

RESEARCH

Open Access

Global dynamics for an SIR patchy model with susceptibles dispersal

Luju Liu*, Weiyun Cai and Yusen Wu

*Correspondence:
lujuliu@gmail.com
School of Mathematics and
Statistics, Henan University of
Science and Technology, Luoyang,
471003, P.R. China

Abstract

An *SIR* epidemiological model with susceptibles dispersal between two patches is addressed and discussed. The basic reproduction numbers R_{01} and R_{02} are defined as the threshold parameters. It shows that if both R_{01} and R_{02} are below unity, the disease-free equilibrium is shown to be globally asymptotically stable by using the comparison principle of the cooperative systems. If R_{01} is above unity and R_{02} is below unity, the disease persists in the first patch provided $S_2^{1*} < S_2^{2*}$. If R_{02} is above unity, R_{01} is below unity, and $S_1^{2*} < S_1^{1*}$, the disease persists in the second patch. And if R_{01} and R_{02} are above unity, and further $S_2^{1*} > S_2^{2*}$ and $S_1^{2*} > S_1^{1*}$ are satisfied, the unique endemic equilibrium is globally asymptotically stable by constructing the Lyapunov function. Furthermore, it follows that the susceptibles dispersal in the population does not alter the qualitative behavior of the epidemiological model.

1 Introduction

The development of economic globalization and the progression of science and technology yield more and more frequent contact and communication between people in different countries and regions, which further directly accelerates the development of global economy and fosters the prosperity and flourishing of a society. However, the bad things may occur simultaneously, such as, the spread of 2003 SARS and 2009 H1N1 influenza almost throughout the world. SARS involved 30 countries and regions, caused more than 8,000 patients, and 774 deaths [19, 20]. The H1N1 influenza virus quickly spread worldwide due to airplane travel. As of May 6, 2009, the virus had invaded in 23 countries including Mexico and the United States, and a total of 1,882 people were confirmed to be infected by it [3]. It then follows that the studies on the influence of infectious diseases transmission on the global population that formulates patchy models are more and more significant and practical.

A great number of mathematical patchy models have been proposed and analyzed to illustrate the influence of the transmission of infectious diseases on the local population among many countries and regions [1, 2, 7, 12, 18]. But for many mathematical models of infectious diseases in a patchy environment, the global stability of the endemic equilibrium is still an open problem. Motivated by this, in the present paper, a class of simple *SIR* models with susceptibles dispersal in a patchy environment is to be formulated and investigated the stability of the endemic equilibrium by constructing the Lyapunov function (also see [5, 6, 9–11, 13, 14]).

The rest of this paper is organized as follows. In Sect. 2, the *SIR* model with susceptibles dispersal between two disjoint patches is formulated, and the existence, uniqueness,

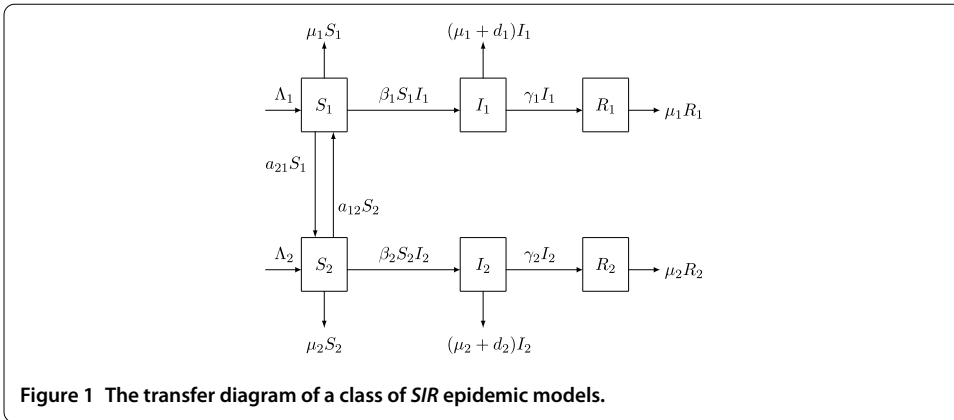


Figure 1 The transfer diagram of a class of SIR epidemic models.

and boundedness of the solutions are analyzed. The existence of equilibria and the basic reproduction numbers are derived in Sect. 3. In Sect. 4, the long-term behavior of the SIR model is studied. The brief conclusions and discussions are given in Sect. 5.

2 Model formulation

In this section, a class of SIR epidemic models for infectious diseases between two patches is developed, in which only susceptible people may disperse between two disjoint patches. All the persons are classified into three compartments: susceptible (S), infectious (I), and removed (R) in each patch, respectively. It is assumed that the mass action incidence is used and there is no birth or death during travel. Based on the transfer diagram of Figure 1, the SIR epidemic model to understand the impact of susceptibles dispersal on the whole population is described by the following system of ordinary differential equations:

$$\begin{aligned}
 \frac{dS_1}{dt} &= \Lambda_1 - \beta_1 S_1 I_1 - \mu_1 S_1 + a_{12} S_2 - a_{21} S_1, \\
 \frac{dI_1}{dt} &= \beta_1 S_1 I_1 - (\mu_1 + d_1 + \gamma_1) I_1, \\
 \frac{dS_2}{dt} &= \Lambda_2 - \beta_2 S_2 I_2 - \mu_2 S_2 + a_{21} S_1 - a_{12} S_2, \\
 \frac{dI_2}{dt} &= \beta_2 S_2 I_2 - (\mu_2 + d_2 + \gamma_2) I_2.
 \end{aligned} \tag{1}$$

Since R_1 and R_2 do not involve in other equations but themselves in system (1), they are not directly taken into account in system (1).

Λ_i ($i = 1, 2$) is the recruitment constant rate of the population in the i th patch. β_i ($i = 1, 2$) represents the transmission rate in the i th patch. μ_i ($i = 1, 2$) represents the natural death rate in the i th patch. d_i ($i = 1, 2$) is the induced-death rate in the i th patch. γ_i ($i = 1, 2$) is the recovery rate of the infectious persons in the i th patch. a_{12} represents the dispersal rate of susceptible individuals from the second patch to the first patch. a_{21} represents the dispersal rate of susceptible individuals from the first patch to the second patch. All the parameters considered in the present paper are nonnegative. $N_i(t)$ ($i = 1, 2$) denotes the number of the total population in the i th patch at time t . Therefore, $N_i = S_i + I_i + R_i$ ($i = 1, 2$).

By applying the Theorem 5.2.1 of [15], it then follows that for any $(S_{10}, I_{10}, S_{20}, I_{20}) \in \mathbb{R}_+^4$, system (1) exists a unique local nonnegative solution $(S_1(t), I_1(t), S_2(t), I_2(t))$ through the initial value $(S_1(0), I_1(0), S_2(0), I_2(0)) = (S_{10}, I_{10}, S_{20}, I_{20})$.

The expressions of N_i and Eq. (1) give rise to the following formula:

$$\begin{aligned} \frac{dN_1}{dt} + \frac{dN_2}{dt} &= \Lambda_1 + \Lambda_2 - \mu_1 N_1 - \mu_2 N_2 - d_1 I_1 - d_2 I_2 \\ &\leq \Lambda_1 + \Lambda_2 - \min\{\mu_1, \mu_2\}(N_1 + N_2). \end{aligned} \tag{2}$$

System (2) implies $\limsup_{t \rightarrow \infty} (N_1 + N_2) \leq (\Lambda_1 + \Lambda_2) / \min\{\mu_1, \mu_2\}$. Therefore, $N_1 + N_2$ is ultimately bounded and all the solutions of system (1) globally exists on the interval $[0, \infty)$. The aforementioned discussions can be summarized into the following results.

Theorem 2.1 *System (1) exists a unique and bounded solution throughout the initial value $(S_{10}, I_{10}, S_{20}, I_{20}) \in \mathbb{R}_+^4$. Further, the compact set*

$$\Omega := \left\{ (S_1, I_1, S_2, I_2) \in \mathbb{R}_+^4 : S_1 + I_1 + S_2 + I_2 \leq \frac{\Lambda_1 + \Lambda_2}{\min\{\mu_1, \mu_2\}} \right\}$$

is a positively invariant set and attracts all positive orbits in \mathbb{R}_+^4 .

Note that the long-time behaviors of the solutions of system (1) are investigated in region Ω instead of the space \mathbb{R}_+^4 .

3 Equilibria and the basic reproduction numbers

In this section, the existence of equilibria and the basic reproduction numbers are studied. By the direct calculation, system (1) always exhibits one disease-free equilibrium $P_0 = (S_1^0, 0, S_2^0, 0)$ for all parameters, where

$$S_1^0 = \frac{\Lambda_1 a_{12} + \Lambda_2 a_{12} + \Lambda_1 \mu_2}{\mu_1 \mu_2 + \mu_2 a_{21} + \mu_1 a_{12}}, \quad S_2^0 = \frac{\Lambda_2 a_{21} + \Lambda_1 a_{21} + \Lambda_2 \mu_1}{\mu_1 \mu_2 + \mu_2 a_{21} + \mu_1 a_{12}}.$$

Applying the next generation matrix approach developed in [4] gives rise to the following formulas:

$$\mathcal{F} = \begin{bmatrix} \beta_1 S_1^0 & 0 \\ 0 & \beta_2 S_2^0 \end{bmatrix} =: \begin{bmatrix} F_1 & 0 \\ 0 & F_2 \end{bmatrix},$$

and

$$\mathcal{V} = \begin{bmatrix} \mu_1 + d_1 + \gamma_1 & 0 \\ 0 & \mu_2 + d_2 + \gamma_2 \end{bmatrix} =: \begin{bmatrix} V_1 & 0 \\ 0 & V_2 \end{bmatrix}.$$

Therefore, the basic reproduction number is defined as

$$\begin{aligned} R_0 &= \rho(\mathcal{F}\mathcal{V}^{-1}) = \max\{\rho(F_1 V_1^{-1}), \rho(F_2 V_2^{-1})\} \\ &= \max\{R_{01}, R_{02}\} = \max\left\{ \frac{\beta_1 S_1^0}{\mu_1 + d_1 + \gamma_1}, \frac{\beta_2 S_2^0}{\mu_2 + d_2 + \gamma_2} \right\}, \end{aligned}$$

where $\rho(\mathbf{M})$ denotes for the spectral radius of the matrix \mathbf{M} , R_{01} and R_{02} correspond to the basic reproduction numbers of the first and the second patch when there is no dispersal between two patches, respectively. The proof process of [4], Theorem 2 implies the following statements.

Lemma 3.1 *There hold*

- (1) Let $M_1 = F_1 - V_1$ and $s(M_1)$ be the maximum real part of all the eigenvalues of the matrix M_1 . Then $s(M_1) < 0$ if and only if $R_{01} < 1$, and $s(M_1) > 0$ if and only if $R_{01} > 1$;
- (2) Let $M_2 = F_2 - V_2$ and $s(M_2)$ be the maximum real part of all the eigenvalues of the matrix M_2 . Then $s(M_2) < 0$ if and only if $R_{02} < 1$, and $s(M_2) > 0$ if and only if $R_{02} > 1$.

Furthermore, if $R_{01} > 1$ and $R_{02} < 1$, there exists a nontrivial boundary equilibrium $P_{1^*} = (S_1^{1^*}, I_1^{1^*}, S_2^*, 0)$, where

$$S_1^{1^*} = \frac{\mu_1 + d_1 + \gamma_1}{\beta_1}, \quad S_2^* = \frac{\Lambda_2 + a_{21}S_1^{1^*}}{\mu_2 + a_{12}},$$

$$I_1^{1^*} = \frac{(\mu_1\mu_2 + \mu_1a_{12} + \mu_2a_{21})S_1^{1^*}(R_{01} - 1)}{(\mu_2 + a_{12})(\mu_1 + d_1 + \gamma_1)}.$$

If $R_{02} > 1$ and $R_{01} < 1$, there exists another nontrivial boundary equilibrium $P_{2^*} = (S_1^{2^*}, 0, S_2^{2^*}, I_2^{2^*})$, where

$$S_1^{2^*} = \frac{\Lambda_1 + a_{12}S_2^{2^*}}{\mu_1 + a_{21}}, \quad S_2^{2^*} = \frac{\mu_2 + d_2 + \gamma_2}{\beta_2},$$

$$I_2^{2^*} = \frac{(\mu_1\mu_2 + \mu_1a_{12} + \mu_2a_{21})S_2^{2^*}(R_{02} - 1)}{(\mu_1 + a_{21})(\mu_2 + d_2 + \gamma_2)}.$$

If $R_{01} > 1, R_{02} > 1, S_1^{2^*} > S_1^{1^*}$, and $S_2^{2^*} > S_2^*$, system (1) admits exactly one endemic equilibrium $P^{**} = (S_1^{**}, I_1^{**}, S_2^{**}, I_2^{**})$, where

$$S_1^{**} = S_1^{1^*}, \quad I_1^{**} = \frac{S_1^{2^*} - S_1^{**}}{(\mu_1 + a_{21})\beta_1 S_1^{**}}, \quad S_2^{**} = S_2^*, \quad I_2^{**} = \frac{S_2^{1^*} - S_2^{**}}{(\mu_2 + a_{12})\beta_2 S_2^{**}}.$$

4 Threshold dynamics

In this section, the stability of equilibria is to be formulated. First of all, the global stability of the disease-free equilibrium P_0 is to be discussed. There holds the following result.

Theorem 4.1 *If the basic reproduction number R_0 is less than one, the disease-free equilibrium P_0 is globally asymptotically stable; while if the basic reproduction number R_0 is greater than one, the disease-free equilibrium P_0 is unstable.*

Proof If $R_0 < 1$, [4], Theorem 2, yields that P_0 is locally asymptotically stable. Thus, it is sufficient to prove the global attractivity of P_0 when $R_0 < 1$. The first and third equations of system (1) implies

$$\frac{dS_1}{dt} \leq \Lambda_1 - (\mu_1 + a_{21})S_1 + a_{12}S_2,$$

$$\frac{dS_2}{dt} \leq \Lambda_2 - (\mu_2 + a_{12})S_2 + a_{21}S_1.$$

It is easy to see the following linear system:

$$\begin{aligned} \frac{d\hat{S}_1}{dt} &= \Lambda_1 - (\mu_1 + a_{21})\hat{S}_1 + a_{12}\hat{S}_2, \\ \frac{d\hat{S}_2}{dt} &= \Lambda_2 - (\mu_2 + a_{12})\hat{S}_2 + a_{21}\hat{S}_1, \end{aligned} \tag{3}$$

has a positive equilibrium $\hat{S}_0 = (S_1^0, S_2^0)$ and \hat{S}_0 is globally asymptotically stable for system (3) in \mathbb{R}_+^2 . Consequently, the comparison principle of cooperative systems [16], Theorem B.1, yields that for any $\varepsilon > 0$, $S_i(t) < S_i^0 + \varepsilon$ ($i = 1, 2$) is satisfied, for sufficiently large t . Thus, if t is sufficiently large, the second and fourth equations of system (1) admit

$$\begin{aligned} \frac{dI_1}{dt} &< \beta_1 I_1 (S_1^0 + \varepsilon) - (\mu_1 + d_1 + \gamma_1) I_1, \\ \frac{dI_2}{dt} &< \beta_2 I_2 (S_2^0 + \varepsilon) - (\mu_2 + d_2 + \gamma_2) I_2. \end{aligned}$$

Thus, it suffices to prove the following system:

$$\begin{aligned} \frac{d\tilde{I}_1}{dt} &= (\beta_1 S_1^0 - (\mu_1 + d_1 + \gamma_1))\tilde{I}_1 + \varepsilon\beta_1\tilde{I}_1, \\ \frac{d\tilde{I}_2}{dt} &= (\beta_2 S_2^0 - (\mu_2 + d_2 + \gamma_2))\tilde{I}_2 + \varepsilon\beta_2\tilde{I}_2, \end{aligned} \tag{4}$$

tends to the zero solution as t goes to infinity. Let $\bar{M}_1 = \beta_1$, and $\bar{M}_2 = \beta_2$. $R_0 < 1$ implies $R_{01} < 1$ and $R_{02} < 1$. Lemma 3.1 implies $s(M_1) < 0$ and $s(M_2) < 0$. By the continuity of $s(M_1 + \varepsilon\bar{M}_1)$ and $s(M_2 + \varepsilon\bar{M}_2)$ in ε , ε can be chosen small enough so that $s(M_1 + \varepsilon\bar{M}_1) < 0$ and $s(M_2 + \varepsilon\bar{M}_2) < 0$. Consequently, the solutions of system (4) approach to zero with t going to infinity. The comparison principle of cooperative systems [16], Theorem B.1, implies $I_1(t) \rightarrow 0$ and $I_2(t) \rightarrow 0$ as $t \rightarrow \infty$. Therefore, the theory of asymptotically autonomous systems [17], Theorem 1.2, shows that $\lim_{t \rightarrow \infty} S_i(t) = S_i^0$ ($i = 1, 2$).

In the case of $R_0 > 1$, [4], Theorem 2, admits that P_0 is unstable, which finishes the theorem. □

Next, the two results regarding the stability of the boundary equilibria are given by applying the so-called Routh-Hurwitz criterion.

Theorem 4.2 *If $R_{01} > 1$ and $R_{02} < 1$, the boundary equilibrium P_{1^*} is stable when $S_2^{1^*} < S_2^{2^*}$; while the boundary equilibrium P_{1^*} is unstable when $S_2^{1^*} > S_2^{2^*}$.*

Proof $R_0 > 1$ and $R_{02} < 1$ imply that system (1) has a boundary equilibrium P_{1^*} . The Jacobian matrix of the right-hand side of system (1) at the equilibrium P_{1^*} , ordering coordinates as (S_1, S_2, I_1, I_2) , is given by

$$M(P_{1^*}) = \begin{bmatrix} -\beta_1 I_1^{1^*} - (\mu_1 + a_{21}) & a_{12} & -\beta_1 S_1^{1^*} & 0 \\ a_{21} & -(\mu_2 + a_{12}) & 0 & -\beta_2 S_2^{1^*} \\ \beta_1 I_1^{1^*} & 0 & 0 & 0 \\ 0 & 0 & 0 & \hat{b} \end{bmatrix},$$

where $\hat{b} = \beta_2 S_2^{1*} - (\mu_2 + d_2 + \gamma_2) = \beta_2(S_2^{1*} - S_2^{2*})$. Therefore, the eigenvalues are: \hat{b} and the solutions of the following cubic equation:

$$\lambda^3 + b_1 \lambda^2 + b_2 \lambda + b_3 = 0, \tag{5}$$

where

$$\begin{aligned} b_1 &= (\mu_1 + a_{21}) + (\mu_2 + a_{12}) + \beta_1 I_1^{1*} > 0, \\ b_2 &= \mu_1 \mu_2 + \mu_2 a_{21} + \mu_2 \beta_1 I_1^{1*} + \mu_1 a_{12} + a_{12} \beta_1 I_1^{1*} + \beta_1 S_1^{1*} \beta_1 I_1^{1*} > 0, \\ b_3 &= \beta_1 S_1^{1*} \beta_1 I_1^{1*} (\mu_2 + a_{12}) > 0. \end{aligned}$$

Since

$$\begin{aligned} b_1 b_2 - b_3 &= (\mu_1 + a_{21} + \beta_1 I_1^{1*}) b_2 \\ &\quad + (\mu_2 + a_{12})(\mu_1 \mu_2 + \mu_2 a_{21} + \mu_2 \beta_1 I_1^{1*} + \mu_1 a_{12} + a_{12} \beta_1 I_1^{1*}) > 0, \end{aligned}$$

Routh-Hurwitz criterion implies all the roots of Eq. (5) have a negative real part.

Therefore, $S_2^{1*} < S_2^{2*}$ yields the boundary equilibrium P_1^* is locally stable; while $S_2^{1*} > S_2^{2*}$ demonstrates the boundary equilibrium P_1^* is unstable. \square

Theorem 4.3 *If $R_{02} > 1$ and $R_{01} < 1$, the boundary equilibrium P_2^* is stable when $S_1^{2*} < S_1^{1*}$; while the boundary equilibrium P_2^* is unstable when $S_1^{2*} > S_1^{1*}$.*

Proof Because $R_{02} > 1$ and $R_{01} < 1$, there exists another boundary equilibrium P_2^* for system (1). The Jacobian matrix of the right-hand side of system (1) at the equilibrium P_2^* is denoted by

$$M(P_2^*) = \begin{bmatrix} -(\mu_1 + a_{21}) & -\beta_1 S_1^{2*} & a_{12} & 0 \\ 0 & \hat{c} & 0 & 0 \\ a_{21} & 0 & -\beta_2 I_2^{2*} - (\mu_2 + a_{12}) & -\beta_2 S_2^{2*} \\ 0 & 0 & \beta_2 I_2^{2*} & 0 \end{bmatrix},$$

where $\hat{c} = \beta_1 S_1^{2*} - (\mu_1 + d_1 + \gamma_1) = \beta_1(S_1^{2*} - S_1^{1*})$.

It is easy to see that all the eigenvalues of the matrix $M(P_2^*)$ are: \hat{c} and the roots of the following equation:

$$\lambda^3 + c_1 \lambda^2 + c_2 \lambda + c_3 = 0, \tag{6}$$

where

$$\begin{aligned} c_1 &= (\mu_1 + a_{21}) + (\mu_2 + a_{12}) + \beta_2 I_2^{2*} > 0, \\ c_2 &= \mu_1 \mu_2 + \mu_1 a_{12} + \mu_1 \beta_2 I_2^{2*} + \mu_2 a_{21} + a_{21} \beta_2 I_2^{2*} + \beta_2 S_2^{2*} \beta_2 I_2^{2*} > 0, \\ c_3 &= (\mu_1 + a_{21}) \beta_2 S_2^{2*} \beta_2 I_2^{2*} > 0. \end{aligned}$$

Because

$$c_1c_2 - c_3 = (\mu_2 + a_{12} + \beta_2I_2^{2*})c_3 + (\mu_1 + a_{21})(\mu_1\mu_2 + \mu_1a_{12} + \mu_1\beta_2I_2^{2*} + \mu_2a_{21} + a_{21}\beta_2I_2^{2*}) > 0,$$

by using the Routh-Hurwitz criterion, it then follows that the real part of all the solutions of (6) is negative.

Furthermore, it is easier to see that if $S_1^{2*} < S_1^{1*}$, the boundary equilibrium P_{2^*} is locally stable; while if $S_1^{2*} > S_1^{1*}$, the boundary equilibrium P_{2^*} is unstable. \square

Now we are in the position to discuss the global stability of the endemic equilibrium.

Theorem 4.4 *If the following statements hold:*

- (i) $R_{01} > 1$;
- (ii) $R_{02} > 1$;
- (iii) $S_1^{2*} > S_1^{1*}$;
- (iv) $S_2^{1*} > S_2^{2*}$;

*then the endemic equilibrium P_{**} is globally asymptotically stable.*

Proof Conditions (i)-(iv) imply system (1) exists the endemic equilibrium P_{**} . Next, we study the stability of the endemic equilibrium P_{**} by using the Lyapunov approach.

The following equations are derived at the endemic equilibrium P_{**} :

$$\begin{aligned} \Lambda_1 &= \beta_1S_1^{**}I_1^{**} + \mu_1S_1^{**} - a_{12}S_2^{**} + a_{21}S_1^{**}, \\ \mu_1 + d_1 + \gamma_1 &= \beta_1S_1^{**}, \\ \Lambda_2 &= \beta_2S_2^{**}I_2^{**} + \mu_2S_2^{**} - a_{21}S_1^{**} + a_{12}S_2^{**}, \\ \mu_2 + d_2 + \gamma_2 &= \beta_2S_2^{**}. \end{aligned} \tag{7}$$

Construct the following Lyapunov function:

$$U = S_1 - S_1^{**} \ln S_1 + I_1 - I_1^{**} \ln I_1 + A(S_2 - S_2^{**} \ln S_2) + A(I_2 - I_2^{**} \ln I_2), \tag{8}$$

where $A = \frac{a_{12}S_2^{**}}{a_{21}S_1^{**}}$.

Differentiating the function V along with the solutions of system (1) with respect to time t gives

$$\frac{dU}{dt} \Big|_{(1)} = \left(1 - \frac{S_1^{**}}{S_1}\right) \frac{dS_1}{dt} + \left(1 - \frac{I_1^{**}}{I_1}\right) \frac{dI_1}{dt} + A \left(1 - \frac{S_2^{**}}{S_2}\right) \frac{dS_2}{dt} + A \left(1 - \frac{I_2^{**}}{I_2}\right) \frac{dI_2}{dt}.$$

Combining system (1) admits

$$\begin{aligned} \frac{dU}{dt} \Big|_{(1)} &= \left(1 - \frac{S_1^{**}}{S_1}\right) (\Lambda_1 - \beta_1S_1I_1 - \mu_1S_1 + a_{12}S_2 - a_{21}S_1) \\ &\quad + \left(1 - \frac{I_1^{**}}{I_1}\right) [\beta_1S_1I_1 - (\mu_1 + d_1 + \gamma_1)I_1] \end{aligned}$$

$$\begin{aligned}
 &+ A \left(1 - \frac{S_2^{**}}{S_2}\right) (\Lambda_2 - \beta_2 S_2 I_2 - \mu_2 S_2 + a_{21} S_1 - a_{12} S_2) \\
 &+ A \left(1 - \frac{I_2^{**}}{I_2}\right) [\beta_2 S_2 I_2 - (\mu_2 + d_2 + \gamma_2) I_2].
 \end{aligned}$$

Applying Eq. (7) shows

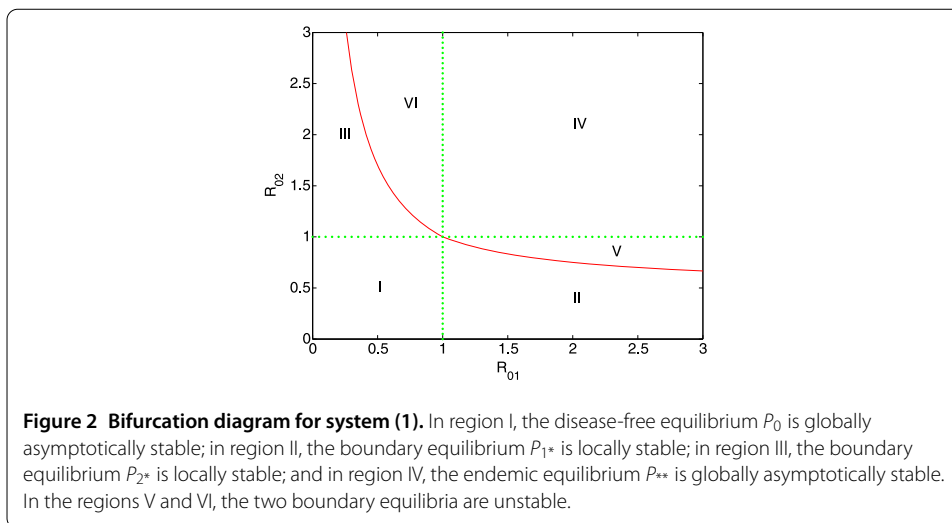
$$\begin{aligned}
 \left. \frac{dU}{dt} \right|_{(1)} &= \left(1 - \frac{S_1^{**}}{S_1}\right) (\beta_1 S_1^{**} I_1^{**} + \mu_1 S_1^{**} - a_{12} S_2^{**} + a_{21} S_1^{**} - \beta_1 S_1 I_1 - \mu_1 S_1 + a_{12} S_2 - a_{21} S_1) \\
 &+ A \left(1 - \frac{S_2^{**}}{S_2}\right) (\beta_2 S_2^{**} I_2^{**} + \mu_2 S_2^{**} - a_{21} S_1^{**} + a_{12} S_2^{**} \\
 &- \beta_2 S_2 I_2 - \mu_2 S_2 + a_{21} S_1 - a_{12} S_2) \\
 &+ \left(1 - \frac{I_1^{**}}{I_1}\right) (\beta_1 S_1 I_1 - \beta_1 S_1^{**} I_1) + A \left(1 - \frac{I_2^{**}}{I_2}\right) (\beta_2 S_2 I_2 - \beta_2 S_2^{**} I_2).
 \end{aligned}$$

Rearranging the above equation, it then follows

$$\begin{aligned}
 \left. \frac{dU}{dt} \right|_{(1)} &= \mu_1 S_1^{**} \left(1 - \frac{S_1^{**}}{S_1}\right) \left(1 - \frac{S_1}{S_1^{**}}\right) + \beta_1 S_1^{**} I_1^{**} \left(1 - \frac{S_1^{**}}{S_1}\right) \left(1 - \frac{S_1 I_1}{S_1^{**} I_1^{**}}\right) \\
 &+ a_{12} S_2^{**} \left(1 - \frac{S_1^{**}}{S_1}\right) \left(\frac{S_2}{S_2^{**}} - 1\right) + a_{21} S_1^{**} \left(1 - \frac{S_1^{**}}{S_1}\right) \left(1 - \frac{S_1}{S_1^{**}}\right) \\
 &+ \beta_1 S_1^{**} I_1^{**} \left(1 - \frac{I_1^{**}}{I_1}\right) \left(\frac{S_1 I_1}{S_1^{**} I_1^{**}} - \frac{I_1}{I_1^{**}}\right) + A \mu_2 S_2^{**} \left(1 - \frac{S_2^{**}}{S_2}\right) \left(1 - \frac{S_2}{S_2^{**}}\right) \\
 &+ A \beta_2 S_2^{**} I_2^{**} \left(1 - \frac{S_2^{**}}{S_2}\right) \left(1 - \frac{S_2 I_2}{S_2^{**} I_2^{**}}\right) + A a_{21} S_1^{**} \left(1 - \frac{S_2^{**}}{S_2}\right) \left(\frac{S_1}{S_1^{**}} - 1\right) \\
 &+ A a_{12} S_2^{**} \left(1 - \frac{S_2^{**}}{S_2}\right) \left(1 - \frac{S_2}{S_2^{**}}\right) + A \beta_2 S_2^{**} I_2^{**} \left(1 - \frac{I_2^{**}}{I_2}\right) \left(\frac{S_2 I_2}{S_2^{**} I_2^{**}} - \frac{I_2}{I_2^{**}}\right).
 \end{aligned}$$

By denoting $x := \frac{S_1}{S_1^{**}}$, $y := \frac{I_1}{I_1^{**}}$, $z := \frac{S_2}{S_2^{**}}$, and $w := \frac{I_2}{I_2^{**}}$, the above formula can be rewritten as

$$\begin{aligned}
 \left. \frac{dU}{dt} \right|_{(1)} &= \mu_1 S_1^{**} \left(1 - \frac{1}{x}\right) (1 - x) + \beta_1 S_1^{**} I_1^{**} \left(1 - \frac{1}{x}\right) (1 - xy) + a_{12} S_2^{**} \left(1 - \frac{1}{x}\right) (z - 1) \\
 &+ a_{21} S_1^{**} \left(1 - \frac{1}{x}\right) (1 - x) + \beta_1 S_1^{**} I_1^{**} \left(1 - \frac{1}{y}\right) (xy - y) + A \mu_2 S_2^{**} \left(1 - \frac{1}{z}\right) (1 - z) \\
 &+ A \beta_2 S_2^{**} I_2^{**} \left(1 - \frac{1}{z}\right) (1 - zw) + A a_{21} S_1^{**} \left(1 - \frac{1}{z}\right) (x - 1) \\
 &+ A a_{12} S_2^{**} \left(1 - \frac{1}{z}\right) (1 - z) + A \beta_2 S_2^{**} I_2^{**} \left(1 - \frac{1}{w}\right) (zw - w) \\
 &= (\mu_1 S_1^{**} + a_{21} S_1^{**}) \left(2 - x - \frac{1}{x}\right) + \beta_1 S_1^{**} I_1^{**} \left(2 - x - \frac{1}{x}\right) \\
 &+ A (\mu_2 S_2^{**} + a_{12} S_2^{**}) \left(2 - z - \frac{1}{z}\right) + A \beta_2 S_2^{**} I_2^{**} \left(2 - z - \frac{1}{z}\right) \\
 &+ a_{12} S_2^{**} \left(z - 1 - \frac{z}{x} + \frac{1}{x}\right) + A a_{21} S_1^{**} \left(x - 1 - \frac{x}{z} + \frac{1}{z}\right).
 \end{aligned}$$



Because of $A = \frac{a_{12}S_2^{**}}{a_{21}S_1^{**}}$, the above equation can be rewritten down as

$$\begin{aligned} \left. \frac{dU}{dt} \right|_{(1)} &= (\mu_1 S_1^{**} + a_{21} S_1^{**} + \beta_1 S_1^{**} I_1^{**}) \left(2 - x - \frac{1}{x} \right) \\ &\quad + A (\mu_2 S_2^{**} + a_{12} S_2^{**} + \beta_2 S_2^{**} I_2^{**}) \left(2 - z - \frac{1}{z} \right) \\ &\quad + a_{12} S_2^{**} \left[\left(z + \frac{1}{z} - 2 \right) + \left(2 - \frac{z}{x} - \frac{x}{z} \right) + \left(x + \frac{1}{x} - 2 \right) \right] \\ &= (\mu_1 S_1^{**} + a_{21} S_1^{**} + \beta_1 S_1^{**} I_1^{**} - a_{12} S_2^{**}) \left(2 - x - \frac{1}{x} \right) + a_{12} S_2^{**} \left(2 - \frac{z}{x} - \frac{x}{z} \right) \\ &\quad + A (\mu_2 S_2^{**} + a_{12} S_2^{**} + \beta_2 S_2^{**} I_2^{**} - a_{21} S_1^{**}) \left(2 - z - \frac{1}{z} \right). \end{aligned}$$

Using Eqs. (7) gives rise to

$$\left. \frac{dU}{dt} \right|_{(1)} = \Lambda_1 \left(2 - x - \frac{1}{x} \right) + A \Lambda_2 \left(2 - z - \frac{1}{z} \right) + a_{12} S_2^{**} \left(2 - \frac{z}{x} - \frac{x}{z} \right).$$

The inequality of arithmetic-geometric mean implies $dU/dt|_{(1)} \leq 0$. The equality holds if and only if $x = z = 1$. That is, when $S_1 = S_1^{**}$ and $S_2 = S_2^{**}$, $dV/dt|_{(1)} = 0$. By using the LaSalle invariant principle [8], the endemic equilibrium P^{**} is globally asymptotically stable. \square

Theorems 4.1-4.4 can be summarized in Figure 2. The basic reproduction numbers R_{01} and R_{02} are two important threshold parameters. It shows that if both R_{01} and R_{02} are less than one, the disease-free equilibrium P_0 is globally asymptotically stable and the disease eventually dies out (the region I in Figure 2); if R_{01} is greater than one, R_{02} is less than one, and $S_2^{1^*} < S_2^{2^*}$, the boundary equilibrium P_{1^*} is locally stable and the disease persists in patch one but can be eradicated in patch two (the region II in Figure 2); if R_{02} is greater than one, R_{01} is less than one, and $S_1^{2^*} < S_1^{1^*}$, the boundary equilibrium P_{2^*} is locally stable and the disease persists in patch two but can be eradicated in patch one (the region III in Figure 2); and if $R_{01} > 1$, $R_{02} > 1$, $S_2^{1^*} > S_2^{2^*}$, and $S_1^{2^*} > S_1^{1^*}$, there is exactly one endemic equi-

librium P^{**} , which is globally asymptotically stable by applying Lyapunov method, and the disease persists in two patches (the region IV in Figure 2). In addition, the two boundary equilibria are unstable in the regions V and VI.

5 Conclusions and discussions

In this paper, an *SIR* infectious diseases model with susceptibles dispersal between two disjoint patches has been proposed and analyzed to investigate the impact of susceptibles dispersal on diseases transmission in the whole population. The existence of equilibria is obtained and the basic reproduction numbers R_{01} , R_{02} , and R_0 are defined. It is indicated that R_{01} and R_{02} are two important threshold parameters to determine the long-term behavior of the solutions of system (1). The disease-free equilibrium is globally asymptotically stable and the disease ultimately dies out by applying the comparison principle of cooperative systems if the basic reproduction numbers both R_{01} and R_{02} are below unity. The disease persists in patch one and can be eradicated in patch two if R_{01} is above one, R_{02} is below one, and $S_2^{1*} < S_2^{2*}$. The disease persists in patch two and can be eradicated in patch one if R_{02} is above one, R_{01} is below one, and $S_1^{2*} < S_1^{1*}$. While the disease uniformly persists in the whole population and the endemic equilibrium is globally asymptotically stable by using the Lyapunov approach if the conditions $R_{01} > 1$, $R_{02} > 1$, $S_2^{1*} > S_2^{2*}$, and $S_1^{2*} > S_1^{1*}$ are satisfied.

System (1) almost shares the same qualitative behavior as the simple *SIR* epidemic model if dispersal can not be considered in the population. The patchy models need not be considered if only susceptibles disperse among patches. Furthermore, all the patches can be thought of as just one patch and susceptibles dispersal has no influence on disease transmission.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

LL proposed the model and completed the main part of this manuscript, WC checked all the theorems and polished the language, and YW enhanced the revised version. All the authors read and approved the manuscript.

Acknowledgements

The authors are grateful to the referee for her/his valuable remarks which led to improvement of the article. This work was supported in part by the National Nature Science Foundation of China (NSFC 11001215, 11101126, and 11101127), the Scientific Research Foundation for Doctoral Scholars of Haust (09001535), and the Foundation of Shaanxi Educational Committee (12JK0859).

Received: 10 January 2012 Accepted: 20 July 2012 Published: 1 August 2012

References

1. Brauer, F, van den Driessche, P: Models for transmission of disease with immigration of infectives. *Math. Biosci.* **171**, 143-154 (2001)
2. Brauer, F, van den Driessche, P, Wang, L: Oscillations in a patchy environment disease model. *Math. Biosci.* **215**, 1-10 (2008)
3. Centers for Disease Control and Prevention: Update: Novel influenza A (H1N1) virus infections-worldwide, May 6, 2009. *Morb. Mort. Wkly. Rep.* **58**, 453-458 (2009)
4. van den Driessche, P, Watmough, J: Reproduction numbers and sub-threshold endemic equilibria for compartmental models of disease transmission. *Math. Biosci.* **180**, 29-48 (2002)
5. Guo, H, Li, MY: Global dynamics of a staged progression model for infectious diseases. *Math. Biosci. Eng.* **3**, 513-525 (2006)
6. Guo, H, Li, MY: Global dynamics of a staged-progression model with amelioration for infectious diseases. *J. Biol. Dyn.* **2**, 154-168 (2008)
7. Hethcote, HW: Qualitative analysis of communicable disease models. *Math. Biosci.* **28**, 335-356 (1976)
8. LaSalle, JP: *The Stability of Dynamical Systems*. SIAM, Philadelphia (1976)
9. Li, M, Shuai, Z: Global-stability problem for coupled systems of differential equations on networks. *J. Differ. Equ.* **248**, 1-20 (2010)

10. Li, MY, Shuai, Z, Wang, C: Global stability of multi-group epidemic models with distributed delays. *J. Math. Anal. Appl.* **361**, 38-47 (2010)
11. Liu, L, Zhou, Y, Wu, J: Global dynamics in a TB model incorporating case detection and two treatment stages. *Rocky Mt. J. Math.* **38**, 1541-1559 (2008)
12. Lloyd, A, May, RM: Spatial heterogeneity in epidemic models. *J. Theor. Biol.* **179**, 1-11 (1996)
13. Mccluskey, CC: Lyapunov functions for tuberculosis models with fast and slow progression. *Math. Biosci. Eng.* **3**, 603-614 (2006)
14. O'Regan, SM, Kelly, TC, Korobeinikov, A, O'Callaghan, MJA, Pokrovskii, AV: Lyapunov functions for *SIR* and *SIRS* epidemic models. *Appl. Math. Lett.* **23**, 446-448 (2010)
15. Smith, HL: *Monotone Dynamical Systems: An Introduction to the Theory of Competitive and Cooperative Systems*. Mathematical Surveys and Monographs, vol. 41. Am. Math. Soc, Providence (1995)
16. Smith, HL, Walman, P: *The Theory of the Chemostat*. Cambridge Univ. Press, Cambridge (1995)
17. Thieme, HR: Convergence results and a Poincaré-Bendixon trichotomy for asymptotical autonomous differential equations. *J. Math. Biol.* **30**, 755-763 (1992)
18. Wang, W, Zhao, X-Q: An epidemic model in a patchy environment. *Math. Biosci.* **190**, 97-112 (2004)
19. World Health Organization: Summary table of SARS cases by country, 1 November 2002-7 August 2003, http://www.who.int/csr/sars/country/2003_8_5/en. Accessed 15 August 2003
20. World Health Organization: WHO guidelines for the global surveillance of severe acute respiratory syndrome (SARS). http://www.who.int/csr/resources/publications/WHO_CDS_CSR_ARO_2004_1/en/index.html

doi:10.1186/1687-1847-2012-131

Cite this article as: Liu et al.: Global dynamics for an SIR patchy model with susceptibles dispersal. *Advances in Difference Equations* 2012 2012:131.

Submit your manuscript to a SpringerOpen[®] journal and benefit from:

- ▶ Convenient online submission
- ▶ Rigorous peer review
- ▶ Immediate publication on acceptance
- ▶ Open access: articles freely available online
- ▶ High visibility within the field
- ▶ Retaining the copyright to your article

Submit your next manuscript at ▶ springeropen.com
