1-\epsilon)kI + (1+\delta_2)V}. \end{aligned}$$

Taking the limit inferior of Eq (3.12) as $t \rightarrow 0$ yields

$$\begin{aligned} \liminf_{t \rightarrow \infty} \frac{\delta_2}{\delta_1 t} \int_0^t I(s) ds + \frac{c}{\delta_1 t} \int_0^t V(s) ds &\geq \liminf_{t \rightarrow \infty} \frac{\delta_2}{(1-\eta)(1-\epsilon)\beta\Lambda k} [c(R_1^s - 1)], \\ \liminf_{t \rightarrow \infty} \frac{1}{t} \left[\int_0^t I(s) ds + \int_0^t V(s) ds \right] &\geq \liminf_{t \rightarrow \infty} \frac{\delta_2/(\delta_2 \vee c)}{(1-\eta)(1-\epsilon)\beta\Lambda k} [\delta_1 c(R_1^s - 1)]. \end{aligned}$$

This completes the proof. \square

4. Numerical simulation

In this section, we carry out a numerical simulation in order to provide an interpretation of the solution. We use the Euler-Maruyama method to determine the solution of system (2.2). The discretization of system (2.2) is given as follows

$$\begin{cases} T_{i+1} = T_i + (\Lambda - \delta_1 T_i - (1-\eta)\beta V_i T_i) \Delta t - \sigma(1-\eta)V_i T_i \sqrt{\Delta t} \zeta \\ I_{i+1} = I_i + ((1-\eta)\beta V_i T_i - \delta_2 I_i) \Delta t + \sigma(1-\eta)V_i T_i \sqrt{\Delta t} \zeta \\ V_{i+1} = V_i + ((1-\epsilon)kI_i - cV_i) \Delta t \end{cases} \quad (4.1)$$

where $t \in [t_0, t_N]$, $\Delta t = \frac{t_N - t_0}{N}$, and ζ is normally distributed $N(0,1)$.

Three simulations are conducted. The first simulation is deterministic system. The second simulation is to represent the solution of a stochastic model in the extinction condition. The third simulation is illustrating the solution of a stochastic model when Lemma 3.1 and Theorem 3.2 are violated. The parameters and initial values of the model in each simulation are given in Table 2.

Figure 1 shows the solution of a deterministic model ($\sigma = 0$) with $R_0^d > 1$. It can be seen that the solution is tend to the endemic equilibrium point $E_1 = (6.6 \times 10^6, 2.6 \times 10^6, 7 \times 10^5)$ which means that HCV disease is persistent.

Figure 2 illustrates the stochastic model with $\sigma = 4 \times 10^{-7}$ the condition

$$\sigma = \sqrt{1.6 \times 10^{-13}} = 4 \times 10^{-7} > \frac{\beta}{\sqrt{2(\delta_2 - (1-\epsilon)k \wedge c)}} = 0.76 \times 10^{-7}$$

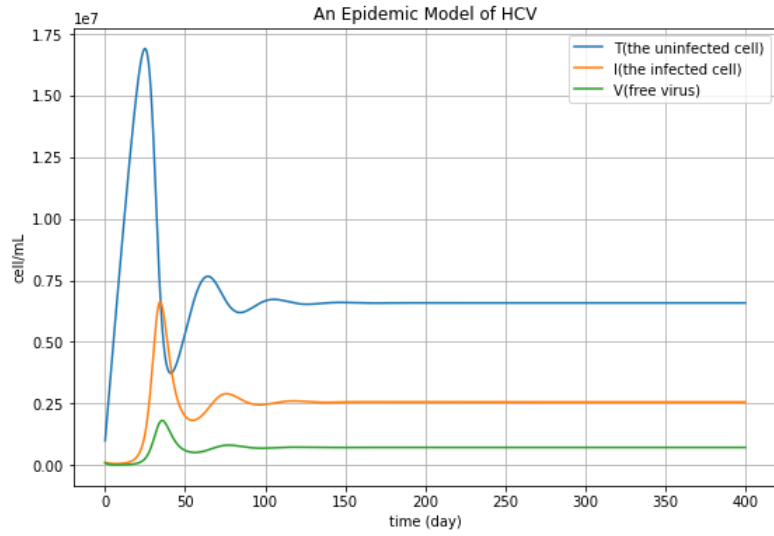
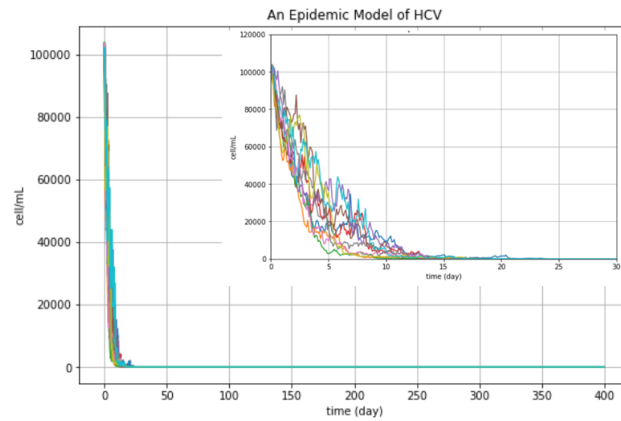
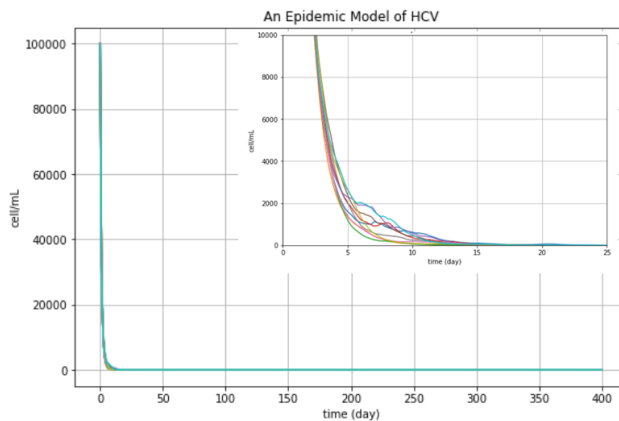


Figure 1. Simulation 1: a deterministic model of the system (2.1).



(a) infected cell



(b) free virus

Figure 2. Simulation 2: a stochastic model of the system (2.2) with $\sigma = 4 \times 10^{-7}$, 10 paths.

Table 2. Values and parameters of the model.

	Sim.1	Sim.2	Sim.3	Unit	Reference
$T(0)$	10^6	10^6	10^6	cell/mL	[45]
$I(0)$	10^5	10^5	10^5	cell/mL	[45]
$V(0)$	10^5	10^5	10^5	virion/mL	[45]
Λ	8×10^5	8×10^5	8×10^5	cell/day×mL	[17]
η	0.7	0.7	0.7	–	[17]
k	0.9	0.2	0.9	virion/cell×day	[17]
ϵ	0.75	0.75	0.75	–	[17]
β	5.4×10^{-8}	5.4×10^{-8}	5.4×10^{-7}	mL/virion × day	[17]
δ_1	0.0047	0.1	0.0047	1/day	[17, 46]
δ_2	0.3	0.3	0.3	1/day	[17]
c	0.8	1	0.8	1/day	[17]
σ	–	4×10^{-7}	9×10^{-8}	–	–

and $R_0^s < 1$ where $\sigma^2 = 1.6 \times 10^{-13} < \frac{4\delta_1\delta_2\beta}{(1-\epsilon)(1-\eta)k\Lambda} = 5.4 \times 10^{-13}$ satisfies Lemma 3.1 and Theorem 3.2 respectively. More precisely, when the chosen parameters satisfy Lemma 3.1 and Theorem 3.2, the point $E_0 = (8 \times 10^6, 0, 0)$ is almost exponentially stable. It can be seen from Figure 2 where the concentration of infected cells and free viruses go to zero. Therefore, HCV will go extinct almost surely.

Figure 3 describes the condition when basic reproduction number $R_0^s = 0.0214 < 1$ and $\sigma^2 = 4.9 \times 10^{-15} < \frac{4\delta_1\delta_2\beta}{(1-\epsilon)(1-\eta)k\Lambda} = 5.4 \times 10^{-13}$. This is satisfying Theorem 3.2 and violating Lemma 3.1. That figure shows a stochastic noise impact the disease extinction.

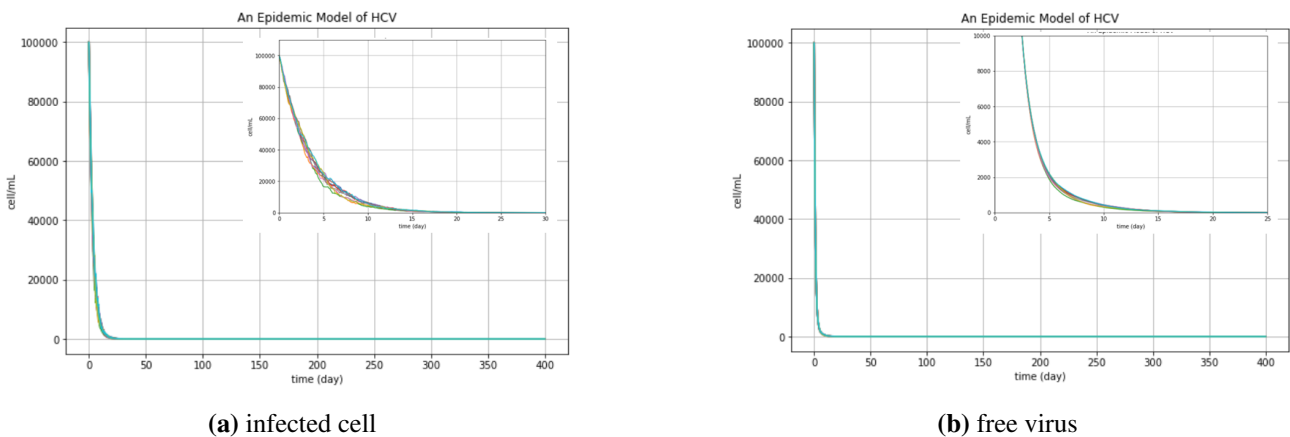


Figure 3. Simulation 2: a stochastic model of the system (2.2) with $\sigma = 7 \times 10^{-8}$, 10 paths

Figure 4 shows that if the intensity of the disturbance is smaller where $\sigma^2 = 8.1 \times 10^{-16} < 1.9 \times 10^{-12}$, thus Lemma 3.1 is violated. Taking the time interval $[0, 400]$ with 4000 points and 10 paths, it appears in Figure 4(a) that the concentration of uninfected cells rises to a peak around at $t = 30$ days, then drops sharply to around $t = 45$ days and then they rise slowly. The graphics in Figures 4(b),(c) show the concentration of infected cells and free virus after $t = 50$ days which tend to certain interval values. This shows that hepatitis C remains exist.

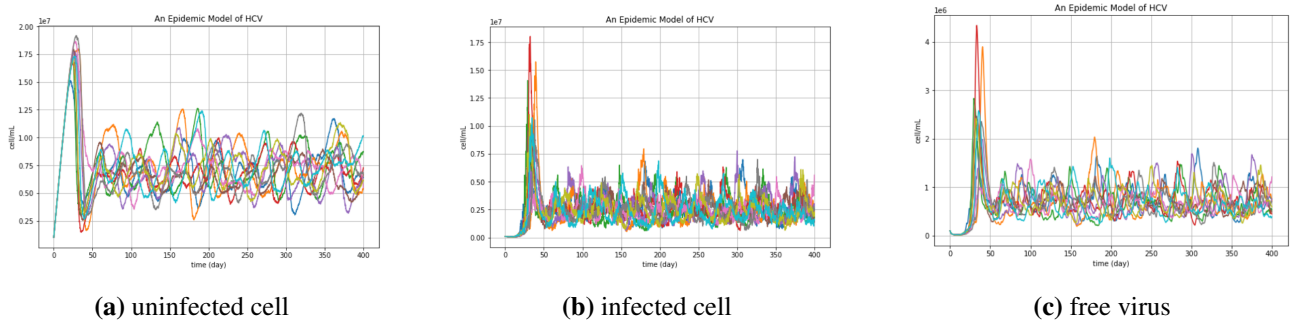


Figure 4. Simulation 3: a stochastic model of the system (2.2) with $\sigma = 9 \times 10^{-8}$, 10 paths.

Figure 5 describes the condition of the HCV extinction when Lemma 3.1 is satisfied while Theorem 3.2 is violated. It can be seen that figures show the concentration of infected cells and free virus from initial value that tends to zero for a long period of time.

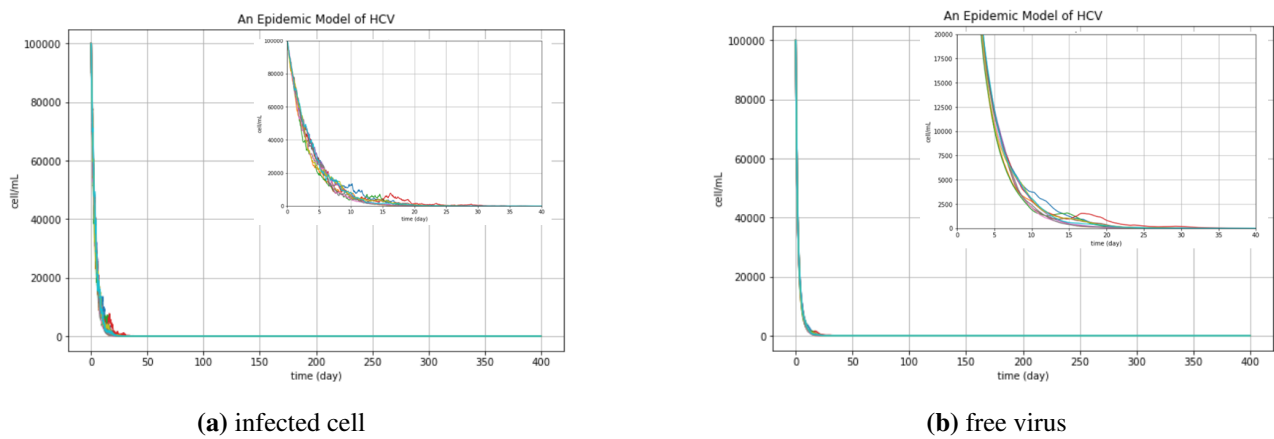


Figure 5. Simulation 3: a stochastic model of the system (2.2) with $\beta = 2.5 \times 10^{-8}$, 10 paths.

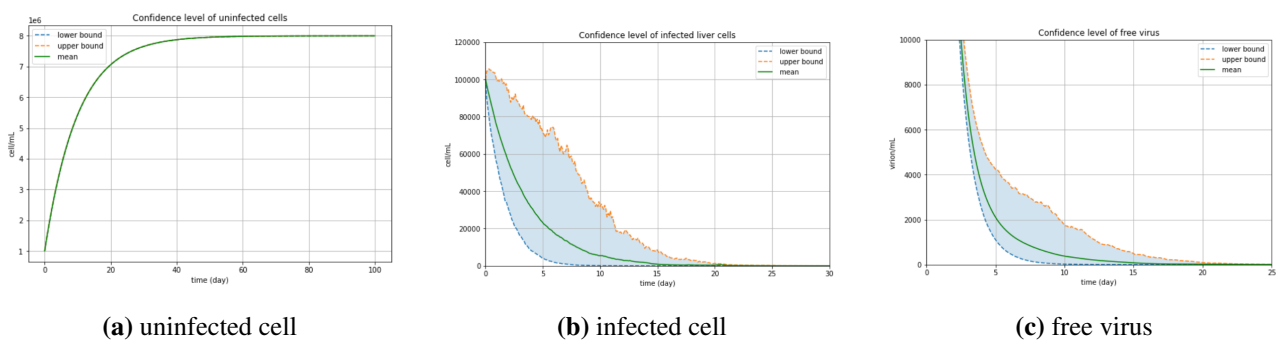


Figure 6. A stochastic model of the system (2.2) with confidence interval on simulation 2.

Figure 6(a) shows the distribution value of uninfected cells in simulation 2 with a confidence interval of 95 %. After time t about 50 days the concentration of uninfected cells will go to a positive constant value (8×10^6). From Figure 6(b), the simulation at $t = 5$ is more diverse than $t = 10$. After time t 20 days, the concentration of infected cells will go to zero. Figure 6(c) shows the free virus concentration at the time t from 3 to 15 days is in the certain interval value. After time t 20 days, the virus concentration will

go to zero. Therefore, from simulation 2, it can be concluded that stochastic disturbances can affect the behavior of the spread of hepatitis C. When conditions are met according to Lemma 3.1 and Theorem 3.2, at a certain time t hepatitis C will disappear. Mathematically, in this case, the basic reproduction number in the extinction condition is less than one.

Figure 7 shows the confidence intervals for simulation 3 where we reduced the sigma values while preserving the other parameter values. Figure 7(a) shows the concentration of uninfected cells will reach peaks around 23 to 30 days and then decline. After 75 days the concentration will move to a certain interval value. The green lines show the mean value of the simulations. Figures 7(b),(c) show that after 75 days, the concentration of infected cells and free virus lead to a certain positive number interval value with a confidence interval of approximately 95 %. In other words, hepatitis C disease will persist.

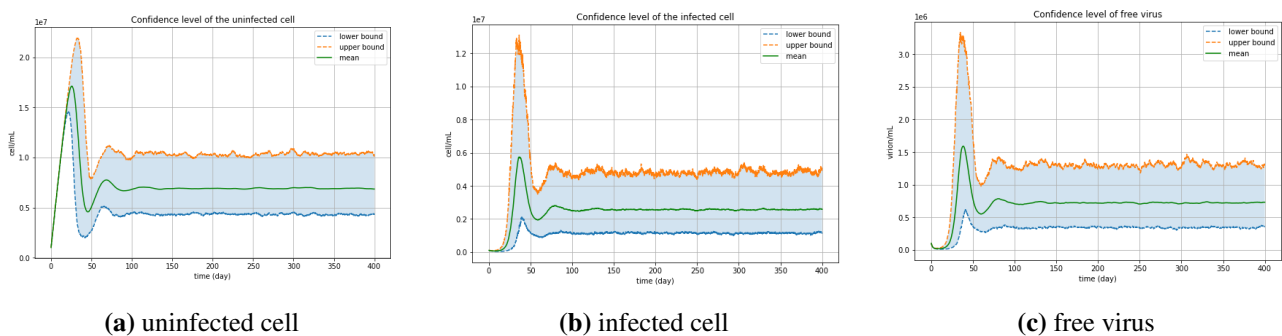


Figure 7. A stochastic model of the system (2.2) with confidence interval on simulation 2.

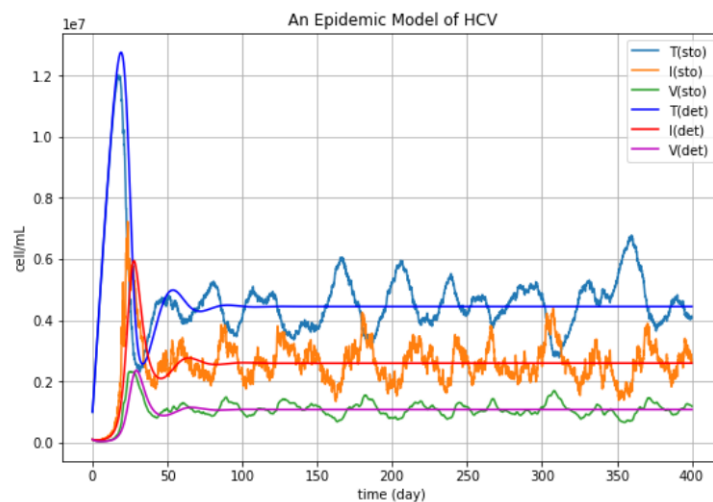


Figure 8. A stochastic model of the system (2.2) on persistence in mean simulation compared to a deterministic.

Lastly, we conduct a simulation for persistence in mean condition where $\beta = 5.4 \times 10^{-7}$, $\delta_1 = 0.0047$, $k = 1$, $c = 0.6$, $\sigma = 9 \times 10^{-8}$, and the rest of parameters and initial values state in Table 2. Figure 8 illustrates the solution of the stochastic model compared to the deterministic. Figure 9 shows the number of concentrations of healthy cells, infected cells, and free virus at a certain value which are

more than zero. When $t \geq 50$, it appears that the concentrations of uninfected cells, infected cells and free viruses go to an equilibrium state. According to the numerical calculations, we obtain $R_1^s > 1$ and the endemic equilibrium point $E_1 = (T, I, V) = (4.4 \times 10^6, 2.6 \times 10^6, 1.08 \times 10^6)$. According to Theorem 3.4 hepatitis C remains to exist which are also depicted in Figure 9. Figure 10 describes the condition if the noise intensity is increasing, then the solution of model (2.2) will be strongly oscillating around the endemic equilibrium point of the model (2.1).

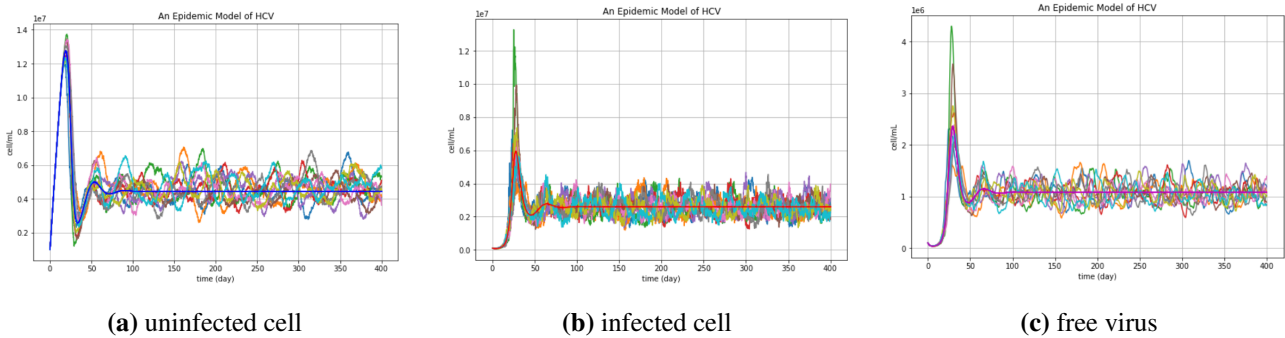


Figure 9. A stochastic model of the system (2.2) with $\sigma = 9 \times 10^{-8}$, 10 paths on simulation persistence in mean.

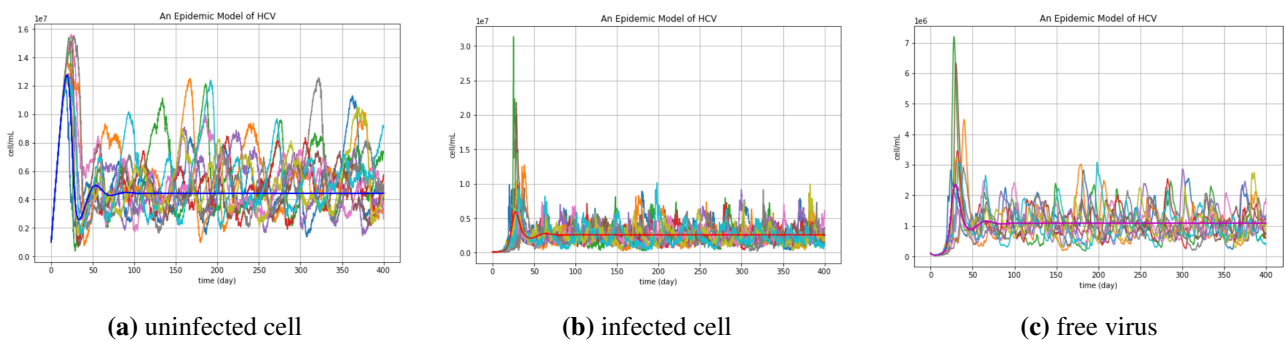


Figure 10. A stochastic model of the system (2.2) with $\sigma = 2 \times 10^{-7}$, 10 paths on simulation persistence in mean.

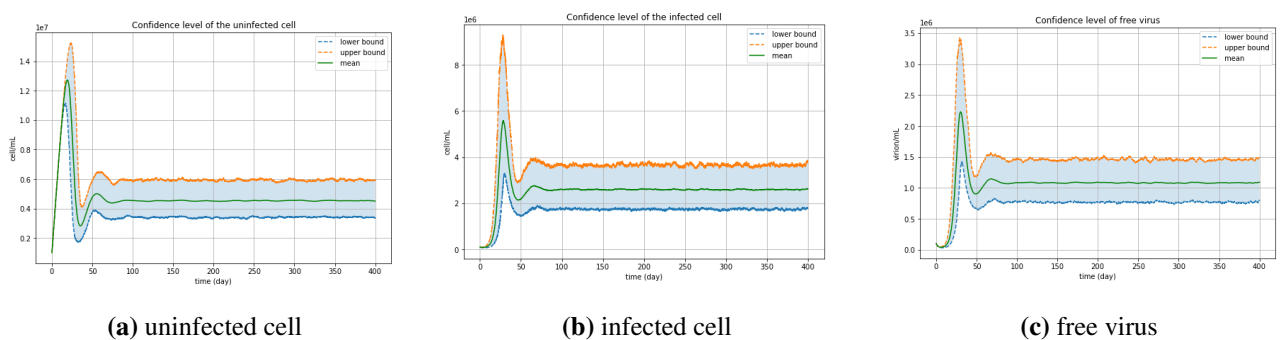


Figure 11. A stochastic model of the system (2.2) with confidence interval on persistence in mean simulation.

Furthermore, in Figure 11, we generate a 95% confidence interval for $t \in [0, 400]$ with 4000 paths.

Figure 11(a) describes the number of uninfected cells that tends to a consistent interval value after 75 days. Figure 11(b),(c) show the confidence intervals of infected liver cells and free viruses. In the time t span of 0 to 400 days the number of concentrations in a row increases to a peak around day 27 for infected cells and free viruses, then decreases. After 30 days, leading to a consistent value for infected cells from 1.9×10^6 to 3.9×10^6 , and free viruses from 0.75×10^6 to 1.5×10^6 .

5. Conclusions

In this paper, we propose an epidemic model for HCV transmission at the cellular level incorporating statistical noise. The results extend the paper of Neumann, et.al. [24] in understanding the dynamics of a deterministic hepatitis C virus. Various types of Lyapunov functions are designed to study the persistence in mean conditions as well as the extinction of the stochastic system. Based on this model, there exists a unique time-global solution for any given positive initial value. In addition, numerical simulations are carried out to describe the solution behaviour of the model. We analyze that if the noise intensity is increasing, then the disease will go to extinction. If the basic reproduction number of the extinction condition is less than one, then hepatitis C will extinct. If the basic reproduction number of the persistence in the mean condition is more than one, then hepatitis C remains to exist. When the noise intensity increases, the solution will give a strong oscillation under the condition of persistence in the mean. It can provide some useful strategies for controlling the dynamics of that disease. In future research, we can consider an optimal control of the stochastic model for HCV, research on memory effect, and fractional derivatives [47]. Furthermore, it needs to investigate the stochastic noise for other parameters apart from infection rate. It is also important to consider a stochastic model for HCV transmission involving the response immune system.

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Conflict of interest

The authors declare that they have no competing financial interest or personal relationships that could have appeared to influence the work reported in this paper.

References

1. C. J. Burrell, C. R. Howard, F. A. Murphy, Viral Syndromes, *Fenner White Med. Virol.*, **2017** (2017), 537–556. doi: 10.1016/B978-0-12-375156-0.00039-4.

2. World Health Organization (WHO), *Guidelines for the Screening Care and Treatment of Persons with Chronic Hepatitis C Infection Updated Version April 2016: Guidelines*, 2016. Available from: <https://apps.who.int/iris/handle/10665/205035>.
3. S. W. Wibawa, *Mengenal hepatitis C, Infeksi Bisu Yang Menghantui Indonesia*, 2017. Available from: <https://sains.kompas.com/read/2017/08/16/221249323/mengenal-hepatitis-c-infeksi-bisu-yang-menghantui-indonesia?page=all>.
4. Ministry of Health Republic of Indonesia, *Guidelines for Controlling Viral Hepatitis*, 2012. Available from: <http://hukor.kemkes.go.id/>.
5. World Health Organization (WHO), *Hepatitis C*, 2021. Available from: <https://www.who.int/news-room/fact-sheets/detail/hepatitis-c>.
6. Alhawaris, Hepatitis C: Epidemiologi, Etiologi, dan Patogenitas, *Jurnal Sains Dan Kesehatan*, **2** (2019), 139–150. doi: 10.25026/jsk.v2i2.132.
7. R. Sanju´an, P. Domingo-Calap, Mechanisms of viral mutation, *Cell. Mol. Life Sci.*, **73** (2016), 4433–4448. doi: 10.1007/s00018-016-2299-6.
8. I. Dontwi, N. Frempong, D. Bentil, I. Adetunde, E. Owusu-Ansah, Mathematical modeling of hepatitis C virus transmission among injecting drug users and the impact of vaccination, *Am. J. Sci. Ind. Res.*, **1** (2010), 41–46.
9. D. Lestari, L. Candrawati, Global stability of SACR epidemic model for hepatitis C on injecting drug users, in *Proceeding of 3rd International Conference on Research, Implementation and Education of Mathematics and Science*, 2016. Available from: <http://seminar.uny.ac.id/icriems/sites/seminar.uny.ac.id/icriems/files/prosiding/M-21.pdf>.
10. M. Imran, M. Hassan, M. Dur-E-Ahmad, A. Khan, A comparison of a deterministic and stochastic model for hepatitis C with an isolation stage, *J. Biol. Dyn.*, **7** (2013), 276–301. doi: 10.1080/17513758.2013.859856.
11. A. B. Pitcher, A. Borquez, B. Skaathun, N. K. Martin, Mathematical modeling of hepatitis C virus (HCV) prevention among people who inject drugs: A review of the literature and insights for elimination strategies, *J. Theor. Biol.*, **481** (2019), 194–201. doi: 10.1016/j.jtbi.2018.11.013.
12. E. H. Elbasha, Model for hepatitis C virus transmissions, *Math. Biosci. Eng.*, **10** (2013), 1045–1065. doi: 10.3934/mbe.2013.10.1045.
13. A. G. Lim, H. Qureshi, H. Mahmood, S. Hamid, C. F. Davies, A. Trickey, et.al., Curbing the hepatitis C virus epidemic in Pakistan: the impact of scaling up treatment and prevention for achieving elimination, *Int. J. Epidemiol.*, **47** (2018), 550–560. doi: 10.1093/ije/dyx270.
14. M. Shen, Y. Xiao, W. Zhou, Z. Li, Global dynamics and applications of an epidemiological model for hepatitis c virus transmission in China, *Discrete Dyn. Nat. Soc.*, **2015** (2015), 1–13. doi: 10.1155/2015/543029.
15. J. Khodaei-Mehr, S. Tangestanizadeh, M. Sharifi, R. Vatankhah, M. Eghtesad, Hepatitis C virus epidemic control using a nonlinear adaptive strategy, preprint, arXiv:2007.13522.
16. R. Avendan˜o, L. Esteva, J. Flores, J. F. Allen, G. G´omez, J. L´opez-Estrada, A mathematical model for the dynamics of hepatitis C, *J. Theor. Med.*, **4** (2002), 109–118. doi: 10.1080/10273660290003777.

17. H. Dahari, J. E. Layden-Almer, E. Kallwitz, R. M. Ribeiro, S. J. Cotler, T. J. Layden, et al., A mathematical model of hepatitis C virus dynamics in patients with high baseline viral loads or advanced liver disease, *Gastroenterology*, **136** (2009), 1402–1409. doi: 10.1053/j.gastro.2008.12.060.
18. J. Guedj, L. Rong, H. Dahari, A. S. Perelson, A perspective on modelling hepatitis C virus infection, *J. Viral Hepatitis*, **17** (2010), 825–833. doi: 10.1111/j.1365-2893.2010.01348.x.
19. I. Zada, M. N. Jan, N. Ali, D. Alrowail, K. S. Nisar, G. Zaman, Mathematical analysis of hepatitis B epidemic model with optimal control, *Adv. Differ. Equations*, **1** (2021), 1–29. doi: 10.1186/s13662-021-03607-2.
20. T. S. Shaikh, N. Fayyaz, N. Ahmed, N. Shahid, M. Rafiq, I. Khan, et al., Numerical study for epidemic model of hepatitis-B virus, *Eur. Phys. J. Plus*, **136** (2021), 1–22. doi: 10.1140/epjp/s13360-021-01248-8.
21. J. E. Pearson, P. Krapivsky, A. S. Perelson, Stochastic theory of early viral infection: continuous versus burst production of virions, *PLoS Comput. Biol.*, **7** (2011), e1001058. doi: 10.1371/journal.pcbi.1001058.
22. T. Nguyen, J. Guedj, HCV kinetic models and their implications in drug development, *CPT Pharmacometrics Syst. Pharmacol.*, **4** (2015), 231–242. doi: 10.1002/psp4.28.
23. Indonesian Heart Research Association (Perhimpunan Peneliti Hati Indonesia/PPHI), *Consensus on the Management of Hepatitis C in Indonesia, Jakarta: PPHI*, 2017.
24. A. U. Neumann, N. P. Lam, H. Dahari, D. R. Gretch, T. E. Wiley, T. J. Layden, et al., Hepatitis C viral dynamics in vivo and the antiviral efficacy of interferon-alpha therapy, *Science*, **282** (1998), 103–107. doi: 10.1126/science.282.5386.103.
25. L. J. S. Allen, G. E. Lahodny, Extinction thresholds in deterministic and stochastic epidemic models, *J. Biol. Dyn.*, **6** (2012), 590–611. doi: 10.1080/17513758.2012.665502.
26. T. Feng, Z. Qiu, X. Meng, Dynamics of a stochastic hepatitis C virus system with host immunity, *Discrete Contin. Dyn. Syst. B*, **24** (2019), 6367–6385. doi: 10/3934/dcdsb.2019143.
27. C. J. Mode, C. K. Sleeman, Stochastic processes in epidemiology: HIV/AIDS, other infectious diseases and computers, in *World Scientific*, (2000).
28. M. Merdan, Z. Bekiryazici, T. Kesemen, T. Khaniyev, Deterministic stability and random behavior of a hepatitis C model, *PloS One*, **12** (2017), e0181571. doi: 10.1371/journal.pone.0181571.
29. Z. U. A. Zafar, N. Ali, S. Younas, S. F. Abdelwahab, K. S. Nisar, Numerical investigations of stochastic HIV/AIDS infection model, *Alexandria Eng. J.*, **60** (2021), 5341–5363. doi: 10.1016/j.aej.2021.04.027.
30. M. A. Noor, A. Raza, M. S. Arif, M. Rafiq, K. S. Nisar, I. Khan, et al., Non-standard computational analysis of the stochastic Covid-19 pandemic model: an application of computational biology, *Alexandria Eng. J.*, **61** (2022), 619–630. doi: 10.1016/j.aej.2021.06.039.
31. K. Hattaf, M. Mahrouf, J. Adnani, N. Yousfi, Qualitative analysis of a stochastic epidemic model with specific functional response and temporary immunity, *Phys. A*, **490** (2018), 591–600. doi: 10.1016/j.physa.2017.08.043.
32. M. Mahrouf, K. Hattaf, N. Yousfi, Dynamics of a stochastic viral infection model with immune response, *Math. Modell. Nat. Phenom.*, **12** (2017), 15–32.

33. P. Van den Driessche, J. Watmough, Reproduction numbers and sub-threshold endemic equilibria for compartmental models of disease transmission, *Math. Biosci.*, **180** (2002), 29–48. doi: 10.1016/S0025-5564(02)00108-6.
34. B. Øksendal, *Stochastic differential equations: an introduction with applications*, Springer, 2003.
35. G. Hu, K. Wang, Existence and uniqueness theorem for stochastic differential equations with self-exciting switching, *Discrete Dyn. Nat. Soc.*, **2011** (2011), 1–12. doi: 10.1155/2011/549651.
36. Y. Zhou, W. Zhang, S. Yuan, Survival dan stationary distribution of a SIR epidemic model with stochastic perturbations, *Appl. Math. Comput.*, **244** (2014), 118–131. doi: 10.1016/j.amc.2014.06.100.
37. X. Mao, G. Marion, E. Renshaw, Environmental brownian noise suppresses explosions in population dynamics, *Stochastic Processes Their Appl.*, **97** (2002), 95–110. doi: 10.1016/S0304-4149(01)00126-0.
38. L. Wang, H. Huang, A. Xu, W. Wang, Stochastic extinction in an SIRS epidemic model incorporating media coverage, *Abstr. Appl. Anal.*, **2013** (2013), 1–8. doi: 10.1155/2013/891765.
39. F. Rao, Dynamics analysis of a stochastic SIR epidemic model, *Abstr. Appl. Anal.*, **2014** (2014), 1–9. doi: 10.1155/2014/356013.
40. Y. Zhang, Y. Li, Q. Zhang, A. Li, Behavior of a stochastic SIR epidemic model with saturated incidence and vaccination rules, *Phys. A*, **501** (2018), 178–187. doi: 10.1016/j.physa.2018.02.191.
41. X. Mao, *Stochastic Differential Equations and Applications*, Elsevier, 2007.
42. T. Feng, Z. Qiu, Global analysis of a stochastic TB model with vaccination and treatment, *Discrete Contin. Dyn. Syst. B*, **24** (2019), 2923–2939. doi: 10.3934/dcdsb.2018292.
43. Q. Liu, D. Jiang, N. Shi, T. Hayat, A. Alsaedi, Dynamics of a stochastic tuberculosis model with constant recruitment and varying total population size, *Phys. A*, **469** (2017), 518–530. doi: 10.1016/j.physa.2016.11.053.
44. B. Boukanjime, M. E. Fatini, A stochastic hepatitis B epidemic model driven by Lévy noise, *Phys. A*, **521** (2019), 796–806. doi: 10.1016/j.physa.2019.01.097.
45. L. Rong, R. M. Ribeiro, A. S. Perelson, Modeling quasispecies and drug resistance in hepatitis C patients treated with a protease inhibitor, *Bull. Math. Biol.*, **74** (2012), 1789–1817. doi: 10.1007/s11538-012-9736-y.
46. D. Wodarz, Hepatitis c virus dynamics and pathology: the role of CTL and antibody responses, *J. Gen. Virol.*, **84** (2003), 1743–1750.
47. K. Hattaf, A new generalized definition of fractional derivative with non-singular kernel, *Computation*, **8** (2020), 49. doi: 10.3390/computation8020049.



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