Existence of weak solutions for two point boundary value problems of Schrödingerian predator-prey system and their applications

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Abstract

By means of a variational analysis and the theory of variable exponent Sobolev spaces, the existence of weak solutions for two point boundary value problems of Schrödingerian predator-prey system with latent periods is investigated either analytically or numerically. More precisely, the local stability of the Schrödingerian equilibrium and endemic equilibrium of the model are discussed in detail. And we specially analyzed the existence and stability of the Schrödingerian Hopf bifurcation by using the center manifold theorem and the bifurcation theory. As applications, theoretic analysis and numerical simulations show that the Schrödingerian predator-prey system with latent period has very rich dynamic characteristics.

Keywords: existence; stability; Schrödingerian predator-prey system; boundary value problem

1 Introduction

The role of mathematical modeling has been intensively growing in the study of epidemiology. Various epidemic models have been proposed and explored extensively and great progress has been achieved in the studies of disease control and prevention. Many authors have investigated the autonomous epidemic models. May and Odter [1] proposed a time-periodic reaction-diffusion epidemic model which incorporates a simple demographic structure and the latent period of an infectious disease. Guckenheimer and Holmes [2] examined an SIR epidemic model with a non-monotonic incidence rate, and they also analyzed the dynamical behavior of the model and derived the stability conditions for the disease-free and the endemic equilibrium. Berryman and Millstein [3] investigated an SVEIS epidemic model for an infectious disease that spreads in the host population through horizontal transmission, and they have shown that the model exhibits two equilibria, namely, the disease-free equilibrium and the endemic equilibrium. Hassell et al. [4] presented four discrete epidemic models with the nonlinear incidence rate by using the forward Euler and backward Euler methods, and they discussed the effect of two discretizations on the stability of the endemic equilibrium for these models. Shilnikov et al. [5] proposed an VEISV network worm attack model and derived global stability of a worm-free state and local stability of a unique worm-epidemic state by using the reproduction rate. Robinson and Holmes [6] discussed the dynamical behaviors of a Schrödingerian...
predator-prey system, and they showed that the model undergoes a flip bifurcation and Hopf bifurcation by using the center manifold theorem and bifurcation theory. Bacaër and Dads [7] investigated an SVEIS epidemic model for an infectious disease that spreads in the host population through horizontal transmission.

Recently, Yan et al. [8, 9] and Xue [10] discussed the threshold dynamics of a time-periodic reaction-diffusion epidemic model with latent period. In this paper, we will study the existence of the disease-free equilibrium and endemic equilibrium, and the stability of the disease-free equilibrium and the endemic equilibrium for this system. Conditions will be derived for the existence of a flip bifurcation and a Hopf bifurcation by using the center manifold theorem [11] and bifurcation theory [12–14].

The rest of this paper is organized as follows. A discrete SIR epidemic model with latent period is established in Section 2. In Section 3 we obtain the main results: the existence and local stability of fixed points for this system. We show that this system undergoes the flip bifurcation and the Hopf bifurcation by choosing a bifurcation parameter in Section 4. A brief discussion is given in Section 5.

2 Model formulation
In 2015, Yan et al. [10] discussed the threshold dynamics of a time-periodic reaction-diffusion epidemic model with latent period. We consider the continuous-time SIR epidemic model described by the differential equations

\[
\begin{align*}
\frac{dS}{dt} &= \beta S(t)I(t), \\
\frac{dI}{dt} &= \beta S(t)I(t) - \gamma I(t), \\
\frac{dR}{dt} &= \gamma I(t),
\end{align*}
\]  

(1)

where \(S(t), I(t)\) and \(R(t)\) denote the sizes of the susceptible, infected and removed individuals, respectively, the constant \(\beta\) is the transmission coefficient, and \(\gamma\) is the recovery rate. Let \(S_0 = S(0)\) be the density of the population at the beginning of the epidemic with everyone susceptible. It is well known that the basic reproduction number \(R_0 = \beta S_0/\gamma\) completely determines the transmission dynamics (an epidemic occurs if and only if \(R_0 > 1\)); see also [11]. It should be emphasized that system (1) has no vital dynamics (births and deaths) because it was usually used to describe the transmission dynamics of disease within a short outbreak period. However, for an endemic disease, we should incorporate the demographic structure into the epidemic model. The classical endemic model is the following SIR model with vital dynamics:

\[
\begin{align*}
\frac{dS}{dt} &= \mu N - \mu S(t) - \frac{\beta S(t)I(t)}{N}, \\
\frac{dI}{dt} &= \frac{\beta S(t)I(t)}{N} - \gamma I(t) - \mu I(t), \\
\frac{dR}{dt} &= \gamma I(t) - \mu I(t),
\end{align*}
\]  

(2)

which is almost the same as the SIR epidemic model (1) above, except that it has an inflow of newborns into the susceptible class at rate \(\mu N\) and deaths in the classes at rates \(\mu N, \mu I\) and \(\mu R\), where \(N\) is a positive constant denoting the total population size. For this model, the basic reproduction number is given by

\[R_0 = \frac{\beta}{\gamma + \mu},\]
which is the contact rate times the average death-adjusted infectious period $\frac{1}{\gamma + \mu}$. The disease-free equilibrium $E_0(N, 0, 0)$ of model (2) is as follows:

$$
\begin{cases}
S_{n+1} = S_n + h(\mu N - \mu S_n - \frac{\beta S_n I_n}{N} ), \\
I_{n+1} = I_n + h(\frac{\beta S_n I_n}{N} - \gamma I_n - \mu I_n), \\
R_{n+1} = R_n + h(\gamma I_n - \mu I_n),
\end{cases}
$$

(3)

where $h$, $N$, $\mu$, $\beta$ and $\gamma$ are all defined in (2).

**Remark 1** If the basic reproductive rate $R_0 < 1$, then model (2) has only a disease-free equilibrium $E_1(N, 0)$. If the basic reproductive rate $R_0 > 1$, then model (2) has two equilibria: a disease-free equilibrium $E_1(N, 0)$ and an endemic equilibrium $E_2(S^*, I^*)$, where

$$
S^* = \frac{N(\gamma + \mu)}{\beta} \quad \text{and} \quad I^* = \frac{N(\beta \mu - \mu(\gamma + \mu))}{\beta(\gamma + \mu)}.
$$

**3 Main results**

We firstly discuss the existence of the equilibria of model (2) by using a linearization method and the Jacobian matrix. The Jacobian matrix of $E_1$ is defined by

$$
J(E_1) = \begin{pmatrix}
1 - h\mu & -h\beta \\
0 & 1 + h\beta - h(\gamma + \mu)
\end{pmatrix}.
$$

If we take the two eigenvalues of $J(E_1)$

$$
\omega_1 = 1 - h\mu \quad \text{and} \quad \omega_2 = 1 + h\beta - h(\gamma + \mu),
$$

then we have the following results from Remark 1 and a simple calculation.

**Theorem 1** Let $R_0$ be the basic reproductive rate such that $R_0 < 1$. Then:

(1) If

$$
0 < h < \min \left\{ \frac{2}{\mu}, \frac{2}{(\gamma + \mu) - \beta} \right\},
$$

then $E_1(N, 0)$ is asymptotically stable.

(2) If

$$
h > \max \left\{ \frac{2}{\mu}, \frac{2}{(\gamma + \mu) - \beta} \right\} \quad \text{or} \quad \frac{2}{\mu} < h < \frac{2}{(\gamma + \mu) - \beta}
$$

or

$$
\frac{2}{(\gamma + \mu) - \beta} < h < \frac{2}{\mu},
$$

then $E_1(N, 0)$ is unstable.
If
\[ h = \frac{2}{\mu} \quad \text{or} \quad h = \frac{2}{(\gamma + \mu) - \beta}, \]
then \( E_1(N, 0) \) is non-hyperbolic.

The Jacobian matrix of model (2) at \( E_2(S^*, I^*) \) is
\[
J(E_2) = \begin{pmatrix}
1 - \frac{h \mu \beta}{\gamma + \mu} & -h(\gamma + \mu) \\
\frac{h \mu}{\gamma + \mu} (\beta - \gamma - \mu) & 1
\end{pmatrix},
\]
which gives
\[
F(\omega) = \omega^2 - \text{tr} J(E_2) \omega + \text{det} J(E_2), \tag{4}
\]
where
\[
\text{tr} J(E_2) = 2 - \frac{h \mu \beta}{\gamma + \mu} \tag{5}
\]
and
\[
\text{det} J(E_2) = 1 - \frac{h \mu \beta}{\gamma + \mu} + h^2 \left[ \mu \beta - (\gamma + \mu) \right]. \tag{6}
\]

The two eigenvalues of \( J(E_2) \) are
\[
\omega_{1,2} = 1 + \frac{1}{2} \left( -\frac{h \mu \beta}{\gamma + \mu} \pm \sqrt{(h R_0)^2 - 4[\mu \beta - \mu(\gamma + \mu)]} \right). \tag{7}
\]

Next we obtain the following result for \( E_2(S^*, I^*) \) by Remark 1 and a simple calculation.

**Theorem.** Let \( R_0 \) be the basic reproductive rate such that \( R_0 < 1 \). Then:

1. **Put**
   - (A) \( 0 < h < h_*, \) and \( (\mu R_0)^2 - 4[\mu \beta - \mu(\gamma + \mu)] \geq 0, \)
   - (B) \( 0 < h < h_{**} \) and \( (\mu R_0)^2 - 4[\mu \beta - \mu(\gamma + \mu)] < 0. \)
   
   If one of the above conditions holds, then we see that \( E_2(S^*, I^*) \) is asymptotically stable.

2. **Put**
   - (A) \( h > h_{**} \) and \( (\mu R_0)^2 - 4[\mu \beta - \mu(\gamma + \mu)] \geq 0; \)
   - (B) \( 0 < h < h_{**} \) and \( (\mu R_0)^2 - 4[\mu \beta - \mu(\gamma + \mu)] < 0; \)
   - (C) \( h_1 < h < h_{**} \) and \( (\mu R_0)^2 - 4[\mu \beta - \mu(\gamma + \mu)] \geq 0. \)
   
   If one of the above conditions holds, then \( E_2(S^*, I^*) \) is unstable.

3. **Put**
   - (A) \( h = h_* \) or \( h = h_{**} \) and \( (\mu R_0)^2 - 4[\mu \beta - \mu(\gamma + \mu)] \geq 0; \)
   - (B) \( h = h_{**} \) and \( (\mu R_0)^2 - 4[\mu \beta - \mu(\gamma + \mu)] < 0; \)
where
\[ h^* = \frac{\mu \beta - \mu (\gamma + \mu) \sqrt{\mu R_0^2 - 4[\mu \beta - \mu (\gamma + \mu)]}}{(\gamma + \mu)[\mu \beta - \mu (\gamma + \mu)]}, \]
\[ h^{**} = \frac{\mu \beta}{(\gamma + \mu)[\mu \beta - \mu (\gamma + \mu)]}, \]
and
\[ h^{***} = \frac{\mu \beta + \mu (\gamma + \mu) \sqrt{\mu R_0^2 - 4[\mu \beta - \mu (\gamma + \mu)]}}{(\gamma + \mu)[\mu \beta - \mu (\gamma + \mu)]}. \]

If one of the above conditions holds, then \( E_2(S^*, I^*) \) is non-hyperbolic.

By a simple calculation, Conditions (A) in Theorem 2 can be written in the following form:

\[ (\mu, N, \beta, h, \gamma) \in M_1 \cup M_2, \]

where
\[ M_1 = \{(\mu, N, \beta, h, \gamma) : h = h^*, N > 0, \Delta > 0, 0 < \mu, \beta, \gamma < 1\} \]

and
\[ M_2 = \{(\mu, N, \beta, h, \gamma) : h = h^{***}, N > 0, R_0 > 1, 0 < \mu, \beta, \gamma < 1\}. \]

It is well known that if \( h^* \) varies in a small neighborhood of \( h^* \) or \( h^{***} \) and \( (\mu, N, \beta, h, \gamma) \in M_1 \) or \( (\mu, N, \beta, h^{***}, \gamma) \in M_2 \), then there may be a flip bifurcation of equilibrium \( E_2(S^*, I^*) \).

4 Bifurcation Analysis

If \( h \) varies in a neighborhood of \( h^* \) and \( (\mu, N, \beta, h^*, \gamma) \in M_1 \), then we derive the flip bifurcation of model (2) at \( E_2(S^*, I^*) \). In particular, in the case that \( h \) changes in the neighborhood of \( h^{***} \) and \( (\mu, N, \beta, h^{***}, \gamma) \in M_2 \) we need to make a similar calculation.

Set
\[ (\mu, N, \beta, h, \gamma) = (\mu_1, N_1, \beta_1, h_1, \gamma_1) \in M_1. \]

If we give the parameter \( h_1 \) a perturbation \( h^* \), model (2) is considered as follows:

\[ \begin{aligned}
S_{n+1} &= S_n + (r^* + h_1)(\mu_1 N_1 S_n - \frac{\beta_1 S_n I_n}{N_1}), \\
I_{n+1} &= I_n + (h^* + h_1)(\frac{\beta_1 S_n I_n}{N_1} - \gamma I_n - \mu_1 I_n),
\end{aligned} \]

where \(|h^*| < 1\).

Put \( U_n = S_n - S^* \) and \( V_n = I_n - I^* \). We have

\[ \begin{aligned}
U_{n+1} &= a_{11} U_n + a_{12} V_n + a_{13} U_n V_n + b_{11} U_n h^* + b_{12} V_n h^* + b_{13} U_n V_n h^*, \\
V_{n+1} &= a_{21} U_n + a_{22} V_n + a_{23} U_n V_n + b_{21} U_n h^* + b_{22} V_n h^* + b_{23} U_n V_n h^*,
\end{aligned} \]
where

\[
\begin{align*}
    a_{11} &= 1 - h_1 \left( \mu_1 + \frac{\beta_1 I^*}{N_1} \right), \\
    a_{12} &= -\frac{h_1 \beta_1 S^*}{N_1}, \\
    a_{13} &= -\frac{h_1 \beta_1}{N_1}, \\
    b_{11} &= - \left( \mu_1 + \frac{\beta_1 I^*}{N_1} \right), \\
    b_{12} &= -\frac{\beta_1 S^*}{N_1}, \\
    b_{13} &= -\frac{\beta_1}{N_1}, \\
    a_{21} &= \frac{h_1 \beta_1 I^*}{N_1}, \\
    a_{22} &= 1, \\
    a_{23} &= -\frac{\beta_1 h_1}{N_1}, \\
    b_{21} &= \frac{\beta_1 I^*}{N_1}, \\
    b_{22} &= 0, \\
    b_{23} &= \frac{\beta_1}{N_1}.
\end{align*}
\]

If we define matrix \( T \) as follows:

\[
T = \begin{pmatrix}
    a_{12} & a_{12} \\
    -1 - a_{11} & \omega_2 - a_{11}
\end{pmatrix},
\]

then we know that \( T \) is invertible. If we use the transformation

\[
\begin{pmatrix}
    U_n \\
    V_n
\end{pmatrix} = T \begin{pmatrix}
    X_n \\
    Y_n
\end{pmatrix}
\]

then model (2) becomes

\[
\begin{align*}
    X_{n+1} &= -X_n + F(U_n, V_n, h^*), \\
    Y_{n+1} &= -\omega_2 Y_n + G(U_n, V_n, h^*). \\
\end{align*}
\]

Thus

\[
W^c(0, 0) = \{(X_n, Y_n) : Y_n = a_1 X_n^2 + a_2 X_n h^* + o((|X_n| + |h^*|)^2)\},
\]

where \( o((|X_n| + |h^*|)^2) \) is a transform function, and

\[
a_1 = \frac{a_{12}(1 + a_{11})}{\omega_2 + 1}
\]

and

\[
a_2 = \frac{b_{12}(1 + a_{11})^2}{a_{12}(\omega_2 + 1)^2} - \frac{a_{12} b_{12} + b_{11}(1 + a_{11})}{(\omega_2 + 1)^2}.
\]

Further we find that the manifold \( W^c(0, 0) \) has the following form:

\[
\begin{align*}
    c_1 &= \frac{a_{11}(1 + a_{11})(\omega_2 - a_{11} + 1)}{\omega_2 + 1}, \\
    c_2 &= -\frac{b_{11}(\omega_2 - a_{11}) - a_{12} b_{21}}{\omega_2 + 1} - \frac{b_{12}(\omega_2 - a_{11})(1 + a_{11})}{a_{12}(\omega_2 + 1)}, \\
    c_3 &= \frac{a_{13}(\omega_2 - 2a_{11} - 1)(\omega_2 - a_{11} + 1) - b_{13}(1 + a_{11})(\omega_2 - a_{11} + 1)}{\omega_2 + 1},
\end{align*}
\]
and
\[ c_4 = 0, \quad c_5 = \frac{a_1a_{13}(\omega_2 - 2a_{11} - 1)(\omega_2 - a_{11} + a_{12})}{\omega_2 + 1}. \]
Therefore the map \( G^* \) with respect to \( W^c(0, 0) \) can be defined by
\[
G^*(X_n) = -X_n + c_1X_n^3 + c_2X_nh^* + c_3X_nh^2 + c_4X_nh^2
+ c_5X_n^3 + o\left(\left(|X_n| + |h^*|\right)^3\right).
\]
In order to calculate map (11), we need two quantities \( \alpha_1 \) and \( \alpha_2 \) which are not zero,
\[
\alpha_1 = \left(\frac{G^*_{X_n,h^*} + \frac{1}{2}G^*_{X_n}}{G^*_{X_n}}\right)_{0,0},
\]
and
\[
\alpha_2 = \left(\frac{1}{6}G^*_{X_nX_nX_n} + \left(\frac{1}{2}G^*_{X_n}\right)^2\right)_{0,0}.
\]
By a simply calculation, we obtain
\[
\alpha_1 = c_2 = -\frac{2}{h_1},
\]
\[
\alpha_2 = c_5 + c_1^2 = \frac{h_1\beta_1}{N_1(\omega_2 + 1)}\left\{2 - h_1\beta_1\mu_1(2 - h_1\gamma_1)\right\}^2,
\]
where
\[
c_1 = \frac{h_1\beta_1\mu_1}{\gamma_1\mu_1}\left[\gamma_1(h_1 + \mu_1) - 2\right\{2 + \left[h_1(\gamma_1 + \mu_1) + \frac{h_1\beta_1\mu_1}{\gamma_1\mu_1}\right] \right].
\]
Therefore we have the following result.

**Theorem 3.** Let \( h \) change in a neighborhood of the origin. If \( \alpha_2 \neq 0 \), then the model (9) has a flip bifurcation at \( E_2(S^*, I^*) \). If \( \alpha_2 > 0 \), then the period-2 points that bifurcation from \( E_2(S^*, I^*) \) are stable. If \( \alpha_2 < 0 \), then it is unstable.

We further consider the bifurcation of \( E_2(S^*, I^*) \) if \( h \) varies in a neighborhood of \( h_{\text{max}} \).

Taking the parameters \((\mu, N, \beta, h, \gamma) = (\mu_2, N_2, \beta_2, h_2, \gamma_2) \in N^* \) arbitrarily, and also giving \( h \) a perturbation \( h^* \) at \( h_2 \), then model (2) gets the following form:

\[
\begin{align*}
S_{n+1} &= S_n + (h^* + h_2)(\mu_2N_2 - \mu_2S_n - \frac{\beta_2S_nI_n}{N_2}), \\
I_{n+1} &= I_n + (h^* + h_2)(\frac{\beta_2S_nI_n}{N_2} - \gamma_2I_n - \mu_2I_n).
\end{align*}
\]

Put \( U_n = S_n - S^* \) and \( V_n = I_n - I^* \). We change the equilibrium \( E_2(S^*, I^*) \) of model (9) and have the following result:

\[
\begin{align*}
U_{n+1} &= U_n + (h^* + h_2)(-\mu_2U_n - \frac{\beta_2}{N_2}U_nV_n - \frac{\beta_2}{N_2}U_nI^* - \frac{\beta_2}{N_2}V_nS^*), \\
V_{n+1} &= V_n + (h^* + h_2)(\frac{\beta_2}{N_2}U_nV_n - (\gamma_1 + \mu_1)V_n + \frac{\beta_2}{N_2}U_nI^* + \frac{\beta_2}{N_2}V_nS^*),
\end{align*}
\]
which gives

\[ \omega_2 + P(h^*) \omega + Q(h^*) = 0, \]

where

\[ 2 + P(h^*) = \frac{\beta_2 \mu_2 (h_2 + h^*)}{\gamma_2 \mu_2} \]

and

\[ Q(h^*) = 1 - \frac{\beta_2 \mu_2 (h_2 + h^*)}{\gamma_2 \mu_2} + (h_2 + h^*)^2 \left[ \mu_2 \beta_2 - \mu_2 (\mu_2 + \gamma_2) \right]. \]

It is easy to see that

\[ \omega_{1,2} = \frac{-P(h^*) \pm \sqrt{(P(h^*))^2 - 4Q(h^*)}}{2}, \]

which yields

\[ |\omega_{1,2}| = \sqrt{Q(h^*)}, \quad k = \left. \frac{d|\omega_{1,2}|}{dh^*} \right|_{h^*=0} = \frac{\mu_2 \beta_2}{2(\mu_2 + \gamma_2)}. \]

We remark that \((\mu_2, N_2, h_2, \gamma_2) \in \mathbb{N}^+ \Delta < 1\) and then we have

\[ \frac{(\mu_2 \beta_2)^2}{(\gamma_2 + \mu_2)^2[\mu_2 \beta_2 - \mu_2 (\mu_2 + \gamma_2)]} \Delta. \]

Thus

\[ P(0) = -2 + \frac{(\mu_2 \beta_2)^2}{(\gamma_2 + \mu_2)^2[\mu_2 \beta_2 - \mu_2 (\mu_2 + \gamma_2)]} \neq \pm 2, \]

which means that

\[ \frac{\mu_2 \beta_2}{(\gamma_2 + \mu_2)^2[\mu_2 \beta_2 - \mu_2 (\mu_2 + \gamma_2)]} \neq \frac{1}{j}, \quad j = 2, 3. \quad (14) \]

Hence, the eigenvalues \(\omega_{1,2}\) of equilibrium \((0,0)\) of model \((14)\) do not lay in the intersection when \(h^* = 0\) and \((14)\) holds.

When \(h^* = 0\) we may begin to study the model \((14)\). Put

\[ \alpha = \frac{(\mu_2 \beta_2)^2}{2(\gamma_2 + \mu_2)^2[\mu_2 \beta_2 - \mu_2 (\mu_2 + \gamma_2)]}, \]

\[ \beta = \frac{\mu_2 \beta_2 \sqrt{4[\mu_2 \beta_2 - \mu_2 (\mu_2 + \gamma_2)] - (\mu_2 \beta_2)^2}}{2(\gamma_2 + \mu_2)[\mu_2 \beta_2 - \mu_2 (\mu_2 + \gamma_2)]}, \]

and

\[ T = \begin{pmatrix} 0 & 1 \\ \beta & \alpha \end{pmatrix}, \]

where \(T\) is invertible.
If we use the transformation
\[
\begin{pmatrix} U_n \\ V_n \end{pmatrix} = T \begin{pmatrix} X_n \\ Y_n \end{pmatrix},
\]
then the model (14) gets the following form:
\[
\begin{cases}
X_{n+1} = \alpha X_n - \beta Y_n + \bar{F}(X_n, Y_n), \\
Y_{n+1} = \beta X_n + \alpha Y_n + \bar{G}(X_n, Y_n),
\end{cases}
\]
where
\[
\bar{F}(X_n, Y_n) = \frac{h_2 \beta (1 + \alpha)(\beta X_n Y_n + \alpha Y_n^2)}{N_2 \beta},
\]
and
\[
\bar{G}(X_n, Y_n) = \frac{-h_2 \beta (\beta X_n Y_n + \alpha Y_n^2)}{N_2}.
\]
Moreover,
\[
\begin{align*}
\bar{F}_{X_n X_n} &= 0, & \bar{F}_{Y_n Y_n} &= \frac{2 h_2 \beta (1 + \alpha)}{N_2 \beta}, & \bar{F}_{X_n Y_n} &= \frac{h_2 \beta (1 + \alpha)}{N_2}, \\
\bar{G}_{X_n X_n} &= 0, & \bar{G}_{Y_n Y_n} &= \frac{2 h_2 \beta \alpha}{N_2}, & \bar{G}_{X_n Y_n} &= \frac{-h_2 \beta}{N_2}, \\
\bar{G}_{X_n X_n X_n} &= \bar{G}_{X_n X_n Y_n} = \bar{G}_{X_n Y_n Y_n} = \frac{2 h_2 \beta}{N_2}, & \bar{G}_{X_n Y_n Y_n} &= \frac{-h_2 \beta}{N_2},
\end{align*}
\]
Thus we have
\[
a = \frac{1}{8} \left[ 1 - \frac{2 \beta}{\omega} \xi_{11} \xi_{20} \right] - \frac{1}{2} \| \xi_{11} \|^2 - \| \xi_{02} \|^2 + \text{Re}(\bar{F} \xi_{21}),
\]
where
\[
\begin{align*}
\xi_{02} &= \frac{1}{8} \left[ (\bar{F}_{X_n X_n} - \bar{F}_{Y_n Y_n}) - 2(\bar{G}_{X_n X_n} + 2 \bar{F}_{X_n Y_n}) \right], \\
\xi_{11} &= \frac{1}{4} \left[ (\bar{F}_{X_n X_n} + \bar{F}_{Y_n Y_n}) + (\bar{G}_{X_n X_n} + \bar{G}_{Y_n Y_n}) \right], \\
\xi_{20} &= \frac{1}{8} \left[ (\bar{F}_{X_n X_n} - \bar{F}_{Y_n Y_n} + 2 \bar{G}_{X_n Y_n}) + (\bar{G}_{X_n X_n} - \bar{G}_{Y_n Y_n} - 2 \bar{F}_{X_n Y_n}) \right],
\end{align*}
\]
and
\[
\xi_{21} = \frac{1}{16} (\bar{F}_{X_n X_n X_n} + \bar{F}_{X_n X_n Y_n} + \bar{G}_{X_n X_n Y_n} + \bar{G}_{Y_n Y_n Y_n}).
\]
Therefore we have the following result.
Theorem 4 Let $a \neq 0$ and $h^*$ change in a neighborhood of $h_{***}$. If the condition (15) holds, then model (13) undergoes a Hopf bifurcation at $E_2(S^*, I^*)$. If $a > 0$, then the repelling invariant closed curve bifurcates from $E_2$ for $h^* < 0$. If $a < 0$, then an attracting invariant closed curve bifurcates from $E_2$ for $h^* > 0$.

5 Conclusions
The paper investigated the basic dynamic characteristics of a Schrödingerian predator-prey system with latent period. First, we applied the forward Euler scheme to a continuous-time SIR epidemic model and obtained the Schrödingerian predator-prey system. Then, the existence and local stability of the disease-free equilibrium and endemic equilibrium of the model were discussed. In addition, we chose $h$ as the bifurcation parameter and studied the existence and stability of flip bifurcation and Hopf bifurcation of this model by using the center manifold theorem and the bifurcation theory. Numerical simulation results show that the model (2) shows a flip bifurcation and Hopf bifurcation when the bifurcation parameter $h$ passes through the respective critical value, and the direction and stability of flip bifurcation and Hopf bifurcation can be determined by the sign of $a_2$ and $a_1$, respectively. Apparently there are more interesting problems as regards this Schrödingerian predator-prey system with latent period which deserve further investigation.

Acknowledgements
The authors would like to express their deep-felt gratitude for the reviewer’s detailed reviewing and useful comments.

Competing interests
The authors declare that they have no competing interests.

Authors’ contributions
The first draft was prepared by TÜ. The final version was prepared by FL, which was verified and improved by TÜ. Both authors read and approved the final manuscript.

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