

Research Article

Dynamical Complexity of a Spatial Phytoplankton-Zooplankton Model with an Alternative Prey and Refuge Effect

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The spatiotemporal dynamics of a phytoplankton-zooplankton model with an alternative prey and refuge effect is investigated mathematically and numerically. The stability of the equilibrium point and the traveling wave solution of the phytoplankton-zooplankton model are described based on theoretical mathematical work, which provides the basis of the numerical simulation. The numerical analysis shows that refuges have a strong effect on the spatiotemporal dynamics of the model according to the pattern formation. These results may help us to understand prey-predator interactions in water ecosystems. They are also relevant to research into phytoplankton-zooplankton ecosystems.

1. Introduction

In recent years, the degradation of water ecosystems has exacerbated certain algal species in aquatic systems [1]. Increasing attention is being paid to preventing this exacerbation from reaching a critical condition because of adverse impacts on fisheries and aquaculture [1]. In aquatic systems, there is an ecological threshold below which species remains essentially undetected, whereas they seize the opportunity for unchecked growth if the ecological threshold is exceeded [2]. Predator-prey models have been studied widely and it is well known that these model can directly reflect changes in the sizes of populations, which can be used to determine the ecological threshold for certain algal species in aquatic systems.

Since the pioneering work of Lotka and Volterra, the predator-prey model has developed significantly in mathematical ecology [3]. A variety of models have been studied including the Wangersky-Cunningham model, the classical prey-dependent predator-prey model, and the ratio-dependent function response model [4–13]. A previous model studied the global dynamics of a predator-prey model with a nonconstant death rate and diffusion [14]. This model incorporated Holling type II and Leslie-Gower functional responses [15, 16]. Most authors have focused their attention on a single prey item known as the focal prey in these studies, although some studies have show that the presence of alternative food sources can affect biological control via a variety of mechanisms [17]. The population of a prey species may be affected negatively by another prey species because they may lead to higher predation rates for both prey items if the prey shares the predator [17]. However, alternative prey can also offset predation on the focal prey if the predator has a predilection for the alternative prey [17, 18]. In this case, the alternative prey may have a positive effect on the population of the focal prey [17].

In a predator-prey model, there are solution states in some cases. However, the transformation between these states is important. Thus, many researchers have studied the presence of traveling wave solutions in predator-prey models [19–25]. In the previous works, we mainly discussed the dynamics of prey-predator system with impulsive control strategy [3, 26–28].

In this paper, we discuss a two-species predator-prey model and the effect of alternative prey. This paper is organized as follows. In Section 2, we introduce the predator-prey model and the reaction-diffusion term. Section 3 presents the mathematical analysis, including the existence of an equilibrium point and boundedness, a stability analysis of the equilibrium point, and traveling wave solutions. In Section 4, we describe a numerical simulation of our analytical results, which helps us to understand the feasibility of the theorem. The final section provides our conclusions.

2. The Model

First, we introduce the following model:

$$\frac{dX}{d\tau} = r_1 X \left(1 - \frac{X}{k_1} \right) - \frac{a_1 \left(1 - m_1 \right) X}{b_1 + \left(1 - m_1 \right) X} Y,$$

$$\frac{dY}{d\tau} = \frac{a_1 e_1 \left(1 - m_1 \right) X}{b_1 + \left(1 - m_1 \right) X} Y + s_1 Y \left(1 - \frac{X}{k_1} \right) - d_1 Y,$$
(1)

where *X* and *Y* are the densities of phytoplankton and zooplankton at time τ and position (x, y), respectively. The parameters of the model (1) can be expressed as follows: r_1 represents the inherent growth rate of the phytoplankton without any environment limitations; k_1 is the carrying capacity of the phytoplankton in the presence of predators and harvesting; a_1 represents the consumption rate; b_1 is the half-saturation constant that determines how quickly the maximum is attained when the phytoplankton density increases; e_1 is the conversion factor denoting the number of new herbivores produced for each capture; s_1 is the growth rate of micrograzers due to the alternative prey; d_1 denotes the foodindependent zooplankton mortality rate where $d_1 > s_1$; m_1 is the refuge protect the prey, where $m_1 \in [0, 1)$ is constant [29–32].

For simplicity, using the scaling $X = k_1 N$, $Y = (r_1 k_1 / a_1)P$, and $\tau = t/r_1$, we have

$$\frac{dN}{dt} = N(1-N) - \frac{N}{b+N}P,$$

$$\frac{dP}{dt} = \frac{eN}{b+N}P + sP(1-N) - dP,$$
(2)

where $b = b_1/(1 - m_1)k_1$, $e = a_1e_1/r_1$, $s = s_1/r_1$, and $d = d_1/r_1$. In model (2), we state that the phytoplankton and zooplankton generally experience the same homogeneous environment. Based on a large number of experimental observations in actual environments, it is known that the distribution of individual organisms often depends on interactions with the physical environment and other organisms. A growing body of research indicates that space can change the dynamics of populations and the structure of communities [33]. Thus, we assume that the phytoplankton and zooplankton move randomly, which is described using random Brownian motion [34, 35]. The system can be described using the following set of differential equations:

$$\frac{\partial N}{\partial t} = N (1 - N) - \frac{N}{b + N} P + D_1 \nabla^2 N,$$

$$\frac{\partial P}{\partial t} = \frac{eN}{b + N} P + sP (1 - N) - dP + D_2 \nabla^2 P,$$
(3)

where $\nabla^2 = \partial^2/\partial x^2 + \partial^2/\partial y^2$ is the usual Laplacian operator in two-dimensional space. It is assumed that the environment is uniform and all parameters of the model (3) are independent of space or time, that is, these parameters are constant. Parameters D_1 and D_2 are positive diffusion coefficients for the phytoplankton and zooplankton, respectively. From the viewpoint of biology, we are only interested in the dynamics of models (2) and (3) in the first quadrant R_+^2 . Thus, model (3) is analyzed using the following initial conditions:

$$N_0 = N(x, y \cdot 0) \ge 0,$$

$$P_0 = P(x, y \cdot 0) \ge 0,$$

$$(x, y) \in \Omega = [0, L] \times [0, L]$$
(4)

and the homogeneous Neumann boundary condition:

$$\frac{\partial N}{\partial n} = \frac{\partial P}{\partial n} = 0, \qquad (x, y) \in \partial\Omega,$$
 (5)

which indicates that there are no population fluxes through the boundary and that no external input is imposed from the outside [19]. *n* is the outward unit normal vector for the boundary $\partial\Omega$, which is usually assumed to be smooth [35].

3. Mathematical Analysis

First, we will prove that all solutions of model (3) are bounded.

Theorem 1. Let (N, P) be any solution of model (3). Thus, we can find

$$\lim_{t \to \infty} \sup_{x \in \overline{\Omega}} \max_{x \in \overline{\Omega}} N(x, t) \le 1,$$

$$\lim_{t \to \infty} \sup_{x \in \overline{\Omega}} \max_{x \in \overline{\Omega}} P(x, t) \le b + 1.$$
(6)

Proof. Based on the first equation in model (3), we derive

$$\frac{\partial N}{\partial t} - D_1 \nabla^2 N \le N \left(1 - N \right). \tag{7}$$

Based on the comparison principle, let z(t) be the solution of the equation

$$z'(t) = z(t)(1 - z(t)),$$

$$z(0) = \max_{x \in \overline{\Omega}} N(x, 0).$$
(8)

Thus, we can obtain $N(x,t) \leq z(t)$. For any $\delta > 0$, there is a $T_{\delta} > 0$, such that $z(t) < 1 + \delta$ for any $t \geq T_{\delta}$. Therefore, N(x,t) is defined for all t > 0 and $\lim_{t\to\infty} \sup \max_{x\in\overline{\Omega}} N(x,t) \leq 1$.

Next, we will discuss the boundedness of P(x, t) based on the second equation of model (3). We let $\beta = d - s > 0$ and we find that

$$\frac{\partial P}{\partial t} - D_2 \nabla^2 P \le \frac{eNP}{b+N} - \beta P. \tag{9}$$

Assuming that there is a $T = T(M_0, \varphi) > 0$ such that

$$P(x,t) \ge M_0, \quad \forall t \ge T, \ x \in \overline{\Omega}, \text{ where } M_0 = b+1, \quad (10)$$

we can determine $P/(b+N) \ge 1$, for all $t \ge T^*$, $x \in \overline{\Omega}$, where $T^* = \max \{T_{\delta}, T\}$.

Based on the formulae above and the first equation in model (3), we can obtain

$$\frac{\partial N}{\partial t} - D_1 \nabla^2 N = -N^2 + N \left(1 - \frac{p}{b+N} \right)$$

$$\leq -N^2, \quad \forall t \ge T^*$$
(11)

so we find that

$$0 \le N(x,t) \le z(t) \longrightarrow 0, \quad t \longrightarrow \infty, \ x \in \overline{\Omega},$$
(12)

where z(t) is the solution of the equation $z'(t) = -z^2(t)$ and $z(T^*) = \max_{x \in \overline{\Omega}} N(x, T^*)$.

Then, $N(x,t) \to 0$ as $t \to \infty$, $x \in \overline{\Omega}$. Thus, there is a $t_0 \ge T^*$ such that

$$\frac{eN}{b+N} \le \frac{\beta}{2}, \quad \forall t \ge t_0, \ x \in \overline{\Omega}.$$
(13)

Substituting inequality (13) into the second equation of model (3), we can get

$$\begin{aligned} \frac{\partial P}{\partial t} - D_2 \nabla^2 P &\leq \frac{\beta}{2} P - \beta P \\ &= -\frac{\beta}{2} P, \quad \forall t \geq t_0, \ x \in \overline{\Omega}. \end{aligned}$$
(14)

Based on the comparison principle, we can find that $P(x,t) \rightarrow 0$ as $t \rightarrow \infty$, $x \in \overline{\Omega}$, although this is clearly contradictory.

Let $\Psi(x,t) = P(x,t) - M_0$, t > 0, $x \in \overline{\Omega}$. If there is a $t_0 > 0$, such that $\Psi(x,t) \neq 0$, for all $t > t_0$, $x \in \overline{\Omega}$, then it is considered to be bounded and obviously we can find that $0 < P(x,t) < M_0$ for for all $t > t_0$, $x \in \overline{\Omega}$.

If $\Psi(x,t) = 0$ for a sequence of t_n tending to $+\infty$ as $n \to \infty$, we can say that it is unbounded, that is, the *t*-axis is divided by the zeroes into a sequence of intervals L_n , n > 0. Given this assumption, $\Psi(x,t) \ge 0$, for all $t \in L_n$, $n = 1, 3, 5, \ldots$ For the intervals L_{2n-1} , arguing the inequality (10) is not possible, so we find that $P(x,t) \ge M_0 \exists A \in N^+$ if n > A, for all $t \in L_{2n-1}$, $x \in \overline{\Omega}$, $\partial P/\partial t - D_2 \nabla^2 P \le -(\beta/2)P$.

Based on the comparison principle, $P(x,t) \leq z(t)$ for $t \in L_{2n-1}$, where z(t) is the solution of the equation $z'(t) = -(\beta/2)z(t)$ given suitable initial conditions. Thus,

$$P(x,t) \le z(t) \le z(t_{2n-1}) = M_0,$$

$$n > A, \ \forall t \in L_{2n-1}, \ x \in \overline{\Omega}.$$
(15)

Therefore, we can find $\lim_{t\to\infty} \sup \max_{x\in\overline{\Omega}} P(x,t) \le M_0 = b + 1.$

Among the conditions that satisfy Theorem 1, we mainly address the steady states and their stabilities for (3), as follows.

(i) $E_0 = (0, 0)$, which corresponds to the extinction of the phytoplankton and zooplankton, which is apparently an unstable saddle point.

- (ii) $E_1 = (1,0)$, which corresponds to the extinction of the zooplankton.
- (iii) Their coexistence will be observed in different states $E_2 = (n_2, p_2)$ and $E_3 = (n_3, p_3)$, where

$$n_{2} = \frac{s + e - d - bs - \sqrt{(s + e - d - bs)^{2} + 4bs(s - d)}}{2s},$$
$$n_{3} = \frac{s + e - d - bs + \sqrt{(s + e - d - bs)^{2} + 4bs(s - d)}}{2s}$$
(16)

and $p_i = (n_i + b)(1 - n_i)$ (i = 2.3). Obviously, if s + e - d - bs < 0, n_2 and n_3 are negative. In this study, we assume that s + e - bs > d > s is satisfied. $E_2 = (n_2, p_2)$ and $E_3 = (n_3, p_3)$ have similar properties, thus we focus only on the positive equilibrium point $E_2 = (n_2, p_2)$.

In an actual environment with changing parameters, we will observe different states. In this study, the main aim is to address changes in the state of the equilibrium points E_i (i = 0, 1, 2). Next, we determine the stability of the equilibrium points E_i (i = 0, 1, 2) for model (3). Substituting

$$\begin{pmatrix} N(x, y \cdot t) \\ P(x, y \cdot t) \end{pmatrix} = E_i^T + \begin{pmatrix} U_1(x, y \cdot t) \\ U_2(x, y \cdot t) \end{pmatrix}$$
(17)

into model (3), where $|U_i| \ll 1(i = 1, 2)$, and selecting all terms that are linear about *U*,

$$\frac{\partial U}{\partial t} = D\nabla^2 U + J(E_i)U, \qquad (18)$$

where

$$D = \begin{pmatrix} D_1 & 0 \\ 0 & D_2 \end{pmatrix}, \qquad J(E_i) = \begin{pmatrix} J_{11} & J_{12} \\ J_{21} & J_{22} \end{pmatrix} \Big|_{E_i}, \qquad (19)$$

where

$$J_{11} = 1 - 2N - \frac{bP}{(b+N)^2},$$

$$J_{12} = -\frac{N}{b+N},$$

$$J_{21} = \frac{beP}{(b+N)^2} - sP,$$

$$J_{22} = \frac{eN}{b+N} + s - d - sN.$$
(20)

Substituting (19) into (18), we obtain:

$$\frac{\partial U_1}{\partial t} = D_1 \nabla^2 U_1 + J_{11} U_1 + J_{12} U_2,$$

$$\frac{\partial U_2}{\partial t} = D_2 \nabla^2 U_2 + J_{21} U_1 + J_{22} U_2.$$
(21)

3.1. Stability Analysis

Theorem 2. (*i*) $E_0 = (0, 0)$ *is unstable*;

(*ii*) if d > e/(b+1), then $E_1 = (1, 0)$ for model (3) is asymptotically stable;

(iii) $E_2 = (n_2, p_2)$ is locally asymptotically stable when s + e - bs > d > s and

Proof. The stability of the equilibrium state is determined based on the nature of the eigenvalues in the Jacobian matrix $J(E_i)$. Substituting E_0 into (19), we can determine the eigenvalues of the Jacobian matrix $J(E_0)$, 1 and s - d. Apparently s - d < 0, so E_0 is unstable. The stability of E_1 can be established in a similar manner as E_2 , so we only discuss the stability of E_2 .

We can use λ_i to determine the eigenvalues of the operator $-\Delta$ on Ω , which satisfy the homogeneous Neumann boundary condition and $\cdots > \lambda_n > \cdots > \lambda_4 > \lambda_3 > \lambda_2 > \lambda_1 = 0$. Γ_i is the eigenvector relative to λ_i (*i* = 1, 2, 3, ...), so

$$\lambda_i \Gamma_i + \Delta \Gamma_i = 0 \quad \text{in } \Omega,$$

$$\frac{\partial \Gamma_i}{\partial n} = 0 \quad \text{on } \partial \Omega.$$
 (22)

By substituting $E_2 = (n_2, p_2)$ into (18) and (19), and expanding the solution *U* of (18),

$$U = \sum_{i=1}^{\infty} \tau_i(t) \Gamma_i, \quad \tau_i(t) \in \mathbb{R}^2.$$
(23)

Substituting (23) into (18) and balancing the coefficients about each Γ_i , we can obtain $d\tau_i(t)/dt = T_i\tau_i$ where the matrix T_i satisfies $T_i = J(E_2) - \lambda_i D$.

Thus, $E_2 = (n_2, p_2)$ is stable if and only if each $\tau_i(t)$ decays to zero, that is, there are two eigenvalues with negative real parts for each matrix T_i . Based on simple algebraic computations, the characteristic polynomial of T_i is determined as follow,

$$\gamma^{2} - \operatorname{tr}(T_{i})\gamma + \det(T_{i}) = 0,$$

$$\operatorname{tr}(T_{i}) = J_{11} + J_{22} - \lambda_{i}(D_{1} + D_{2}),$$

$$\det(T_{i}) = \lambda_{i}^{2}D_{1}D_{2} - \lambda_{i}(J_{11}D_{2} + J_{22}D_{1})$$

$$+ J_{11}J_{22} - J_{12}J_{21}.$$
(24)

If tr $(T_i) < 0$ and det $(T_i) > 0$ is satisfied, the above conditions are met. Therefore, with a random $\lambda_i \ge 0$, tr $(T_i) < 0$ and det $(T_i) > 0$ hold if $J_{11} < 0$ and $J_{12}J_{21} < 0$.

Thus, we can obtain:

(a) if
$$b < 1$$
, then $(1 - b)/2 < n_2 < \min(1, \sqrt{be/s} - b)$,

(b) if $b \ge 1$, then $0 \le n_2 < \min(1, \sqrt{be/s} - b)$. This completes the proof.

Theorem 3. If $1 < b < (e - 2s + \sqrt{e^2 - 4es})/2s$, $e \ge 4s$ and (B - s)b > s hold, the steady state $E_2 = (n_2, p_2)$ of the model (3) is globally asymptotically stable, where $B = (en_2)/(b + n_2) + s(1 - n_2)$.

Proof. From La Salle's theorem [36], the Lyapunov function of model (3) needs to be analyzed to prove the theorem. Thus, the following function is structured,

$$V(N,P) = \int_{n_2}^{N} e + \left(d(1-u) - \frac{en_2}{b+n_2} - d(1-n_2) \right) \\ \times \left(1 + \frac{b}{u} \right) du + \int_{p_2}^{P} \frac{u-p_2}{u} du,$$
(25)

assuming $0 < b < (e - 2s + \sqrt{e^2 - 4es})/2s$ and $e \ge 4s$ are satisfied.

$$W(t) = \iint_{\Omega} V(N, P) \, d\Omega \tag{26}$$

V(N, P) is positive for all (N, P) in the positive quadrant, except for $E_2 = (n_2, p_2)$ where $V(n_2, p_2) = 0$. If dW/dt < 0 is proven, then W(t) is the Lyapunov function. Calculating the rate of change of W(t) for the solution of model (3), we find that,

$$\begin{split} \frac{dW}{dt} = &\iint_{\Omega} \left(\left(e + \left(s(1-N) - \frac{en_2}{b+n_2} - s\left(1-n_2\right) \right) \left(1 + \frac{b}{N} \right) \right) \right. \\ &\times \left(N\left(1-N\right) - \frac{NP}{b+N} + D_1 \nabla^2 N \right) + \frac{P-p_2}{P} \\ &\times \left(\frac{eNP}{b+N} + sP\left(1-N\right) - dP + D_2 \nabla^2 P \right) \right) d\Omega \\ &= \iint_{\Omega} \left(\left(e + \left(s(1-N) - \frac{en_2}{b+n_2} - s\left(1-n_2\right) \right) \left(1 + \frac{b}{N} \right) \right) \right. \\ &\times \left(N\left(1-N\right) - \frac{NP}{b+N} \right) + \frac{P-p_2}{P} \\ &\times \left(\frac{eNP}{b+N} + sP\left(1-N\right) - dP \right) \\ &+ \left(e + \left(s\left(1-N\right) - \frac{en_2}{b+n_2} - s\left(1-n_2\right) \right) \left(1 + \frac{b}{N} \right) \right) \right) \\ &+ \frac{P-p_2}{P} D_2 \nabla^2 P \right) d\Omega \\ &= H_1 + H_2. \end{split}$$

Based on simple algebraic computations, H_1 becomes

$$H_{1} = \iint_{\Omega} \left(f\left(N\right) - f\left(n_{2}\right) \right) \left(g\left(N\right) - g\left(n_{2}\right) \right) d\Omega, \quad (28)$$

(27)

where f(x) = ex/(b+x) + s(1-x) and g(x) = (1-x)(b+x).

Obviously, f(x) increases strictly monotonically, so H_1 is negative for all (N, P) in the positive quadrant if and only if

g(x) decreases strictly monotonically, and we find that 1 < $b < (e - 2s + \sqrt{e^2 - 4es})/2s$ and $e \ge 4s$ holds.

In order to determine H_2 , we can use the Green formula. If it satisfies the homogeneous Neumann boundary conditions, we find

$$H_{2} = -\iint_{\Omega} \left(D_{1} |\nabla N|^{2} \frac{d}{dN} \times \left(e + \left(s \left(1 - N\right) - \frac{en_{2}}{b + n_{2}} - s \left(1 - n_{2}\right) \right) \left(1 + \frac{b}{N}\right) \right) + D_{2} |\nabla P|^{2} \frac{d}{dP} \left(\frac{P - p_{2}}{P}\right) d\Omega$$
$$= -\iint_{\Omega} \left(D_{1} |\nabla N|^{2} \left(\frac{(B - s)b}{N^{2}} - s \right) + D_{2} |\nabla P|^{2} \frac{p_{2}}{P^{2}} \right) d\Omega,$$
(29)

where $B = en_2/(b + n_2) + s(1 - n_2)$. Based on Theorem 1, we know 0 < N < 1. If (B - s)b > s is found, $H_2 < 0$. If 1 < b < b $(e-2s+\sqrt{e^2-4es})/2s, e \ge 4s$ and (B-s)b > s hold, so dW/dtis negative. Therefore, $E_2 = (n_2, p_2)$ is globally asymptotically stable.

3.2. Traveling Wave Solutions. In the previous section, we discussed the stability of the equilibrium points E_i (i = 0, 1, 2) for model (3). It can be got that $E_0 = (0,0)$ is unstable, $E_1 = (1,0)$ is unstable for model (3), and the steady state $E_2 = (n_2, p_2)$ of the model (3) is globally asymptotically stable in suitable conditions. However, this is an essential condition that leads to traveling wave solutions. In this section, the main aim is to find a heteroclinic orbit between $E_1 = (1, 0)$ and $E_2 = (n_2, p_2)$ or $E_0 = (0, 0)$ and $E_2 = (n_2, p_2)$.

To find the heteroclinic orbit between $E_1 = (1, 0)$ and $E_2 = (n_2, p_2)$, we need to show that model (3) has a solution with the special form N(x,t) = N(x+ct), P(x,t) = P(x+t)ct), where the wave speed parameter c is positive. After substituting N(x,t) = N(x + ct), P(x,t) = P(x + ct), and s = x + ct into (3), we get the following wave equations

$$cN' = N(1 - N) - \frac{N}{b + N}P + D_1 N'',$$

$$cP' = \frac{eN}{b + N}P + sP(1 - N) - dP + D_2 P''.$$
(30)

If N(s) and P(s) are nonnegative and they satisfy the following boundary conditions

$$N(-\infty) = 1,$$
 $N(+\infty) = n_2,$
 $P(-\infty) = 0,$ $P(+\infty) = p_2,$
(31)

which shows there is a traveling wave solution between E_1 = (1, 0) and $E_2 = (n_2, p_2)$. Substituting u = N' and w = P' into (30),

$$N'=u,$$

$$u' = \frac{c}{D_1}u - \frac{N(1-N)}{D_1} + \frac{NP}{D_1(b+N)},$$
$$P' = w,$$
$$w' = \frac{c}{D_2}w - \frac{eNP}{D_2(b+N)} - \frac{sP(1-N)}{D_2} + \frac{dP}{D_2}.$$
(32)

After computing the Jacobian matrix for model (32) at (1, 0, 0, 0), we get the following matrix:

$$J_{(1,0,0,0)} = \begin{bmatrix} 0 & 1 & 0 & 0\\ \frac{1}{D_1} & \frac{c}{D_1} & \frac{1}{D_1(b+1)} & 0\\ 0 & 0 & 0 & 1\\ 0 & 0 & \frac{d}{D_2} - \frac{1}{D_2(b+1)} & \frac{c}{D_2} \end{bmatrix}.$$
 (33)

Therefore, the characteristic equation of (1, 0, 0, 0) is given by $|\eta E - J_{(1,0,0,0)}| = 0$, and η_i are its eigenvalues, as follows

$$\eta_{1} = \frac{c + \sqrt{c^{2} + 4D_{1}}}{D_{1}}, \qquad \eta_{2} = \frac{c - \sqrt{c^{2} + 4D_{1}}}{D_{1}},$$
$$\eta_{3} = \frac{c + \sqrt{c^{2} - 4(1/(b+1) - d)}}{D_{2}},$$
$$\eta_{4} = \frac{c - \sqrt{c^{2} - 4(1/(b+1) - d)}}{D_{2}}.$$
(34)

If $0 < c < c^* = 2\sqrt{D_2(e/(b+1)-d)}$, then η_3 and η_4 are a pair of complex conjugate eigenvalues with positive real parts. Based on the theorem given in [37], we can find a two-dimensional unstable manifold, which has the base at (1, 0, 0, 0). If $s \rightarrow -\infty$, the trajectory approaching (1, 0, 0, 0)must have P(s) < 0 for some *s*. However, this is contradictory if 0 < P < b + 1.

This discussion shows that because $0 < c < c^* =$ $2\sqrt{D_2(e/(b+1)-d)}$, traveling wave solutions cannot be found for model (32), whereas if $c > 2\sqrt{D_2(e/(b+1)-d)}$, model (32) has non-negative solutions that satisfy the boundary conditions. In some suitable conditions, therefore, there is a traveling wave solution from (1, 0, 0, 0) to $(n_2, 0, p_2, 0)$. Next, we discuss the Jacobian matrix of model (32) at $(n_2, 0, p_2, 0)$,

$$J_{(n_2,0,p_2,0)} = \begin{bmatrix} 0 & 1 & 0 & 0 \\ a_{21} & \frac{c}{D_1} & a_{23} & 0 \\ 0 & 0 & 0 & 1 \\ a_{41} & 0 & 0 & \frac{c}{D_2} \end{bmatrix},$$
(35)

where $a_{21} = -(1 - 2n_2)/D_1 + b(1 - n_2)/D_1(b + n_2), a_{23} =$ $n_2/D_1(b+n_2)$, and $a_{41} = -eb(1-n_2)/D_2(b+n_2) + s(b+n_2)(1-n_2)/D_2(b+n_2)$ $n_2)/D_2$. Thus, the characteristic equation of $(n_2, 0, p_2, 0)$ is given by $|\eta E - J_{(n_2,0,p_2,0)}| = 0$ and η is its eigenvalues, as follows: $T(\eta) = \eta^4 - (c/D_1 + c/D_2)\eta^3 + (c^2/D_1D_2 - a_{21})\eta^2 + (ca_{21}/D_2)\eta - d_{21}(ca_{21}/D_2)\eta^2 + (ca_{21}/D_2)\eta^2 + (ca$ $a_{23}a_{41}$.

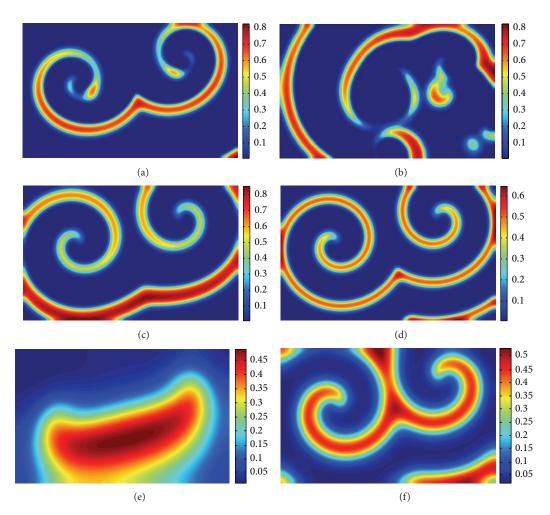


FIGURE 1: Spatiotemporal dynamics of model (3). In ((a), (b)) b = 0.1, ((c), (d)) b = 0.2, and ((e), (f)) b = 0.5. The iteration numbers are 140 in ((a), (c), (e)) and 3000 in ((b), (d), (f)).

Based on the Routh-Hurwitz stability criterion, when $a_{41} < 0$ and $c > \sqrt{D_1 D_2 a_{21}}$, the characteristic equation of $(n_2, 0, p_2, 0)$ always has two eigenvalues with positive real parts and two eigenvalues with negative real parts, that is, to say, there is a stable two-dimensional manifold at $(n_2, 0, p_2, 0)$ and a trajectory from (1, 0, 0, 0) to $(n_2, 0, p_2, 0)$ where $s \rightarrow \infty$. Thus, if $a_{41} < 0$ and $c > \max(2\sqrt{D_2(e/(b+1)-d}), \sqrt{D_1D_2a_{21}})$ hold, there is a traveling wave solution.

4. Numerical Results

In this section, the spatiotemporal dynamics of the phytoplankton-zooplankton model will be simulated, to better understand the theoretical findings. All our numerical simulations use the homogeneous Neumann boundary conditions where (Lx, Ly) = (500, 800). Based on numerical simulations, we found that the spatial pattern distributions of the phytoplankton and zooplankton were always of the same type. This is because the Holling II function reaction term shows that the number of predatory zooplankton is proportional to the number of phytoplankton, while the number of zooplankton is equal to the number of predators. Therefore, we only show the spatial patterns of the phytoplankton and the snapshot with blue parts represents the low density value of phytoplankton N, whereas the snapshot with red parts represents the high density value of phytoplankton N. To investigate the interplay among several critical parameters, we determine the pattern formation using model (3) where other parameters were fixed as d = 0.1, s = 0.01, e = 0.4, $D_1 = 1$, and $D_2 = 0.01$.

We only changed the value of b to test its effects on the spatiotemporal dynamics of the model (3). In Figure 1, we show some snapshots of the numerical simulations where the value of b ranged from 0.1 to 0.5. Figures 1(a) and 1(b) show that the value of b is 0.1, the spatial distribution of prey is a spiral stripy wave, and it is unstable. The instability is uncertain, so we expect that instability can be avoided by increasing the value of b. In Figures 1(c) and 1(d), the value of b is 0.2 and we can observe a periodic spiral wave, because the phytoplankton gather in the form of a continuous spiral stripe to avoid the predator. Finally, in Figures 1(e) and 1(f) the value of b increases to 0.5 and the spatial distribution of prey is a periodic spiral wave, where the width of the stripe increases. Thus, the living space of the prey grows with

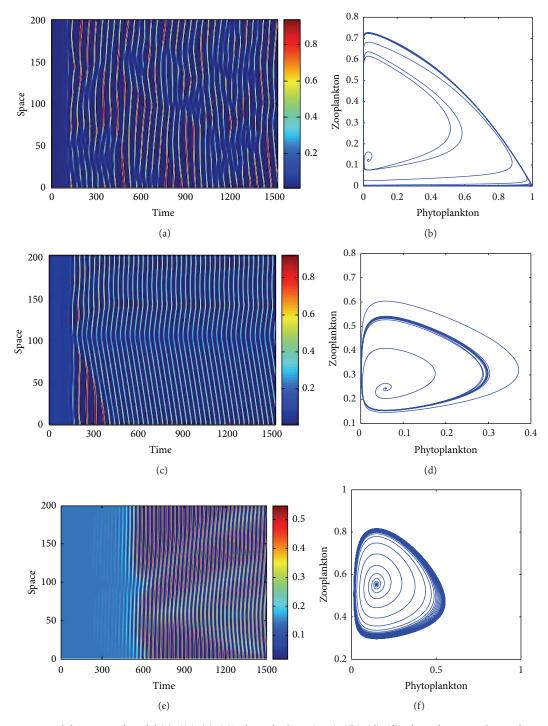


FIGURE 2: Spatiotemporal dynamics of model (3). ((a), (c), (e)) phytoplankton (2-D), ((b), (d), (f)) phase diagram. This analysis used the same set of parameter values as Figure 1.

increasing values of *b*. To better understand the effects of prey refuges, we present the spatiotemporal series and phase diagram in Figure 2, using the same parameter values shown in Figure 1. This series of snapshots also shows that we can increase the number of prey refuges to avoid exacerbating algal growth.

Figure 3 shows the spatial patterns of the phytoplankton when the value of b increases to 1.2. Based on the above

theoretical analysis, we can see that phytoplankton and zooplankton will coexist. With increasing iterations, we can obtain the stable wave pattern shown in the interior of Figure 3(c), which is limited to a rectangular area. Thus, the phytoplankton is limited to this area and its value is almost constant in this area. Figure 3(d) also further that the state of the phytoplankton is changed from its original state to a final stable state.

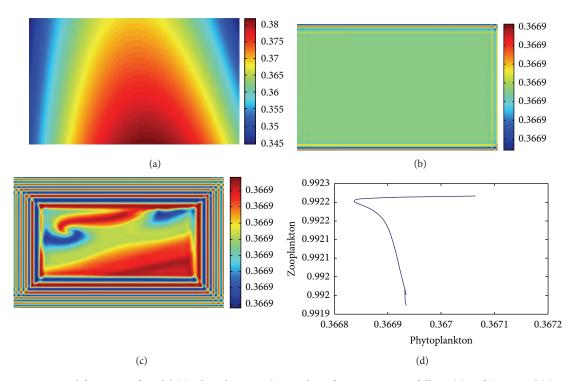


FIGURE 3: Spatiotemporal dynamics of model (3) where b = 1.2. The number of iterations are as follows (a) 1, (b) 200, and (c) 3000. (d) The phase diagram.

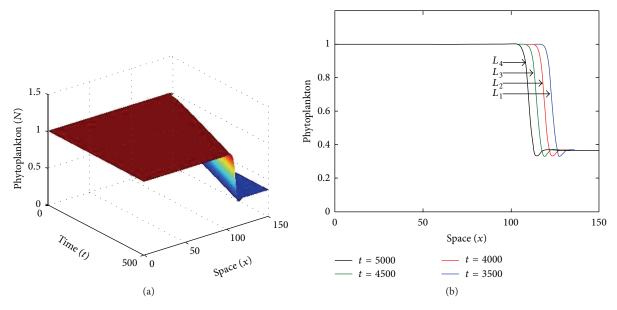


FIGURE 4: Traveling wave solutions for phytoplankton and the change in phytoplankton density at specific times.

Based on the analysis above, we can see that model (3) has traveling wave solutions, which supports all of the results above. If the value of *b* is 1.2, the traveling wave solution of model (3) and the corresponding spatial trend are as shown in Figure 4. It is important to note that the phytoplankton density in Figure 4 changes from $E_1 = (1, 0)$ to $E_2 = (n_2, p_2)$ with spatial variation *x* and time variation *t*, before finally reaching a steady state. We also used different time traveling

waves L_1 , L_2 , L_3 , L_4 at times t = 3500, t = 4000, t = 4500, t = 5000, respectively. Figure 4 shows that the trajectories of the phytoplankton population are different at different times, although all trajectories started at a steady state $E_1 =$ (1,0), before finally moving to a steady state $E_2 = (n_2, p_2)$. Furthermore, it is important to note that model (3) has a traveling wave solution connecting the steady state $E_1 = (1,0)$ and the coexisting equilibrium points $E_2 = (n_2, p_2)$. These results further show that the coexisting equilibrium points $E_2 = (n_2, p_2)$ becomes more stable as the value of *b* increase.

5. Discussion

In this paper, we considered a phytoplankton-zooplankton model given by two coupled reaction-diffusion equations with a Holling's type II functional response and alternative prey and refuge effects with homogeneous Neumann boundary conditions. Mathematical analyses were used to investigate the stability of the equilibrium points and the critical conditions for traveling wave solution. The numerical analysis indicated that parameter b has an important effect on the spatiotemporal dynamics of model (3). Earlier in the article, $m_1 = 1 - b_1/bk_1$. When the carrying capacity k_1 and half-saturation constant b_1 are fixed, the value of b is only determined by the refuge protecting the prey m_1 , which is changed readily by different biological and chemical factors. These result shows that the prey refuge m_1 has an important role in the spatiotemporal dynamics of model (1), where it can make the system more stable thereby stabilizing the density of phytoplankton, which helps to avoid the exacerbation of algal growth. These results may help us to understand the effects of the undoubted susceptibility to diffusion in real ecosystems.

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