

THE THRESHOLD OF A STOCHASTIC SIRS EPIDEMIC MODEL IN A POPULATION WITH VARYING SIZE

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ABSTRACT. In this paper, a stochastic susceptible-infected-removed-susceptible (SIRS) epidemic model in a population with varying size is discussed. A new threshold \tilde{R}_0 is identified which determines the outcome of the disease. When the noise is small, if $\tilde{R}_0 < 1$, the infected proportion of the population disappears, so the disease dies out, whereas if $\tilde{R}_0 > 1$, the infected proportion persists in the mean and we derive that the disease is endemic. Furthermore, when $R_0 > 1$ and subject to a condition on some of the model parameters, we show that the solution of the stochastic model oscillates around the endemic equilibrium of the corresponding deterministic system with threshold R_0 , and the intensity of fluctuation is proportional to that of the white noise. On the other hand, when the noise is large, we find that a large noise intensity has the effect of suppressing the epidemic, so that it dies out. These results are illustrated by computer simulations.

1. Introduction. Studies of epidemic models that incorporate disease-caused death and a varying total population have become one of the important areas in the mathematical theory of epidemiology, largely inspired by the works of Anderson and May (see [1], [2]).

Much of the classical work on epidemiological models has been restricted to situations where the affected population is of constant size. This assumption is relatively valid for diseases of short duration with limited effects on mortality. However, it

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clearly fails to hold for diseases that are endemic in communities with changing populations, and for diseases which raise the mortality rate substantially. Well-known examples of such diseases are malaria in developing countries with growing populations, and the current AIDS pandemic. In such situations, the effects of the disease-induced mortality and of the change in population size are far from negligible, and in fact, may have a crucial influence on whether or not the disease can reach epidemic levels.

In recent years there have been a number of studies on disease transmission models in populations of varying size, and some of these have given a complete global analysis of the model equations. We briefly note the work on such models that is related to ours as including the papers [3, 5, 7, 10, 15, 16, 17, 19].

Busenberg and van den Driessche (1990) [6] discussed a SIRS epidemic model in a population with varying size. The system has the following form:

$$\begin{cases} \dot{S}(t) = bN - \mu S - \frac{\beta SI}{N} + \delta R, \\ \dot{I}(t) = \frac{\beta SI}{N} - (\mu + \alpha + \gamma)I, \\ \dot{R}(t) = \gamma I - (\mu + \delta)R. \end{cases} \quad (1)$$

Here S , I and R denote the total numbers of susceptible, infective and recovered (removed) individuals respectively, and N is the population size. We set $N(t) = S(t) + I(t) + R(t)$ at time t , and $\dot{S}(t)$, $\dot{I}(t)$ and $\dot{R}(t)$ are the corresponding rates of change with respect to time t . We use the following additional notation:

- b : per capita birth rate,
- μ : per capita disease free death rate,
- β : effective per capita contact rate of infective individuals,
- δ : per capita loss of immunity rate of recovered individuals,
- α : excess per capita death rate of infected individuals,
- γ : per capita recovery rate of infected individuals.

All parameter values are assumed to be nonnegative and $b, \gamma > 0$. Note that the system (1) reduces to the SIR model when $\delta = 0$, so there is no loss of immunity. The equation for the total population size is $\dot{N}(t) = (b - \mu)N - \alpha I$. Busenberg and van den Driessche (1990) [6] consider the proportions of individuals in the three epidemiological classes, namely

$$x = \frac{S}{N}, \quad y = \frac{I}{N}, \quad \text{and} \quad z = \frac{R}{N} \quad (2)$$

It is easy to verify that x , y and z satisfy the system of differential equations

$$\begin{cases} \dot{x}(t) = b - bx - \beta xy + \delta z + \alpha xy, \\ \dot{y}(t) = \beta xy - (b + \alpha + \gamma)y + \alpha y^2, \\ \dot{z}(t) = \gamma y - (b + \delta)z + \alpha yz. \end{cases} \quad (3)$$

The feasibility region becomes $\Gamma = \{x \geq 0, y \geq 0, z \geq 0, x + y + z = 1\}$ and $\Gamma_0 = \Gamma - \{(1, 0, 0)\}$. There are two distinct ways of considering a disease as being brought under control in a population of increasing or decreasing total size. The stricter way requires that the total number of infectives $I(t) \rightarrow 0$, while a weaker requirement is that the proportion $y(t) \rightarrow 0$. This distinction is discussed in some detail in Busenberg et al. (1991) [7]. Thus, they showed the conditions for the existence and stability of the endemic proportion steady state (x^*, y^*, z^*) and for

the stability of the disease-free steady state $(x, y, z) = (1, 0, 0)$. The threshold parameter of the system is

$$R_0 = \frac{\beta}{b + \alpha + \gamma} \quad (4)$$

which determines the extinction and persistence of the epidemic.

According to the theory in Busenberg and van den Driessche (1990) [6], then

(a) The disease free equilibrium proportion $(x, y, z) = (1, 0, 0)$ always exists, and is globally asymptotically stable in the feasibility region Γ whenever $R_0 \leq 1$, and it is unstable when $R_0 > 1$.

(b) When $R_0 > 1$, there exists a unique endemic proportion equilibrium $(x, y, z) = (x^*, y^*, z^*)$ with $y^* > 0, z^* > 0$, which is globally asymptotically stable in Γ_0 .

However the deterministic approach has some limitations in the mathematical modelling transmission of an infectious disease. Stochastic differential equation (SDE) models play a significant role in various branches of applied sciences including infectious dynamics, as they provide some additional degree of realism compared to their deterministic counterpart. Recently, Many authors have introduced parameter perturbation into epidemic models and have studied their dynamics [4, 8, 21, 22].

Using stochastic differential equation models is one way to do this. However, compared to deterministic systems, it is extremely difficult to give the threshold of stochastic systems. Recently, Gray et al. [11] discussed the following stochastic SIS epidemic model

$$\begin{cases} dS(t) = [\mu N - \beta S(t)I(t) + \gamma I(t) - \mu S(t)]dt - \sigma S(t)I(t)dB(t), \\ dI(t) = [\beta S(t)I(t) - (\mu + \gamma)I(t)]dt + \sigma S(t)I(t)dB(t), \end{cases} \quad (5)$$

and examined its dynamics. Here $B(t)$ is a standard Brownian motion with $B(0) = 0$, the white noise intensity is $\sigma^2 > 0$, and μ, β and γ are as above. In this model S and I denote the numbers of susceptible and infected individuals in the population, respectively. As $S(t) + I(t) \equiv N$, the authors simplified system (5) into a single equation, and showed that

- If $R_0^S < 1$ and $\sigma^2 \leq \beta/N$, then the disease will die out with probability one.
- If $R_0^S > 1$, then the disease will be persistent, where $R_0^S = R_0^D - \sigma^2 N^2 / 2(\mu + \gamma)$ and $R_0^D = \beta N / \mu + \gamma$ is the threshold of the corresponding deterministic model. The quantity R_0^S can be considered as the threshold of system (5), which is less than the value of R_0^D . The results were illustrated by computer simulations.

Tornatore, Buccellato and Vetro [18] discussed a stochastic SIR system, given by

$$\begin{cases} dS(t) = [\mu - \mu S(t) - \beta S(t)I(t)]dt - \sigma S(t)I(t)dB(t), \\ dI(t) = [\beta S(t)I(t) - (\mu + \gamma)I(t)]dt + \sigma S(t)I(t)dB(t), \\ dR(t) = [\gamma I(t) - \mu R(t)]dt. \end{cases} \quad (6)$$

They proved that $0 < \beta < \min\{\lambda + \mu - \sigma^2/2, 2\mu\}$ is a sufficient condition for the asymptotic stability of the disease-free equilibrium (DFE). Their computer simulations for the SDE SIR model agree well with the analytical results and show that the introduction of noise into the system raises the threshold to $\mu + \gamma + \sigma^2/2$, so if

$$\min\{\mu + \gamma - \frac{\sigma^2}{2}, 2\mu\} < \beta < \mu + \gamma + \frac{\sigma^2}{2},$$

then the DFE $E_0 = (S(0), I(0), R(0)) = (1, 0, 0)$ is stable and the disease does not occur, whereas if $\beta > \mu + \gamma + \sigma^2/2$, then the DFE is unstable.

Ji, Jiang and Shi [13] discussed system (6) further, and obtained the results that

• If $\beta < \gamma + \mu - \sigma^2/2$, then the disease-free equilibrium $(1, 0, 0)$ of system (6) is stochastically asymptotically stable in the large and is exponentially mean-square stable.

• If $\beta > \gamma + \mu$, then the solution of system (6) fluctuates around (S^*, I^*, R^*) , which is the endemic equilibrium of the corresponding deterministic system. They consider that the disease will prevail if the white noise is small, but they did not give an accurate threshold [13, 18].

Recently Yang and Mao [20] studied a class of multi-group SEIR epidemic models with stochastic perturbations. By the method of stochastic Lyapunov functions, they studied the extinction and recurrence in terms of the intensity of the stochastic perturbations and the reproductive number R_0 .

In this paper, we suppose that some stochastic environmental factor acts simultaneously on each individual in the population. In this case β changes to a random variable $\tilde{\beta}$, as in [11]. More precisely each infected individual makes

$$\tilde{\beta}dt = \beta dt + \sigma dB(t)$$

potentially infectious contacts with each other individual in $[t, t + dt)$. Here $dB(t) = B(t + dt) - B(t)$ is the increment of a standard Brownian motion. Thus the number of potentially infectious contacts that a single infected individual makes with another individual in $[t, t + dt)$ is normally distributed with mean βdt and variance $\sigma^2 dt$. Hence $E(\tilde{\beta}dt) = \beta dt$ and $var(\tilde{\beta}dt) = \sigma^2 dt$. As $var(\tilde{\beta}dt) \rightarrow 0$ as $dt \rightarrow 0$ this is a biologically reasonable model. Indeed this is a well-established way of introducing stochastic environmental noise into biologically realistic population dynamic models.

The stochastic model corresponding to the deterministic SIRS model (1) then takes the following form, as an SDE model:

$$\begin{cases} dS = (bN - \mu S - \frac{\beta SI}{N} + \delta R)dt - \sigma \frac{SI}{N} dB(t), \\ dI = [\frac{\beta SI}{N} - (\mu + \alpha + \gamma)I]dt + \sigma \frac{SI}{N} dB(t), \\ dR = [\gamma I - (\mu + \delta)R]dt. \end{cases} \quad (7)$$

The equation for the total population N size is obtained from (7) as:

$$\frac{dN}{dt} = (b - \mu)N - \alpha I,$$

as in model (1). When the proportions x, y, z given by (2) are used as the independent variables, the system (7) becomes

$$\begin{cases} dx = (b - bx - \beta xy + \delta z + \alpha xy)dt - \sigma xy dB(t), \\ dy = [\beta xy - (b + \alpha + \gamma)y + \alpha y^2]dt + \sigma xy dB(t), \\ dz = [\gamma y - (b + \delta)z + \alpha yz]dt. \end{cases} \quad (8)$$

We will try to give a threshold of the system (8), which can easily determine the extinction and persistence of the disease. Using the relation $x = 1 - y - z$, we can omit analysis of the first equation of system (8), and so we discuss the following system

$$\begin{cases} dy = [\beta(1 - y - z)y - (b + \alpha + \gamma)y + \alpha y^2]dt + \sigma(1 - y - z)y dB(t), \\ dz = [\gamma y - (b + \delta)z + \alpha yz]dt, \end{cases} \quad (9)$$

with any given initial value $(y(0), z(0)) \in \mathbb{R}_+^2$ and $y(0) + z(0) < 1$.

This paper is organized as follows. In Section 2, we show there is a unique positive solution of system (7) using the method mentioned in [9, 11]. In Section 3, we deduce the conditions which will cause the disease to die out. The condition for the disease being persistent is given in Section 4. In Section 5, when $R_0 > 1$ and α^2 is sufficiently small, we derive that the solution of system (9) oscillates around the endemic proportion equilibrium $P^*(y^*, z^*)$, and the intensity of fluctuation is proportional to the size of the white noise. The key to our analysis is choosing an appropriate Lyapunov function. Section 6 gives a short conclusion. Throughout the paper, outcomes of numerical simulations are shown in support of our analytical results.

Throughout this paper, unless otherwise specified, let $(\Omega, \{\mathcal{F}_t\}_{t \geq 0}, P)$ be a complete probability space with a filtration $\{\mathcal{F}_t\}_{t \geq 0}$ satisfying the usual conditions (i.e. it is right continuous and \mathcal{F}_0 contains all P-null sets), and let $B(t)$ be a scalar Brownian motion defined on this probability space.

2. Existence and uniqueness of positive solution. To investigate the dynamical behaviour of a population model, the first concern is whether the solution of the model is positive and global. As we know, in order to get a stochastic differential equation for which a unique global solution exists (i.e. there is no explosion within a finite time) for any initial value, the coefficients of the equation should satisfy the linear growth condition and the local Lipschitz condition (cf. Mao [14]). However, the coefficients of system (7) do not satisfy the linear growth condition (as the incidence is nonlinear), so the solution of system (7) may explode at a finite time. In this section, using the Lyapunov analysis method (as mentioned in [9, 11]), we show that the solution of system (7) is positive and global.

Theorem 2.1. *There is a unique solution $(S(t), I(t), R(t))$ of system (7) on $t \geq 0$ for any initial value $(S(0), I(0), R(0)) \in \mathbb{R}_+^3$, and the solution will remain in \mathbb{R}_+^3 with probability 1, namely, $(S(t), I(t), R(t)) \in \mathbb{R}_+^3$ for all $t \geq 0$ almost surely.*

Proof. Consider the diffusion process as follows

$$\begin{cases} du = \left[\frac{b(e^u + e^v + e^w)}{e^u} - \mu - \frac{\beta e^v}{e^u + e^v + e^w} + \frac{\delta e^w}{e^u} - \frac{1}{2} \sigma^2 \left(\frac{e^v}{e^u + e^v + e^w} \right)^2 \right] dt \\ \quad - \frac{\sigma e^v}{e^u + e^v + e^w} dB(t), \\ dv = \left[\frac{\beta e^v}{e^u + e^v + e^w} - (\mu + \alpha + \gamma) - \frac{\sigma^2}{2} \left(\frac{e^u}{e^u + e^v + e^w} \right)^2 \right] dt + \frac{\sigma e^u}{e^u + e^v + e^w} dB(t), \\ dw = \left[\frac{\gamma e^v}{e^u + e^v + e^w} - (\mu + \delta) \right] dt, \end{cases} \quad (10)$$

with initial value $u(0) = \log(S(0))$, $v(0) = \log(I(0))$, $w(0) = \log(R(0))$. Since the coefficients of system (10) are locally Lipschitz continuous, there is a unique local solution $(u(t), v(t), w(t))$ on $t \in [0, \tau_e)$, where τ_e is the explosion time (see Mao [14]). Therefore, by the Itô formula, it is easy to see that $S(t) = e^{u(t)}$, $I(t) = e^{v(t)}$, $R(t) = e^{w(t)}$ is the unique positive local solution of system (7) with initial value $(S(0), I(0), R(0)) \in \mathbb{R}_+^3$.

To show that this solution is global, we need to show that $\tau_e = \infty$ a.s. Let $k_0 \geq 0$ be sufficiently large so that $S(0), I(0), R(0)$ all lie within the interval $[1/k_0, k_0]$. For

each integer $k \geq k_0$, define the stopping time

$$\tau_k = \inf\{t \in [0, \tau_e) : \min\{S(t), I(t), R(t)\} \leq \frac{1}{k} \text{ or } \max\{S(t), I(t), R(t)\} \geq k\},$$

where, throughout this paper, we set $\inf \emptyset = \infty$ (and as usual \emptyset denotes the empty set). According to the definition, τ_k is increasing as $k \rightarrow \infty$. Set $\tau_\infty = \lim_{k \rightarrow \infty} \tau_k$, whence $\tau_\infty \leq \tau_e$ a.s. If we can show that $\tau_\infty = \infty$ a.s., then $\tau_e = \infty$ and $(S(t), I(t), R(t)) \in \mathbb{R}_+^3$ a.s. for all $t \geq 0$. In other words, to complete the proof all we need to show is that $\tau_\infty = \infty$ a.s. For if this statement is false, then there exist a pair of constants $T > 0$ and $\varepsilon \in (0, 1)$ such that

$$P\{\tau_\infty \leq T\} > \varepsilon.$$

Hence there is an integer $k_1 \geq k_0$ such that

$$P\{\tau_k \leq T\} \geq \varepsilon \text{ for all } k \geq k_1. \quad (11)$$

Define a C^2 -function $V : \mathbb{R}_+^3 \rightarrow \bar{\mathbb{R}}_+$, where $\bar{\mathbb{R}}_+ = \{x \in \mathbb{R} : x \geq 0\}$, by

$$V(S, I, R) = (S - 1 - \log S) + (I - 1 - \log I) + (R - 1 - \log R).$$

The nonnegativity of this function can be seen from $u - 1 - \log u \geq 0, \forall u > 0$. Let $k \geq k_0$ and $T > 0$ be arbitrary. Applying the Itô formula, we obtain

$$\begin{aligned} dV(S, I, R) &= (1 - \frac{1}{S})dS + \frac{1}{2S^2}(dS)^2 + (1 - \frac{1}{I})dI + \frac{1}{2I^2}(dI)^2 + (1 - \frac{1}{R})dR \\ &= LVdt + \frac{\sigma(I - S)}{N}dB(t), \end{aligned} \quad (12)$$

where $LV : \mathbb{R}_+^3 \rightarrow \mathbb{R}_+$ is defined by

$$\begin{aligned} LV &= (b - \mu)N + \mu + \beta \frac{I}{N} + \frac{1}{2}\sigma^2 \left(\frac{I}{N}\right)^2 + (\mu + \alpha + \gamma) + \frac{1}{2}\sigma^2 \left(\frac{S}{N}\right)^2 + (\mu + \delta) \\ &\quad - \alpha I - \frac{bN}{S} - \frac{\delta R}{S} - \frac{\beta S}{N} - \frac{\gamma I}{N} \\ &\leq (b - \mu)N + \beta + \sigma^2 + (3\mu + \gamma + \alpha + \delta). \end{aligned}$$

Noting that

$$\begin{aligned} L(e^{-bt}V) &= e^{-bt}(-bV + LV) \\ &= e^{-bt}\{-b[(S - 1 - \log S) + (I - 1 - \log I) + (R - 1 - \log R)] \\ &\quad + (b - \mu)(S + I + R) + \beta + \sigma^2 + (3\mu + \gamma + \alpha + \delta)\} \\ &= e^{-bt}[-\mu(S - b \log S) - \mu(I - b \log I) - \mu(R - b \log R) + 3b + \beta \\ &\quad + \sigma^2 + 3\mu + \gamma + \alpha + \delta]. \end{aligned} \quad (13)$$

Since $S - b - b \log(S/b) \geq 0, \forall S > 0$, we have

$$S - b \log S = (S - b - b \log \frac{S}{b}) + b(1 - \log b) \geq b(1 - \log b), \quad (14)$$

and similarly,

$$I - b \log I \geq b(1 - \log b), \quad R - b \log R \geq b(1 - \log b). \quad (15)$$

Substituting (14) and (15) into (13), we have

$$\begin{aligned} L(e^{-bt}V) &\leq e^{-bt}[-3\mu b(1 - \log b) + 3b + \beta + \sigma^2 + 3\mu + \gamma + \alpha + \delta] \\ &\leq e^{-bt}(3\mu b|1 - \log b| + 3b + \beta + \sigma^2 + 3\mu + \gamma + \alpha + \delta) \\ &:= e^{-bt}H, \end{aligned} \quad (16)$$

where $H := 3\mu b|1 - \log b| + 3b + \beta + \sigma^2 + 3\mu + \gamma + \alpha + \delta$. By (16), we have

$$LV \leq bV + H.$$

Letting $\tilde{V} = V + H/b$, we obtain

$$L\tilde{V} \leq b\tilde{V}.$$

Moreover, substituting this into (12), we obtain

$$d\tilde{V}(S, I, R) \leq b\tilde{V}dt + \frac{\sigma(I - S)}{N}dB(t).$$

By the Itô formula, for any $t \in [0, T]$ and $k \geq k_1$, we have

$$\begin{aligned} & E\tilde{V}(S(t \wedge \tau_k), I(t \wedge \tau_k), R(t \wedge \tau_k)) \\ &= \tilde{V}(S(0), I(0), R(0)) + E \int_0^{t \wedge \tau_k} L\tilde{V}(S(r), I(r), R(r))dr \\ &= \tilde{V}(S(0), I(0), R(0)) + E \int_0^{t \wedge \tau_k} b\tilde{V}(S(r), I(r), R(r))dr \\ &= \tilde{V}(S(0), I(0), R(0)) + b \int_0^t E\tilde{V}(S(r \wedge \tau_k), I(r \wedge \tau_k), R(r \wedge \tau_k))dr, \end{aligned}$$

where E is the expectation operator and $a \wedge b$ denotes the minimum of a and b . The Gronwall inequality yields that

$$E\tilde{V}(S(T \wedge \tau_k), I(T \wedge \tau_k), R(T \wedge \tau_k)) \leq \tilde{V}(S(0), I(0), R(0))e^{bT}. \quad (17)$$

Set $\Omega_k = \{\tau_k \leq t\}$ for $k \geq k_1$ and by (11), $P(\Omega_k) \geq \varepsilon$. Note that for every $\omega \in \Omega_k$, there is at least one of $S(\tau_k, \omega)$, $I(\tau_k, \omega)$ and $R(\tau_k, \omega)$ that equals either k or $1/k$, and hence $\tilde{V}(S(\tau_k, \omega), I(\tau_k, \omega), R(\tau_k, \omega))$ is no less than

$$k - 1 - \log k \text{ or } \frac{1}{k} - 1 - \log \frac{1}{k} = \frac{1}{k} - 1 + \log k.$$

Consequently,

$$\tilde{V}(S(\tau_k, \omega), I(\tau_k, \omega), R(\tau_k, \omega)) \geq [k - 1 - \log k] \wedge \left[\frac{1}{k} - 1 + \log k\right].$$

It then follows from (11) and (17) that

$$\begin{aligned} \tilde{V}(S(0), I(0), R(0))e^{bT} &\geq E[1_{\Omega_k}(\omega)\tilde{V}(S(\tau_k, \omega), I(\tau_k, \omega), R(\tau_k, \omega))] \\ &\geq \varepsilon \left[(k - 1 - \log k) \wedge \left(\frac{1}{k} - 1 + \log k \right) \right], \end{aligned}$$

where $1_{\Omega_k}(\omega)$ is the indicator function of Ω_k . Letting $k \rightarrow \infty$ leads to the contradiction

$$\infty > \tilde{V}(S(0), I(0), R(0))e^{bT} = \infty.$$

Therefore we must have $\tau_\infty = \infty$ a.s., whence the proof of Theorem 2.1 is complete. \square

Remark 1. Theorem 2.1 shows that for any initial value $(S(0), I(0), R(0)) \in \mathbb{R}_+^3$, there is a unique global solution $(S(t), I(t), R(t)) \in \mathbb{R}_+^3$ almost surely of system (7). Hence by (2), the region

$$\Gamma^* = \{(x, y, z) : x > 0, y > 0, z > 0, x + y + z = 1\}$$

is a positively invariant set of system (8), which is similar to the feasibility region Γ of the deterministic system (3). Moreover we can state the following theorem.

Theorem 2.2. *There is a unique solution $(x(t), y(t), z(t))$ of system (8) on $t \geq 0$ for any initial value $(x(0), y(0), z(0)) \in \Gamma^*$, and the solution will remain in Γ^* with probability 1, namely, $(x(t), y(t), z(t)) \in \Gamma^*$ for all $t \geq 0$ almost surely.*

From now on, we always assume that $(x(0), y(0), z(0)) \in \Gamma^*$.

3. Extinction. When studying dynamical systems for epidemics, two of the most important issues are extinction and persistence. We will discuss the extinction of the system (9) in this section but leave its persistence to the next section.

Theorem 3.1. *Let $(y(t), z(t))$ be the solution of system (9) with initial value $(y(0), z(0)) \in \mathbb{R}_+^2$ and $y(0) + z(0) < 1$.*

Case 1. If $\sigma^2 > \max\{(\beta - \alpha)^2/2(b + \gamma), \beta - \alpha\}$, then

$$\limsup_{t \rightarrow \infty} \frac{\log y(t)}{t} \leq -(b + \gamma) + \frac{(\beta - \alpha)^2}{2\sigma^2} < 0 \text{ a.s.} \quad (18)$$

Case 2. If $\beta - \alpha \leq 0$, then

$$\limsup_{t \rightarrow \infty} \frac{\log y(t)}{t} \leq -(b + \gamma) < 0 \text{ a.s.} \quad (19)$$

Case 3. If $\tilde{R}_0 < 1$ and $\sigma^2 \leq \beta - \alpha$, then

$$\limsup_{t \rightarrow \infty} \frac{\log y(t)}{t} \leq \tilde{R}_0 - 1 < 0 \text{ a.s.} \quad (20)$$

where

$$\tilde{R}_0 = R_0 - \frac{\sigma^2}{2(b + \alpha + \gamma)} = \frac{\beta - \frac{\sigma^2}{2}}{b + \alpha + \gamma}, \quad (21)$$

and R_0 is defined in (4). That is, the proportion of infectives $y(t)$ tends to zero exponentially a.s. In other words, the disease will die out with probability one. In addition, for the proportion of recovered individuals $z(t)$,

$$\lim_{t \rightarrow \infty} z(t) = 0 \text{ a.s.} \quad (22)$$

Proof. Applying the Itô formula to system (9) leads to

$$\begin{aligned} & d(\log y) \\ &= [\beta(1 - y - z) - (b + \alpha + \gamma) + \alpha y - \frac{1}{2}\sigma^2(1 - y - z)^2]dt + \sigma(1 - y - z)dB(t) \\ &= [(\beta - \alpha)(1 - y - z) - (b + \alpha + \gamma) + \alpha(1 - z) - \frac{1}{2}\sigma^2(1 - y - z)^2]dt \\ &\quad + \sigma(1 - y - z)dB(t) \\ &\leq [(\beta - \alpha)(1 - y - z) - (b + \gamma) - \frac{1}{2}\sigma^2(1 - y - z)^2]dt + \sigma(1 - y - z)dB(t) \\ &= f(x)dt + \sigma(1 - y - z)dB(t), \end{aligned}$$

where $f : (0, 1) \rightarrow \mathbb{R}$ is defined by

$$\begin{aligned} f(x) &= (\beta - \alpha)x - (b + \gamma) - \frac{1}{2}\sigma^2x^2 \\ &= -\frac{\sigma^2}{2}\left(x - \frac{\beta - \alpha}{\sigma^2}\right)^2 - (b + \gamma) + \frac{(\beta - \alpha)^2}{2\sigma^2}, \quad x = 1 - y - z \in (0, 1). \end{aligned} \quad (23)$$

Integrating this from 0 to t and dividing by t on both sides, we have

$$\frac{\log y(t)}{t} \leq \frac{\log y(0)}{t} + \frac{1}{t} \int_0^t f(x) ds + \frac{1}{t} \int_0^t \sigma(1-y-z) dB(s). \quad (24)$$

Case 1. Compute

$$\begin{aligned} f(x) &= -\frac{\sigma^2}{2} \left(x - \frac{\beta - \alpha}{\sigma^2}\right)^2 - (b + \gamma) + \frac{(\beta - \alpha)^2}{2\sigma^2} \\ &\leq -(b + \gamma) + \frac{(\beta - \alpha)^2}{2\sigma^2}, \end{aligned} \quad (25)$$

which is negative by the condition $\sigma^2 > \max\{(\beta - \alpha)^2/2(b + \gamma), \beta - \alpha\}$. It therefore follows from (24) and (25) that we have

$$\frac{\log y(t)}{t} \leq [-(b + \gamma) + \frac{(\beta - \alpha)^2}{2\sigma^2}] + \frac{M_1(t)}{t} + \frac{\log y(0)}{t},$$

where $M_1(t) := \sigma \int_0^t (1-y-z) dB(r)$. By the large number theorem for martingales (see e.g. [14]),

$$\lim_{t \rightarrow \infty} \frac{M_1(t)}{t} = 0 \text{ a.s.} \quad (26)$$

Taking the limit superior of both sides, that we obtain the desired assertion (18)

$$\limsup_{t \rightarrow \infty} \frac{\log y(t)}{t} \leq -(b + \gamma) + \frac{(\beta - \alpha)^2}{2\sigma^2} < 0 \text{ a.s.}$$

Case 2. We use the same notation as in the proof of Case 1. If $\beta - \alpha \leq 0$, by (23), it is easy to see that

$$f(x) \leq f(0) = -(b + \gamma).$$

It then follows from (24) that

$$\frac{\log y(t)}{t} \leq -(b + \gamma) + \frac{M_1(t)}{t} + \frac{\log y(0)}{t}.$$

In the same way as in the proof of Case 1, we have

$$\limsup_{t \rightarrow \infty} \frac{\log y(t)}{t} \leq -(b + \gamma) < 0 \text{ a.s.}$$

Case 3. We consider the quadratic function $f : (0, 1) \rightarrow R$ defined by (23). By (25) and the condition $\sigma^2 \leq \beta - \alpha$, it is easy to see that

$$\bar{x} = \frac{\beta - \alpha}{\sigma^2} \geq 1,$$

and then $f(x)$ takes its maximum value

$$f(\hat{x}) = f(1) = \beta - \frac{1}{2}\sigma^2 - (b + \alpha + \gamma) = (\tilde{R}_0 - 1)(b + \alpha + \gamma), \quad (27)$$

where \tilde{R}_0 is defined by (21). It follows from (24) and (27) that

$$\frac{\log y(t)}{t} \leq (\tilde{R}_0 - 1)(b + \alpha + \gamma) + \frac{M_1(t)}{t} + \frac{\log y(0)}{t}.$$

If $\tilde{R}_0 < 1$, then

$$\limsup_{t \rightarrow \infty} \frac{\log y(t)}{t} \leq (\tilde{R}_0 - 1)(b + \alpha + \gamma) < 0 \text{ a.s.}$$

For cases 1, 2 and 3, results (18), (19) and (20) respectively imply that

$$\lim_{t \rightarrow \infty} y(t) = 0 \text{ a.s.} \quad (28)$$

Next, let us prove the last assertion (22). According to (28), the last equation of the system (9) is an asymptotically differential system with limit system

$$\frac{dz}{dt} = -(b + \delta)z.$$

So we obtain

$$z(t) = z(0)e^{-(b+\delta)t}.$$

Therefore,

$$\lim_{t \rightarrow \infty} z(t) = 0 \text{ a.s.}$$

This finishes the proof of Theorem 3.1. \square

Remark 2. The condition of Case 3 states that the disease will die out if $\tilde{R}_0 < 1$ and the white noise is not large, while if the white noise is large enough such that the condition of case 1 is satisfied, then the disease will also die out. Moreover, we notice the expression of \tilde{R}_0 in (21). There is a difference compared with the threshold R_0 of system (3). In other words, the conditions for $y(t)$ to become extinct in the SDE model (9) are weaker than in the corresponding undisturbed model. The following examples illustrate this result more explicitly.

Example 1. Throughout the paper we shall assume that the unit of time is one day and the population sizes are measured in units of 1 million. Choose the parameters in system (9) as follows:

$$b = 0.2, \beta = 0.6, \alpha = 0.1, \gamma = 0.2, \delta = 0.2, \text{ and } \sigma = 0.49.$$

Noting that

$$\tilde{R}_0 = \frac{\beta}{b + \alpha + \gamma} - \frac{\sigma^2}{2(b + \alpha + \gamma)} = 0.9599 < 1, \quad \sigma^2 = 0.2401 \leq \beta - \alpha = 0.5,$$

then by Case 3 of Theorem 3.1, the solution $(y(t), z(t))$ of system (9) obeys

$$\limsup_{t \rightarrow \infty} \frac{\log y(t)}{t} \leq \tilde{R}_0 - 1 = -0.0401 < 0, \quad \lim_{t \rightarrow \infty} z(t) = 0 \text{ a.s.}$$

with any initial value $(y(0), z(0)) \in (0, 1) \times (0, 1)$. That is $y(t)$ will tend to zero exponentially with probability one and $z(t)$ will tend to zero a.s.

On the other hand, for the corresponding deterministic model,

$$R_0 = \frac{\beta}{b + \alpha + \gamma} = 1.2 > 1,$$

so the endemic proportion equilibrium (y^*, z^*) is globally asymptotically stable in Γ_0 and the disease persists. Using the Euler-Maruyama (EM) method (see [12]), we give the simulations shown in Figure 1 to support our results.

Example 2. We keep the system parameters the same as in Example 1 but let $\sigma = 0.8$. It is easy to verify that the system parameters obey the condition of Case 1 of Theorem 3.1, as

$$\sigma^2 = 0.64 > \max\left\{\frac{(\beta - \alpha)^2}{2(b + \gamma)}, \beta - \alpha\right\} = 0.5.$$

We can therefore conclude that for any initial value $(y(0), z(0)) \in (0, 1) \times (0, 1)$, the solution of (9) obeys

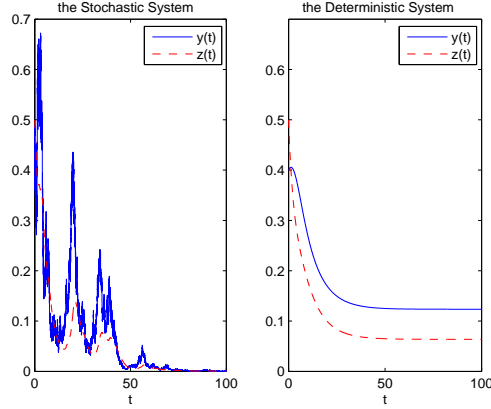


FIGURE 1. Computer simulation of the path $y(t)$, $z(t)$, shown in blue and red respectively, for the SDE SIRS model (9) (left plot) and its corresponding deterministic model (1.10) (right plot), for parameter values $\beta = 0.6$, $b = 0.2$, $\alpha = 0.1$, $\gamma = 0.2$, $\delta = 0.2$ and $\sigma = 0.49$, such that $\tilde{R}_0 < 1$, but $R_0 > 1$. The simulation uses the EM method with step size $\Delta t = 0.001$ and initial value $(y(0), z(0)) = (0.4, 0.5)$.

$$\limsup_{t \rightarrow \infty} \frac{\log y(t)}{t} \leq -(b + \gamma) + \frac{(\beta - \alpha)^2}{2\sigma^2} = -0.2047 < 0, \quad \lim_{t \rightarrow \infty} z(t) = 0 \text{ a.s.}$$

That is, $y(t)$ will tend to zero exponentially with probability one. The computer simulation shown in Figure 2 clearly supports this result, showing extinction of the disease.

4. Persistence.

Definition 4.1. System (9) is said to be persistent in the mean, if

$$\liminf_{t \rightarrow \infty} \frac{1}{t} \int_0^t y(r) dr > 0, \quad \liminf_{t \rightarrow \infty} \frac{1}{t} \int_0^t z(r) dr > 0 \text{ a.s.}$$

For convenience, we introduce some notation, defining $\langle x(t) \rangle$ as

$$\langle x(t) \rangle = \frac{1}{t} \int_0^t x(r) dr.$$

Theorem 4.2. *If*

$$\tilde{R}_0 = \frac{\beta}{b + \alpha + \gamma} - \frac{\sigma^2}{2(b + \alpha + \gamma)} > 1,$$

then for any initial value $(y(0), z(0)) \in \mathbb{R}_+^2$ and $y(0) + z(0) < 1$, the solution $(y(t), z(t))$ of system (9) has the following property:

$$\liminf_{t \rightarrow \infty} \langle y(t) \rangle \geq \tilde{y}^* \text{ and } \liminf_{t \rightarrow \infty} \langle z(t) \rangle \geq \frac{\gamma \tilde{y}^*}{(b + \delta)} \text{ a.s.,}$$

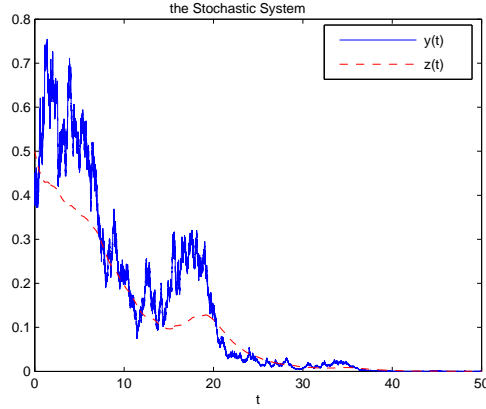


FIGURE 2. Computer simulation of the path $y(t)$, $z(t)$, shown in blue and red respectively, of the SDE SIRS model (9) for parameter values $\beta = 0.6$, $b = 0.2$, $\alpha = 0.1$, $\gamma = 0.2$, $\delta = 0.2$ and $\sigma = 0.8$, such that $\sigma^2 > \max\{(\beta - \alpha)^2/2(b + \gamma), \beta - \alpha\}$. The simulation uses the EM method with step size $\Delta t = 0.001$ and initial value $(y(0), z(0)) = (0.4, 0.5)$.

where

$$\tilde{y}^* = \frac{\{(\beta - \alpha)(b + \delta) + \alpha[\beta - (b + \alpha + \gamma)] + \beta\gamma\} - (b + \delta)\sqrt{\Delta'}}{2\alpha(\beta - \alpha)} < 1$$

and

$$\Delta' = \left\{(\beta - \alpha) + \frac{\beta\gamma + \alpha[\beta - (b + \alpha + \gamma)]}{b + \delta}\right\}^2 - \frac{4\alpha(\beta - \alpha)[\beta - (b + \alpha + \gamma) - \frac{1}{2}\sigma^2]}{b + \delta}.$$

Proof. An integration of the system (9) yields

$$\begin{cases} \frac{y(t) - y(0)}{t} = \beta\langle(1 - y(t) - z(t))y(t)\rangle - (b + \alpha + \gamma)\langle y(t)\rangle + \alpha\langle y^2(t)\rangle \\ \quad + \frac{\sigma}{t} \int_0^t (1 - y(r) - z(r))y(r)dB(r), \\ \frac{z(t) - z(0)}{t} = \gamma\langle y(t)\rangle - (b + \delta)\langle z(t)\rangle + \alpha\langle y(t)z(t)\rangle, \end{cases} \quad (29)$$

According to (29), we have

$$\begin{aligned} & \frac{\alpha(y(t) - y(0))}{t} + \frac{\beta(z(t) - z(0))}{t} - \frac{\alpha\sigma \int_0^t (1 - y(r) - z(r))y(r)dB(r)}{t} \\ & = \{\alpha[\beta - (b + \alpha + \gamma)] + \beta\gamma\}\langle y(t)\rangle - \alpha(\beta - \alpha)\langle y^2(t)\rangle - \beta(b + \delta)\langle z(t)\rangle. \end{aligned}$$

We compute that

$$\langle z(t)\rangle = \frac{\alpha[\beta - (b + \alpha + \gamma)] + \beta\gamma}{\beta(b + \delta)}\langle y(t)\rangle - \frac{\alpha(\beta - \alpha)}{\beta(b + \delta)}\langle y^2(t)\rangle - \varphi(t), \quad (30)$$

where $\varphi(t)$ is defined by

$$\varphi(t) = \frac{1}{\beta(b+\delta)} \left[\frac{\alpha(y(t) - y(0))}{t} + \frac{\beta(z(t) - z(0))}{t} - \frac{\alpha\sigma}{t} \int_0^t (1-y(r)-z(r))y(r)dB(r) \right]. \quad (31)$$

Since $0 < x(t), y(t), z(t) < 1$ and $x = 1 - y - z$, by the large number theorem for martingales (see [14]), we have that

$$\lim_{t \rightarrow \infty} \frac{y(t)}{t} = 0, \quad \lim_{t \rightarrow \infty} \frac{z(t)}{t} = 0, \quad \lim_{t \rightarrow \infty} \frac{\int_0^t x(r)y(r)dB(r)}{t} = 0 \text{ a.s.}$$

Obviously,

$$\lim_{t \rightarrow \infty} \varphi(t) = 0 \text{ a.s.} \quad (32)$$

By the Itô formula, we have

$$d(\log y) = [\beta(1-y-z) - (b+\alpha+\gamma) + \alpha y - \frac{\sigma^2}{2}(1-y-z)^2]dt + \sigma(1-y-z)dB(t).$$

Integrating this from 0 to t and dividing by t on both sides, we have

$$\begin{aligned} & \frac{\log y(t) - \log y(0)}{t} \\ &= -(\beta - \alpha)\langle y(t) \rangle + \beta - (b + \gamma + \alpha) - \beta\langle z(t) \rangle - \frac{1}{2}\sigma^2\langle (1-y(t)-z(t))^2 \rangle + \frac{M_1(t)}{t} \\ &\geq -(\beta - \alpha)\langle y(t) \rangle + [\beta - (b + \gamma + \alpha)] - \beta\langle z(t) \rangle - \frac{1}{2}\sigma^2 + \frac{M_1(t)}{t} \\ &\geq [\beta - (b + \gamma + \alpha) - \frac{1}{2}\sigma^2] - (\beta - \alpha)\langle y(t) \rangle - \frac{\alpha[\beta - (b + \alpha + \gamma)] + \beta\gamma}{b + \delta}\langle y(t) \rangle \\ &\quad + \frac{\alpha(\beta - \alpha)}{b + \delta}\langle y(t) \rangle^2 + \beta\varphi(t) + \frac{M_1(t)}{t}, \end{aligned} \quad (33)$$

where (30) and the Schwarz inequality are used. Therefore, rearranging gives

$$\frac{\alpha(\beta - \alpha)}{b + \delta}\langle y(t) \rangle^2 - \lambda\langle y(t) \rangle + [\beta - (b + \gamma + \alpha) - \frac{1}{2}\sigma^2] \leq \phi(t),$$

where $\lambda = \{(\beta - \alpha) + \frac{\alpha[\beta - (b + \alpha + \gamma)] + \beta\gamma}{b + \delta}\}$ and $\phi(t) = \frac{\log y(t) - \log y(0)}{t} - \frac{M_1(t)}{t} - \beta\varphi(t)$. Noting that $-\infty < \log y(t) < 0$ (as $0 < y(t) < 1$) and using (26) and (32), we see that

$$\lim_{t \rightarrow \infty} \phi(t) = 0 \text{ a.s.}$$

Then for arbitrary $0 < \varepsilon < 1$, there exists $T = T(\omega) > 0$ and a set Ω_ε , such that $P(\Omega_\varepsilon) \geq 1 - \varepsilon$. For all $t \geq T(\omega)$, $\omega \in \Omega_\varepsilon$, then $\phi(t) \leq \varepsilon$, so we have

$$\frac{\alpha(\beta - \alpha)}{b + \delta}\langle y(t) \rangle^2 - \lambda\langle y(t) \rangle + [\beta - (b + \gamma + \alpha) - \frac{1}{2}\sigma^2 - \varepsilon] \leq 0, \quad (34)$$

which has the form $c_1\langle y(t) \rangle^2 + c_2\langle y(t) \rangle + c_3 \leq 0$. If $\tilde{R}_0 > 1$, then

$$\begin{aligned}
\Delta &= c_2^2 - 4c_1 * c_3 \\
&= \lambda^2 - \frac{4\alpha(\beta - \alpha)[\beta - (b + \gamma + \alpha) - \frac{1}{2}\sigma^2 - \varepsilon]}{b + \delta} \\
&= \left\{ (\beta - \alpha) + \frac{\beta\gamma - \alpha[\beta - (b + \alpha + \gamma)]}{b + \delta} \right\}^2 + \frac{4\alpha\beta\gamma[\beta - (b + \alpha + \gamma)]}{(b + \delta)^2} \\
&\quad + \frac{4\alpha(\beta - \alpha)[\frac{1}{2}\sigma^2 + \varepsilon]}{b + \delta} \\
&> 0.
\end{aligned} \tag{35}$$

Let $\varepsilon < \beta - (b + \gamma + \alpha) - \sigma^2/2$ and

$$y_1^\varepsilon = \frac{\{(\beta - \alpha)(b + \delta) + \alpha[\beta - (b + \alpha + \gamma)] + \beta\gamma\} - (b + \delta)\sqrt{\Delta}}{2\alpha(\beta - \alpha)}$$

and

$$y_2^\varepsilon = \frac{\{(\beta - \alpha)(b + \delta) + \alpha[\beta - (b + \alpha + \gamma)] + \beta\gamma\} + (b + \delta)\sqrt{\Delta}}{2\alpha(\beta - \alpha)}.$$

Hence, by (34), we have

$$y_1^\varepsilon \leq \langle y(t) \rangle \leq y_2^\varepsilon, \quad t \geq T(\omega), \quad \omega \in \Omega_\varepsilon. \tag{36}$$

Next, we shall prove the assertion

$$0 < y_1^\varepsilon < \frac{\beta - (b + \alpha + \gamma)}{\beta - \alpha} < 1.$$

According to (35), we can easily obtain that

$$(\beta - \alpha) + \frac{\beta\gamma - \alpha[\beta - (b + \alpha + \gamma)]}{b + \delta} < \sqrt{\Delta}.$$

Some computations yield the expression

$$(\beta - \alpha)(b + \delta) + \beta\gamma + \alpha[\beta - (b + \alpha + \gamma)] - (b + \delta)\sqrt{\Delta} < 2\alpha[\beta - (b + \alpha + \gamma)],$$

thus

$$y_1^\varepsilon = \frac{(b + \delta)(\lambda - \sqrt{\Delta'})}{2\alpha(\beta - \alpha)} < \frac{\beta - (b + \alpha + \gamma)}{\beta - \alpha} < 1.$$

So the assertion is true. Therefore, by (36) we obtain

$$\liminf_{t \rightarrow \infty} \langle y(t) \rangle \geq y_1^\varepsilon, \quad a.s.$$

Letting $\varepsilon \rightarrow 0$, we get

$$\liminf_{t \rightarrow \infty} \langle y(t) \rangle \geq \frac{(b + \delta)(\lambda - \sqrt{\Delta'})}{2\alpha(\beta - \alpha)} := \tilde{y}^* \quad a.s.,$$

where

$$\Delta' := \lambda^2 - \frac{4\alpha(\beta - \alpha)[\beta - (b + \gamma + \alpha) - \frac{1}{2}\sigma^2]}{(b + \delta)},$$

and

$$0 < \tilde{y}^* \leq \frac{\beta - (b + \alpha + \gamma)}{\beta - \alpha} < 1.$$

Finally, according to the last equality of (29), we get

$$\begin{aligned} \frac{z(t) - z(0)}{t} &= \gamma \langle y(t) \rangle - (b + \delta) \langle z(t) \rangle + \alpha \langle y(t)z(t) \rangle \\ &\geq \gamma \langle y(t) \rangle - (b + \delta) \langle z(t) \rangle, \end{aligned}$$

thus

$$\langle z(t) \rangle \geq \frac{\gamma}{(b + \delta)} \langle y(t) \rangle - \frac{z(t) - z(0)}{(b + \delta)t}.$$

So we have

$$\liminf_{t \rightarrow \infty} \langle z(t) \rangle \geq \frac{\gamma}{(b + \delta)} \tilde{y}^* \text{ a.s.}$$

This finishes the proof of Theorem 4.2. \square

Remark 3. From Theorems 3.1 and 4.2, we can see that when the noise is sufficiently small that $\sigma^2 \leq \beta - \alpha$, if the value of \tilde{R}_0 is above 1 or below 1 then the disease will persist or die out respectively. Therefore, we consider \tilde{R}_0 as the threshold of system (9).

Example 3. Assume that the parameters of system (9) are given by

$$b = 0.2, \beta = 0.6, \alpha = 0.1, \gamma = 0.2, \delta = 0.2, \text{ and } \sigma = 0.1.$$

That is, we keep all the parameters the same as in Example 1, except that σ is reduced to 0.1 from 0.49. Note that

$$\tilde{R}_0 = \frac{\beta}{b + \alpha + \gamma} - \frac{\sigma^2}{2(b + \alpha + \gamma)} = 1.2 - 0.01 = 1.19 > 1,$$

so by Theorem 4.2, for any initial value $(y(0), z(0)) \in (0, 1) \times (0, 1)$, we conclude that the solution $(y(t), z(t))$ of system (9) obeys

$$\liminf_{t \rightarrow \infty} \langle y(t) \rangle \geq \tilde{y}^* = 0.1235 \text{ and } \liminf_{t \rightarrow \infty} \langle z(t) \rangle \geq \frac{\gamma}{(b + \delta)} \tilde{y}^* = 0.0637 \text{ a.s.}$$

That is to say, the disease persists. The simulations shown in Figure 3 illustrate this.

5. Asymptotic behavior around the endemic proportion equilibrium. In the deterministic model, the endemic proportion equilibrium $P^*(y^*, z^*)$ is globally asymptotically stable. In this section, we show the effect of stochastic fluctuations of the environment on the endemic proportion equilibrium of the corresponding deterministic system.

Theorem 5.1. *If $R_0 > 1$ and $\alpha^2 < 4\gamma^2 y^*(\beta - \alpha)/\beta z^*$, then for any initial value $(y(0), z(0)) \in \mathbb{R}_+^2$ and $y(0) + z(0) < 1$, the solution $(y(t), z(t))$ of system (9) has the following property:*

$$\limsup_{t \rightarrow \infty} \frac{1}{t} \int_0^t \left(\beta - \alpha - \frac{\beta \alpha \rho}{2\gamma} \right) (y(r) - y^*)^2 + \frac{\beta}{\gamma} \left(\frac{\gamma y^*}{z^*} - \frac{\alpha}{2\rho} \right) (z(r) - z^*)^2 dr \leq \frac{\sigma^2 y^*}{2} \text{ a.s.} \quad (37)$$

where ρ is a positive constant satisfying the inequality $\alpha z^*/2\gamma y^* < \rho < 2\gamma(\beta - \alpha)/\alpha\beta$.

Proof. Since $P^*(y^*, z^*)$ is the endemic equilibrium of the corresponding deterministic model of system (9), we have that

$$b + \gamma + \alpha = \beta(1 - y^* - z^*) + \alpha y^*, \quad \gamma y^* + \alpha y^* z^* = (b + \delta) z^*.$$

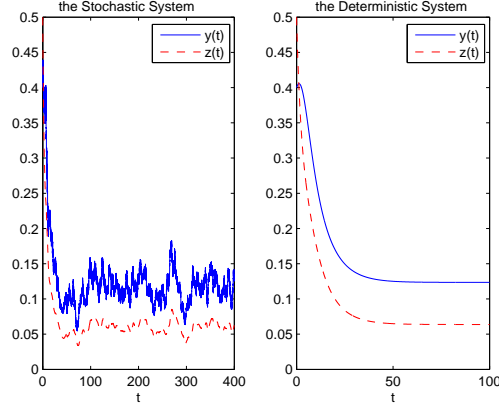


FIGURE 3. Computer simulation of the path $y(t)$, $z(t)$, shown in blue and red respectively, for the SDE SIRS model (9) (left plot) and its corresponding deterministic model (right plot), for parameter values $\beta = 0.6$, $b = 0.2$, $\alpha = 0.1$, $\gamma = 0.2$, $\delta = 0.2$ and $\sigma = 0.1$, such that $\tilde{R}_0 > 1$. The simulation uses the EM method with step size $\Delta t = 0.001$ and initial value $(y(0), z(0)) = (0.4, 0.5)$.

Define a C^2 -function $V : (0, 1) \times (0, 1) \rightarrow \mathbb{R}_+$ by

$$V(y, z) = y - y^* - y^* \log \frac{y}{y^*} + \frac{\beta}{2\gamma} (z - z^*)^2 := V_1 + \frac{\beta}{\gamma} V_2,$$

where $V_1 = y - y^* - y^* \log(y/y^*)$ and $V_2 = (z - z^*)^2/2$. The nonnegativity of this function can be observed from $u - 1 - \log u \geq 0$ on $u > 0$. Let L be the generating operator of system (9). Then we get

$$\begin{aligned} LV_1 &= [\beta(1 - y - z) - (b + \gamma + \alpha) + \alpha y](y - y^*) + \frac{1}{2} \sigma^2 y^* (1 - y - z)^2 \\ &= [-\beta(y - y^*) - \beta(z - z^*) - \alpha(y - y^*)](y - y^*) + \frac{1}{2} \sigma^2 y^* (1 - y - z)^2 \\ &\leq -(\beta - \alpha)(y - y^*)^2 - \beta(y - y^*)(z - z^*) + \frac{1}{2} \sigma^2 y^* \end{aligned}$$

and

$$\begin{aligned} LV_2 &= (z - z^*)[\gamma y - (b + \delta)z + \alpha y z] \\ &= (z - z^*)[\gamma(y - y^*) - (b + \delta)(z - z^*) + \alpha z(y - y^*) + \alpha y^*(z - z^*)] \\ &= (z - z^*)[(\gamma + \alpha z)(y - y^*) - \frac{\gamma y^*}{z^*}(z - z^*)] \\ &= (\gamma + \alpha z)(y - y^*)(z - z^*) - \frac{\gamma y^*}{z^*}(z - z^*)^2. \end{aligned}$$

Then

$$\begin{aligned}
LV &= LV_1 + \frac{\beta}{\gamma} LV_2 \\
&\leq -(\beta - \alpha)(y - y^*)^2 - \beta(y - y^*)(z - z^*) + \frac{\beta}{\gamma}(\gamma + \alpha z)(y - y^*)(z - z^*) \\
&\quad - \frac{\beta y^*}{z^*}(z - z^*)^2 + \frac{1}{2}\sigma^2 y^* \\
&= -(\beta - \alpha)(y - y^*)^2 + \frac{\alpha\beta z}{\gamma}(y - y^*)(z - z^*) - \frac{\beta y^*}{z^*}(z - z^*)^2 + \frac{1}{2}\sigma^2 y^* \\
&\leq -(\beta - \alpha)(y - y^*)^2 + \frac{\alpha\beta}{\gamma} \|y - y^*\| |z - z^*| - \frac{\beta y^*}{z^*}(z - z^*)^2 + \frac{1}{2}\sigma^2 y^* \\
&\leq -(\beta - \alpha - \frac{\beta\alpha\rho}{2\gamma})(y - y^*)^2 - \frac{\beta}{\gamma}(\frac{\gamma y^*}{z^*} - \frac{\alpha}{2\rho})(z - z^*)^2 + \frac{1}{2}\sigma^2 y^* \\
&:= F(t),
\end{aligned}$$

where ρ is a positive constant as specified in Theorem 5.1, and Young's inequality is used. If

$$\alpha^2 < \frac{4\gamma^2 y^*}{\beta z^*}(\beta - \alpha),$$

then

$$\beta - \alpha - \frac{\beta\alpha\rho}{2\gamma} > 0 \text{ and } \frac{\gamma y^*}{z^*} - \frac{\alpha}{2\rho} > 0.$$

Therefore,

$$dV \leq F(t)dt + \sigma(1 - y - z)(y - y^*)dB(t).$$

Integrating both sides of this from 0 to t yields

$$V(t) - V(0) \leq \int_0^t F(s)ds + \int_0^t \sigma(1 - y - z)(y - y^*)dB(s). \quad (38)$$

Let $M_2(t) := \int_0^t \sigma(1 - y - z)(y - y^*)dB(s)$. By the large number theorem for martingales (see e.g. [14]), we obtain

$$\lim_{t \rightarrow \infty} \frac{M_2(t)}{t} = 0 \text{ a.s.},$$

which together with (38) implies

$$\limsup_{t \rightarrow \infty} \frac{\int_0^t F(s)ds}{t} \geq 0 \text{ a.s.}$$

Consequently,

$$\limsup_{t \rightarrow \infty} \frac{1}{t} \int_0^t (\beta - \alpha - \frac{\beta\alpha\rho}{2\gamma})(y(r) - y^*)^2 + \frac{\beta}{\gamma}(\frac{\gamma y^*}{z^*} - \frac{\alpha}{2\rho})(z(r) - z^*)^2 dr \leq \frac{\sigma^2 y^*}{2} \text{ a.s.}$$

Thus, Theorem 5.1 is proved. \square

Remark 4. Theorem 5.1 shows that under some conditions, the distance between the solution $Y(t) = (y(t), z(t))$ of the stochastic system (9) and the endemic proportion equilibrium $P^* = (y^*, z^*)$ has the form

$$\limsup_{t \rightarrow \infty} \frac{1}{t} \int_0^t \|Y(s) - P^*\|^2 ds \leq C \|\sigma\|^2 \text{ a.s.},$$

where C is a positive constant. Therefore, although the solution of system (9) does not have stability like the deterministic system, there is approximate stability, provided that $\|\sigma\|^2$ is sufficiently small.

Example 4. Assume that the parameters of system (9) are given by

$$b = 0.2, \beta = 0.6, \alpha = 0.1, \gamma = 0.2, \delta = 0.2, \text{ and } \sigma = 0.05 \text{ or } \sigma = 0.01.$$

That is, we keep all the parameters the same as in Example 1, except for σ which is now much smaller. Note that

$$R_0 = \frac{\beta}{b + \alpha + \gamma} = 1.2 > 1.$$

Then by Theorem 5.1, for any initial value $(y(0), z(0)) \in (0, 1) \times (0, 1)$, we conclude that the difference between the perturbed solution $(y(t), z(t))$ of system (9) and $P^*(y^*, z^*)$ is related only to the level of the white noise, under the conditions $R_0 > 1$ and $\alpha^2 < 4\gamma^2 y^*(\beta - \alpha)/\beta z^*$. Using the EM method mentioned in [12], we show simulations to support our result. As expected, the solution oscillates around the endemic equilibrium P^* for a long time (see Figure 4). The parameters used for the first two plots in Figure 4 are all the same but with different intensities of the white noise σ , i.e. in the first one $\sigma = 0.05$ and in the second one $\sigma = 0.01$. We observe clearly that as the white noise gets weaker, the fluctuation around P^* gets smaller, which supports the result of Theorem 5.1.

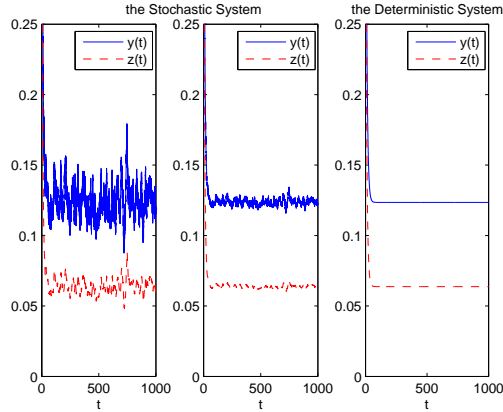


FIGURE 4. Computer simulation of the path $y(t), z(t)$, shown in blue and red respectively, for the SDE SIRS model (9) and its corresponding deterministic model (right plot) for parameter values $\beta = 0.6, b = 0.2, \alpha = 0.1, \gamma = 0.2, \delta = 0.2$ and differing values of $\sigma = 0.05$ (left plot) and $\sigma = 0.01$ (middle plot). The simulation uses the EM method with step size $\Delta t = 0.02$ and initial value $(y(0), z(0)) = (0.4, 0.5)$.

6. Conclusions. In this paper, we have looked at an SDE version of the classical SIRS epidemic model in a population with varying size, with noise introduced into the disease transmission term. We showed that the SDE has a unique positive global solution, and we established conditions for the extinction and persistence of the disease. A key parameter is the threshold \tilde{R}_0 , which is less than the corresponding deterministic version R_0 . Theorem 3.1 shows that if $\tilde{R}_0 < 1$, under mild extra conditions the disease will die out. Theorem 4.2 shows that if $\tilde{R}_0 > 1$, then the disease will persist. We also showed that if the conditions of Theorem 5.1 are

satisfied, then the SDE solution oscillates around the endemic equilibrium of the deterministic system, and the intensity of fluctuation is proportional to the white noise intensity. Throughout the paper we have illustrated our theoretical results with computer simulations.

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REFERENCES

- [1] R. M. Anderson and R. M. May, Population biology of infectious diseases I, *Nature*, **280** (1979), 361–367.
- [2] R. M. Anderson and R. M. May, Population biology of infectious diseases II, *Nature*, **280** (1979), 455–461.
- [3] R. M. Anderson and R. M. May, The population dynamics of microparasites and their invertebrate hosts, *Philos. Trans. R. Soc. Lond. B*, **291** (1981), 451–524.
- [4] M. Bandyopadhyay and J. Chattopadhyay, Ratio-dependent predator-prey model: effect of environmental fluctuation and stability, *Nonlinearity*, **18** (2005), 913–936.
- [5] S. Busenberg, K. L. Cooke and M. A. Pozio, Analysis of a model of a vertically transmitted disease, *J. Math. Biol.*, **17** (1983), 305–329.
- [6] S. Busenberg and P. van den Driessche, Analysis of a disease transmission model in a population with varying size, *J. Math. Biol.*, **28** (1990), 257–270.
- [7] S. Busenberg, K. L. Cooke and H. Thieme, Demographic change and persistence of HIV/AIDS in a heterogeneous population, *Siam J. Appl. Math.*, **51** (1991), 1030–1052.
- [8] M. Carletti, K. Burrage and P. M. Burrage, Numerical simulation of stochastic ordinary differential equations in biomathematical modelling, *Math. Comput. Simulation*, **64** (2004), 271–277.
- [9] N. Dalal, D. Greenhalgh and X. Mao, A stochastic model of AIDS and condom use, *J. Math. Anal. Appl.*, **325** (2007), 36–53.
- [10] M. Fan, M. Y. Li and K. Wang, Global stability of an SEIS epidemic model with recruitment and a varying total population size, *Math. Biosciences*, **170** (2001), 199–208.
- [11] A. Gray, D. Greenhalgh, L. Hu, X. Mao and J. Pan, A stochastic differential equation SIS epidemic model, *Siam J. Appl. Math.*, **71** (2011), 876–902.
- [12] D. J. Higham, An algorithmic introduction to numerical simulation of stochastic differential equations, *Siam Review*, **43** (2001), 525–546.
- [13] C. Y. Ji, D. Q. Jiang and N. Z. Shi, The behavior of an SIR epidemic model with stochastic perturbation, *Stochastic Anal. Appl.*, **30** (2012), 755–773.
- [14] X. Mao, *Stochastic Differential Equations and Applications*, 2nd edition, Horwood, Chichester, UK, 2008.
- [15] M. Martcheva and C. Castillo-Chavez, Diseases with chronic stage in a population with varying size, *Math. Biosciences*, **182** (2003), 1–25.
- [16] R. M. May, R. M. Anderson and A. R. Mclean, Possible demographic consequences of HIV/AIDS epidemics. I. assuming HIV infection always leads to AIDS, *Math. Biosciences*, **90** (1988), 475–505.
- [17] C. J. Sun and Y. H. Hsieh, Global analysis of an SEIR model with varying population size and vaccination, *Applied Mathematical Modelling*, **34** (2010), 2685–2697.
- [18] E. Tornatore, S. M. Buccellato and P. Vetro, Stability of a stochastic SIR system, *Physica A*, **354** (2005), 111–126.
- [19] C. Vargas-De-Leon, On the global stability of SIS, SIR and SIRS epidemic models with standard incidence, *Chaos, Solitons & Fractals*, **44** (2011), 1106–1110.
- [20] Q. Yang and X. Mao, Extinction and recurrence of multi-group SEIR epidemic models with stochastic perturbations, *Nonlinear Anal. RWA*, **14** (2013), 1434–1456.
- [21] Yanan Zhao and Daqing Jiang, The threshold of a stochastic SIRS epidemic model with saturated incidence, *Applied Mathematical Letter*, **34** (2014), 90–93.

- [22] Y. Zhao, D. Jiang and D. O'Regan, The extinction and persistence of the stochastic SIS epidemic model with vaccination, *Physica A*, **392** (2013), 4916–4927.

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