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Research article

Schistosomiasis model with treatment, habitat modification and biological control

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Schistosomiasis is a parasitic disease caused by Schistosoma worm infection. Some Abstract: species of snails can serve as the intermediate hosts for the parasite. Numerous interventions have been performed to repress the snail population. One of them is the use of molluscicide. Nevertheless, it is debated that molluscicide intervention has negative impacts on the ecosystem. To investigate the impact of more environmentally friendly interventions, we develop a schistosomiasis model with treatment, habitat modification and biological control. The biological control agent examined in our model is a snail predator. Moreover, to investigate the impact of snail habitat modification, we assume that the snail population grows logistically. We show that all solutions of our model are non-negative and bounded. We also study the existence and stability conditions of equilibrium points. The basic reproduction numbers are determined using the next-generation operator. Linearization combined with the Routh-Hurwitz criterion is used to prove the local stability condition of disease-free equilibrium points. Bifurcation theory is applied to investigate the local stability condition of the endemic equilibrium points. To examine the global behavior of our model, we use asymptotically autonomous system theory and construct a Lyapunov function. We perform several numerical simulations to validate and support our deductive results. Our results show that early treatment can reduce the basic reproduction number and schistosomiasis cases. In addition, modifying snail habitat and releasing the snail predator at the snail habitat can reduce schistosomiasis prevalence. We suggest using snail predators which can hunt and kill snails effectively as a biological control agent.

Keywords: schistosomiasis model; treatment; snail predator; logistic growth; biological control agent; snail habitat modification; asymptotically autonomous systems

1. Introduction

Schistosomiasis is a neglected tropical disease caused by parasitic helminths of the genus *Schistosoma* [1]. Approximately 250 million people are infected with schistosomiasis [2]. Schistosomiasis is commonly found in parts of the Middle East, South America, Southeast Asia and Africa [1]. This disease is mostly found in low-income neighborhoods that lack access to clean water and adequate sanitation [3]. The three main species infecting humans are *Schistosoma haematobium*, *S. japonicum* and *S. mansoni* [4]. *S. japonicum* causes intestinal schistosomiasis and hepatosplenic schistosomiasis in China, The Philippines and Indonesia, while urogenital schistosomiasis in Africa and parts of the Middle East is caused by *S. haematobium* infection. In Africa, the Arabian peninsula and Latin America, *S. mansoni* causes intestinal and liver disease [2]. Colley et al. [5] stated that *S. japonicum* is a zoonotic species, while *S. mansoni* can infect baboons and rodents. On the other hand, *S. haematobium* is declared to be not a zoonotic species because its definitive host is exclusively humans.

As a parasite, the *Schistosoma* worm has an elaborate life cycle. To accomplish its life cycle, *Schistosoma* requires two types of host, namely, intermediate hosts and definitive hosts. Some species of snails can act as intermediate hosts for the parasite. *S. japonicum* used the *Oncomelania* snail as an intermediate host while the *Bulinus* snail could serve as an intermediate host for *S. haematobium*. Meanwhile, *S. mansoni* can infect and make *Biomphalaria* snails intermediate hosts [2]. Humans can serve as a definitive host for the parasite [6]. In the human body, parasites reproduce sexually. The transmission cycle process occurs when the eggs of adult *Schistosoma* worm pairs are excreted through the feces and urine by infected humans. After some time, the eggs that survive in the wild hatch and release miracidia which can infect suitable snails. The parasites reproduce asexually in the snail body [7]. Inside the snail's body, miracidia develop and then produce cercariae which will be released from the snail's body. Cercariae that successfully infect humans can develop into adult worms [2].

Some researchers stated that molluscicide can be used to manage snail populations and schistosomiasis spread [8-10]. This intervention is very effective in decreasing the prevalence of schistosomiasis. However, it is believed that molluscicide may damage the ecosystem [11]. Some researchers recommended the use of snail predators or snail competitors as biological control agents of the snail population [12–14]. Secor [15] stated that several types of fish are predators of host snails. An example is the Cichlid fish in Senegal. In addition, Sokolow et al. [13] reported that prawns in Senegal are predators of host snails, while Mkoji et al. [16] found that the introduction of Red Swamp Crayfish in schistosomiasis endemic areas in Kenya could reduce snail populations and schistosomiasis cases. Different interventions that can be used to control a snail population are snail habitat modification, such as turning farmland to fish ponds, digging and dredging canals, planting of proper trees such as cotton in high-altitude lands or poplar in marshlands. These interventions can reduce snail habitats, which results in a decrease in snail density and survival [17]. Meanwhile, the elimination of snails in some areas in China is achieved after the construction of lake embankments. The construction of the lake embankment prevents flooding which results in a reduced spread of snails around the lake [18]. This suggests that snail habitat modification is related to the environmental carrying capacity of the snail population.

To study the dynamics of schistosomiasis spread, Macdonald [19] proposed a mathematical model.

It is well recognized that the model is the first mathematical model of schistosomiasis. Since then, many researchers have proposed mathematical models which are related to schistosomiasis. Schistosomiasis models that take into account the parasite density in the environment are discussed in [20–23]. In these works, the parasite is divided into two distinct classes, i.e., miracidia and cercariae. Miracidia and cercariae are the parasite stages that can infect snails and humans, respectively. To control vector-borne disease spread, we can use any biological control agents, e.g., predator, competitor or parasites of the vector [24, 25]. A schistosomiasis model considering competitor resistant snails as biological control agents is discussed by Diaby et al. [26]. They found that the introduction of competing snails that are resistant to Schistosoma infection could reduce the population of snails that could serve as intermediate hosts for Schistosoma. In 2021, Nur et al. [27] extended the model proposed in [23] by adding a snail predator population as a biological control agent. Their results showed that the introduction of snail predators in the host snail habitat could reduce the host snail population, cases of schistosomiasis in humans and snails and the number of parasites in the environment. In this paper, we extend the model developed in [27] by adding treated human compartments. In addition, to study the impact of snail habitat modification on the spread of schistosomiasis, we assume that the snail population grows logistically. The rest of the paper is organized as follows. Section 2 describes the model formulation and some of the basic properties of the model. The stability analysis of the equilibrium points is discussed in Section 3. In Section 4, several numerical simulation results are presented. The paper ends with some conclusions in Section 5.

2. Model formulation and basic properties of the model

2.1. Model formulation

In this subsection, we describe the basis and assumptions used in model formulation. We only consider schistosomiasis caused by S. haematobium. Therefore, in our model, only humans are the definitive hosts, because S. haematobium is not zoonotic. We construct the schistosomiasis model based on the life cycle of the Schistosoma worm [6,7,28]. There is a latent period because the parasites need time to migrate through the body, mature and pair up to begin producing eggs. Further, we constructed the model to study the impact of early treatment of exposed humans (humans who have been infected but are still in the prepatent period). Consequently, it is essential to include the exposed human compartment in the model constructed. Thus, we divide the human population into susceptible humans (S_h) , exposed humans (E_h) , infectious humans (I_h) and treated humans (T_h) . People who have been treated can be reinfected; therefore, in Figure 1 which shows the compartment diagram, there is a flow from the treated human to the susceptible human. Anderson et al. [29] stated that the latent period of the snail intermediate host is very dependent on the ambient temperature. High temperatures cause a short latent period. In this article, we assume that the environment has a high temperature so that the latent period in snails is short enough to be negligible. Thus, the snail population is only divided into susceptible snails (S_v) and infectious snails (I_v) . Similar to [20–23], we only consider two stages of Schistosoma worm development, i.e., miracidia and cercariae. Therefore, the parasite is divided into miracidia (M) and cercariae (C). According to the epidemiology of schistosomiasis [7, 28], miracidia can infect snails and cercariae have the ability to infect humans. Hence, susceptible humans and susceptible snails may get infected after adequate contact with cercariae and miracidia, respectively. We assume that snails that can serve as intermediate host is the only food for the predator. It should be noted that *S. haematobium* in the infectious human can produce 100–200 per day per pair. Some of which will be excreted from the infected human body [30]. Moreover, an infectious snail can release about 200 cercariae into the environment for the cases of *S. haematobium* infection [7]. Therefore, the decrease of the parasites in the environment on account of direct interaction with susceptible humans and susceptible snails is neglected. In addition, we assume that there is no recovery for infectious snail and no disease-induced death. Figure 1 shows the transition and interaction between compartments.



Figure 1. Compartment Diagram.

Based on the assumptions and Figure 1, we get the schistosomiasis model as follows:

$$\frac{dS_h}{dt} = \Omega_h - w_1 \beta_{ch} CS_h + \theta_{ts} T_h - d_h S_h$$

$$\frac{dE_h}{dt} = w_1 \beta_{ch} CS_h - w_2 E_h$$

$$\frac{dI_h}{dt} = \theta_{et} E_h - w_3 I_h$$

$$\frac{dT_h}{dt} = \theta_{et} E_h + \theta_{it} I_h - w_4 T_h$$

$$\frac{dS_v}{dt} = \varphi(S_v + I_v) - \beta_{mv} MS_v - \rho S_v(S_v + I_v) - w_5 S_v - \xi PS_v$$

$$\frac{dI_v}{dt} = \beta_{mv} MS_v - w_5 I_v - \rho I_v(S_v + I_v) - \xi PI_v$$

$$\frac{dC}{dt} = \sigma I_v - d_c C$$

$$\frac{dM}{dt} = w_6 I_h - d_m M$$

$$\frac{dP}{dt} = \tau P(S_v + I_v) - d_p P,$$
(2.1)

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where

 $w_1 = 1 - \phi_e \phi$ $w_2 = \theta_{ei} + \theta_{et} + d_h$ $w_3 = \theta_{it} + d_h$ $w_4 = \theta_{ts} + d_h$ $w_5 = d_v + d_r$ $w_6 = \alpha h_h g_h,$

 $\phi_e, \phi \in [0, 1)$ and the other parameters are positive. The definition of all parameters is given in Table 1.

Parameter	Definition
Ω_h	Recruitment rate of humans
ϕ_e	The effectiveness of education implementation
ϕ	Education coverage
eta_{ch}	Cercariae infection rate on humans
$ heta_{it}$	Average waiting time for infectious humans to receive treatment ⁻¹
θ_{et}	Average waiting time for exposed humans to receive treatment ⁻¹
θ_{ts}	Average treatment time ⁻¹
d_h	Natural death rate of humans
θ_{ei}	Latent period ⁻¹
arphi	Birth rate of snails
β_{mv}	Miracidia infection rate on snails
d_v	Natural death rate of snail
d_r	Molluscicide induced death rate of snails
ξ	Predation rate
ho	Competition rate of snail
au	Conversion rate
σ	Cercaria production rate
α	Schistosoma egg hatch rate
h_h	The number of Schistosoma eggs per ml urine
g_h	Average volume of human urine per day
d_p	Natural death rate of snail predator
d_c	Natural death rate of cercariae
d_m	Natural death rate of miracidia

 Table 1. Definitions of model parameters.

2.2. Non-negativity and Boundedness

Theorem 2.1. Solutions of system (2.1) with non-negative initial conditions are non-negative.

Proof. To prove this theorem, we use the method used in [31, 32]. From system (2.1), we have

$$\begin{split} \frac{dS_h}{dt}\Big|_{\zeta(S_h)} &= \Omega_h + \theta_{ts}T_h > 0, \qquad \frac{dE_h}{dt}\Big|_{\zeta(E_h)} = w_1\beta_{ch}CS_h \ge 0, \\ \frac{dI_h}{dt}\Big|_{\zeta(I_h)} &= \theta_{ei}E_h \ge 0, \qquad \frac{dT_h}{dt}\Big|_{\zeta(E_h)} = \theta_{et}E_h + \theta_{it}I_h \ge 0, \\ \frac{dS_v}{dt}\Big|_{\zeta(S_v)} &= \varphi(S_v + I_v) \ge 0, \qquad \frac{dI_v}{dt}\Big|_{\zeta(I_v)} = \beta_{mv}MS_v \ge 0, \\ \frac{dC}{dt}\Big|_{\zeta(C)} &= \sigma I_v \ge 0, \qquad \frac{dM}{dt}\Big|_{\zeta(M)} = w_6I_h \ge 0, \end{split}$$

where $\zeta(x) = \{x = 0 \text{ and } S_h, E_h, I_h, T_h, S_v, I_v, C, M, P \in C(\mathbb{R}_{+0}, \mathbb{R}_{+0})\}$. Based on Lemma 2 presented in [33], the invariant region of system (2.1) is \mathbb{R}^{9}_{+0} . Hence, solutions of system (2.1) with non-negative initial conditions are non-negative.

Theorem 2.2. Solutions of system (2.1) with non-negative initial values are bounded.

Proof. Here, N_h and N_v are the total number of humans and the total number of snails, respectively. It is clear that $N_h = S_h + E_h + I_h + T_h$ and $N_v = S_v + I_v$. From system (2.1), we get

$$\frac{dN_h}{dt} = \Omega_h - d_h N_h,
\frac{dN_v}{dt} = (\varphi - w_5) N_v - \rho N_v^2 - \xi N_v P,
\frac{dP}{dt} = \tau N_v P - d_p P.$$
(2.2)

Using the technique of an integrating factor, we obtain the solution of the first equation of (2.2).

$$N_h(t) = \frac{\Omega_h}{d_h} + \left(N_h(0) - \frac{\Omega_h}{d_h}\right)e^{-d_h t}.$$

It is clear that $0 \le N_h(t) \le \frac{\Omega_h}{d_h}$ for $t \ge 0$ whenever $0 \le N_h(0) \le \frac{\Omega_h}{d_h}$. Thus, N_h is bounded. Now, let $\Pi = N_v + P$. We will show that $\Pi = N_v + P$ is bounded. From the second and third equations of (2.2), we have

$$\begin{aligned} \frac{d\Pi}{dt} &= \frac{dN_v}{dt} + \frac{dP}{dt} \\ &= \left((\varphi - w_5) N_v \left(1 - \frac{\rho N_v}{\varphi - w_5} \right) - \xi N_v P \right) + \left(\tau N_v P - d_p P \right) \\ &= \left(\varphi N_v - \rho N_v^2 - w_5 N_v - \xi N_v P \right) + \left(\tau N_v P - d_p P \right) \\ &\leq \varphi N_v - (\xi - \tau) P N_v - q (N_v + P) \\ &= \varphi \bar{k} - q \Pi, \end{aligned}$$

where $\xi \ge \tau$, $q = \min\{w_5, d_p\}$ and $\bar{k} = \frac{\varphi - w_5}{\rho}$. According to Gronwall's Inequality Lemma, it is found that $\Pi = N_v + P \leq \frac{\varphi \bar{k}}{q}$. Hence, N_v and P are bounded. Now, we use the previous results to show that C and *M* are bounded. Taking into account that $I_h \leq N_h \leq \frac{\Omega_h}{d_h}$ and $I_v \leq P + N_v \leq \frac{\varphi \bar{k}}{q}$, we have

$$\frac{dC}{dt} = \sigma I_v - d_c C \le \sigma \frac{\varphi k}{q} - d_c C,
\frac{dM}{dt} = \alpha h_h g_h I_h - d_m M \le \alpha h_h g_h \frac{\Omega_h}{d_h} - d_m M.$$

According to Gronwall's Inequality Lemma, we get $C \leq \frac{\sigma \varphi \bar{k}}{q d_c}$ and $M \leq \frac{\alpha h_h g_h \Omega_h}{d_h d_m}$. Thus, the solutions of system (2.1) are bounded.

Therefore, we have the following invariant region of system (2.1).

$$\Xi_{+} = \{ (S_h, E_h, I_h, T_h, S_v, I_v, C, M, P) \in \mathbb{R}^9_{+0} | N_h \leq \frac{\Omega_h}{d_h}; N_v + P \leq \frac{\varphi \bar{k}}{d}; C \leq \frac{\sigma \varphi \bar{k}}{q d_c}; M \leq \frac{\alpha h_h g_h \Omega_h}{d_h d_m} \}.$$

The solutions of system (2.1) with initial value in Ξ_+ are always in Ξ_+ .

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2.3. Equilibrium points and basic reproduction number

System (2.1) has five equilibrium points. The first equilibrium point is the snail and predator extinction point $\mathbf{Y}_0 = \left(S_h^{*0}, E_h^{*0}, I_h^{*0}, T_h^{*0}, S_v^{*0}, I_v^{*0}, C^{*0}, M^{*0}, P^{*0}\right) = \left(\frac{\Omega_h}{d_h}, 0, 0, 0, 0, 0, 0, 0, 0, 0\right)$, which always exists in \mathbb{R}^9_{+0} . The second equilibrium point is predator extinction and disease-free point $\mathbf{Y}_0^a = (S_h^{a*}, E_h^{a*}, I_h^{a*}, T_h^{a*}, S_v^{a*}, I_v^{a*}, C^{a*}, M^{a*}, 0) = \left(\frac{\Omega_h}{d_h}, 0, 0, 0, \frac{\varphi - w_5}{\rho}, 0, 0, 0, 0, 0\right)$, which exists in \mathbb{R}^9_{+0} if $\frac{\varphi}{w_5} > 1$. The third equilibrium point is predator extinction-endemic point $\mathbf{Y}_1^a = (S_h^{a**}, E_h^{a**}, I_h^{a**}, T_h^{a**}, S_v^{a**}, I_v^{a**}, C^{a**}, M^{a**}, 0)$, where

$$\begin{split} S_{h}^{a**} &= \frac{w_{4}w_{3}w_{2}d_{c}\Omega_{h}}{(w_{4}w_{3}w_{2}-(\theta_{ts}w_{3}\theta_{et}+\theta_{ts}\theta_{it}\theta_{et}))w_{1}\beta_{ch}\sigma I_{v}^{a**}+d_{h}w_{4}w_{3}w_{2}d_{c}}, \\ E_{h}^{a**} &= \frac{w_{1}\beta_{ch}\sigma w_{4}w_{3}\Omega_{h}I_{v}^{a**}}{(w_{4}w_{3}w_{2}-(\theta_{ts}w_{3}\theta_{et}+\theta_{ts}\theta_{it}\theta_{et}))w_{1}\beta_{ch}\sigma I_{v}^{a**}+d_{h}w_{4}w_{3}w_{2}d_{c}}, \\ I_{h}^{a**} &= \frac{\theta_{ei}w_{1}\beta_{ch}\sigma w_{4}\Omega_{h}I_{v}^{a**}}{(w_{4}w_{3}w_{2}-(\theta_{ts}w_{3}\theta_{et}+\theta_{ts}\theta_{it}\theta_{et}))w_{1}\beta_{ch}\sigma I_{v}^{a**}+d_{h}w_{4}w_{3}w_{2}d_{c}}, \\ T_{h}^{a**} &= \frac{(w_{3}\theta_{et}+\theta_{it}\theta_{ei})w_{1}\beta_{ch}\sigma\Omega_{h}I_{v}^{a**}}{(w_{4}w_{3}w_{2}-(\theta_{ts}w_{3}\theta_{et}+\theta_{ts}\theta_{it}\theta_{et}))w_{1}\beta_{ch}\sigma I_{v}^{a**}+d_{h}w_{4}w_{3}w_{2}d_{c}}, \\ S_{v}^{a**} &= \frac{\varphi_{v}-w_{5}}{\rho} - I_{v}^{a**}, \\ I_{v}^{a**} &= \frac{\varphi_{v}-w_{5}}{w_{1}\beta_{ch}\sigma(\beta_{mv}w_{6}\theta_{ei}\Omega_{h}w_{4}+d_{m}(w_{5}+\rho N_{v}^{a})(w_{3}w_{4}w_{2}-(\theta_{ts}w_{3}\theta_{et}+\theta_{ts}\theta_{it}\theta_{et})))}, \\ C^{a**} &= \frac{\sigma I_{v}^{a**}}{d_{c}}, \\ M^{a**} &= \frac{w_{6}\theta_{ei}w_{1}\beta_{ch}\sigma w_{4}\Omega_{h}I_{v}^{a**}}{d_{m}((w_{4}w_{3}w_{2}-(\theta_{ts}w_{3}\theta_{et}+\theta_{ts}\theta_{it}\theta_{et}))w_{1}\beta_{ch}\sigma I_{v}^{a**}+d_{h}w_{4}w_{3}w_{2}d_{c})}, \\ R_{e}^{a} &= \frac{\beta_{mv}N_{v}^{a}w_{6}\theta_{ei}w_{1}\beta_{ch}\sigma\Omega_{h}}{d_{m}d_{h}d_{c}w_{3}w_{2}(w_{5}+\rho N_{v}^{a})}, \\ N_{v}^{a} &= \frac{\varphi_{-w_{5}}}{\rho}. \end{split}$$

It is easy to see that \mathbf{Y}_{1}^{a} exists in \mathbb{R}_{+0}^{9} if $R_{e}^{a} > 1$ and $\frac{\varphi}{w_{5}} > 1$. The fourth equilibrium point is disease-free point $\mathbf{Y}_{0}^{b} = (S_{h}^{b*}, E_{h}^{b*}, I_{h}^{b*}, T_{h}^{b*}, S_{v}^{b*}, I_{v}^{b*}, C^{b*}, M^{b*}, P^{b*}) = \left(\frac{\Omega_{h}}{d_{h}}, 0, 0, 0, \frac{d_{p}}{\tau}, 0, 0, 0, \frac{\tau(\varphi - w_{5}) - \rho d_{p}}{\xi \tau}\right)$ which exists in \mathbb{R}_{+0}^{9} if $\frac{\tau(\varphi - w_{5})}{\rho d_{p}} > 1$. The last equilibrium point is endemic point $\mathbf{Y}_{1}^{b} = (S_{h}^{b**}, E_{h}^{b**}, I_{h}^{b**}, S_{v}^{b**}, I_{v}^{b**}, C^{b**}, M^{b**}, P^{b*})$, where

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S_h^{b**}	=	$\frac{w_4w_3w_2d_c\Omega_h}{(w_4w_3w_2-(\theta_{ts}w_3\theta_{et}+\theta_{ts}\theta_{it}\theta_{ei}))w_1\beta_{ch}\sigma I_v^{a**}+d_hw_4w_3w_2d_c},$
E_h^{b**}	=	$\frac{w_1\beta_{ch}\sigma w_4w_3\Omega_h I_v^{b**}}{(w_4w_3w_2 - (\theta_{ts}w_3\theta_{et} + \theta_{ts}\theta_{it}\theta_{ei}))w_1\beta_{ch}\sigma I_v^{b**} + d_hw_4w_3w_2d_c},$
I_h^{b**}	=	$\frac{\theta_{ei}w_1\beta_{ch}\sigma w_4\Omega_h I_v^{b**}}{(w_4w_3w_2 - (\theta_{ts}w_3\theta_{et} + \theta_{ts}\theta_{it}\theta_{ei}))w_1\beta_{ch}\sigma I_v^{a**} + d_hw_4w_3w_2d_c},$
T_h^{b**}	=	$\frac{(w_3\theta_{et}+\theta_{it}\theta_{ei})w_1\beta_{ch}\sigma\Omega_h I_v^{b**}}{(w_4w_3w_2-(\theta_{ts}w_3\theta_{et}+\theta_{ts}\theta_{it}\theta_{ei}))w_1\beta_{ch}\sigma I_v^{b**}+d_hw_4w_3w_2d_c},$
S_v^{b**}	=	$\frac{d_p}{\tau} - I_v^{b**},$
I_v^{b**}	=	$\frac{(R_e^b-1)d_hd_md_cw_4w_3w_2(w_5+\rho N_v^b+\xi P^{b*})}{w_1\beta_{ch}\sigma\left(\beta_{mv}w_6\theta_{ei}\Omega_hw_4+d_m(w_5+\rho N_v^b+\xi P^{b*})(w_3w_4w_2-(\theta_{ts}w_3\theta_{et}+\theta_{ts}\theta_{it}\theta_{ei}))\right)},$
C^{b**}	=	$\frac{\sigma I_{\nu}^{b**}}{d_c},$
M^{b**}	=	$\frac{w_{6}\theta_{ei}w_{1}\beta_{ch}\sigma w_{4}\Omega_{h}I_{v}^{b**}}{d_{m}\left((w_{4}w_{3}w_{2}-(\theta_{ls}w_{3}\theta_{el}+\theta_{ls}\theta_{il}\theta_{ei}))w_{1}\beta_{ch}\sigma I_{v}^{a**}+d_{h}w_{4}w_{3}w_{2}d_{c}\right)},$
R_e^b	=	$\frac{\beta_{m\nu}N_{\nu}^{b}w_{6}\theta_{ei}w_{1}\beta_{ch}\sigma\Omega_{h}}{d_{m}d_{h}d_{c}w_{3}w_{2}(w_{5}+\rho N_{\nu}^{b}+\xi P^{b*})},$
N_v^b	=	$\frac{d_p}{\tau}$.

It is clear that \mathbf{Y}_1^b exists in \mathbb{R}_{+0}^9 if $R_e^b > 1$ and $\frac{\tau(\varphi - w_5)}{\rho d_p} > 1$.

In epidemiology, there is an important number recognized as the basic reproduction number. In this paper, we employ the next-generation matrix method described in [34] to investigate the basic reproduction number. First, we establish \mathscr{F} and \mathscr{V} , explained in [34]. \mathscr{F} relates to the new infection terms, while \mathscr{V} represents the transition terms. We consider that the infected classes in our model are E_h , I_h , T_h , I_v , C and M. Consequently, we obtain the following \mathscr{F} and \mathscr{V} .

$$\mathscr{F} = \begin{pmatrix} w_1 \beta_{ch} CS_h \\ 0 \\ 0 \\ \beta_{mv} MS_v \\ 0 \\ 0 \end{pmatrix} \text{ and } \mathscr{V} = \begin{pmatrix} w_2 E_h \\ w_3 I_h - \theta_{ei} E_h \\ w_4 T_h - \theta_{ei} E_h - \theta_{ii} I_h \\ w_5 I_v + \rho I_v (S_v + I_v) + \xi P I_v \\ d_c C - \sigma I_v \\ d_m M - w_6 I_h \end{pmatrix}.$$

We now determine the Jacobian matrix of \mathscr{F} and \mathscr{V} at arbitrary disease-free equilibrium point $(S_h^*, 0, 0, 0, S_v^*, 0, 0, 0, P^*)$ and obtain the following results.

According to [34], the basic reproduction number is the dominant eigenvalue of $F_0V_0^{-1}$.

Obviously, the dominant eigenvalue of $F_0V_0^{-1}$ is given by

$$\rho(F_0 V_0^{-1}) = \sqrt{\frac{w_1 \beta_{ch} S_h^* \sigma \beta_{m\nu} S_\nu^* w_6 \theta_{ei}}{w_2 w_3 d_m d_c (w_5 + \rho S_\nu^* + \xi P^*)}}.$$
(2.3)

Substituting \mathbf{Y}_0^a into (2.3) produces the basic reproduction number when the snail survives but the predator goes to extinction.

$$R_0^a = \sqrt{\frac{\beta_{mv} N_v^a w_6 \theta_{ei} w_1 \beta_{ch} \sigma \Omega_h}{d_m d_h d_c w_3 w_2 (w_5 + \rho N_v^a)}}$$

Substituting \mathbf{Y}_0^b into (2.3) yields the basic reproduction number when the snail and predator survive.

$$R_0^b = \sqrt{\frac{\beta_{mv}N_v^b w_6 \theta_{ei} w_1 \beta_{ch} \sigma \Omega_h}{d_m d_h d_c w_3 w_2 (w_5 + \rho N_v^b + \xi P^{b*})}}.$$

We notice that $R_e^a = (R_0^a)^2$ and $R_e^b = (R_0^b)^2$. Since R_0^a and R_0^b are always positive, it is clear that \mathbf{Y}_1^a exists in \mathbb{R}^9_{+0} if $R_0^a > 1$ and $\frac{\varphi}{w_5} > 1$. Furthermore, \mathbf{Y}_1^b exists in \mathbb{R}^9_{+0} if $R_0^b > 1$ and $\frac{\tau(\varphi - w_5)}{\rho d_p} > 1$.

3. Stability analysis

3.1. Local stability

The general Jacobian matrix of system (2.1) at arbitrary equilibrium point $\mathscr{E} = (S_h^*, E_h^*, I_h^*, T_h^*, S_v^*, I_v^*, C^*, M^*, P^*)$ is given by

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$$\mathbf{J}_{\mathscr{E}} = \begin{pmatrix} -(w_1\beta_{ch}C^* + d_h) & 0 & 0 & \theta_{ts} & 0 & 0 & -w_1\beta_{ch}S_h^* & 0 & 0 \\ w_1\beta_{ch}C^* & -w_2 & 0 & 0 & 0 & 0 & w_1\beta_{ch}S_h^* & 0 & 0 \\ 0 & \theta_{ei} & -w_3 & 0 & 0 & 0 & 0 & 0 \\ 0 & \theta_{et} & \theta_{it} & -w_4 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & A_1 & \varphi - \rho S_v^* & 0 & -\beta_{mv}S_v^* & -\xi S_v^* \\ 0 & 0 & 0 & 0 & \beta_{mv}M^* & -A_2 & 0 & \beta_{mv}S_v^* & -\xi I_v^* \\ 0 & 0 & 0 & 0 & \sigma & -d_c & 0 & 0 \\ 0 & 0 & w_6 & 0 & 0 & 0 & -d_m & 0 \\ 0 & 0 & 0 & 0 & \tau P^* & \tau P^* & 0 & 0 & \tau (S_v^* + I_v^*) - d_p \end{pmatrix},$$

where

$$A_1 = \varphi - \beta_{mv} M^* - (w_5 + \rho(2S_v^* + I_v^*) + \xi P^*), \quad A_2 = w_5 + \rho(S_v^* + 2I_v^*) + \xi P^*$$

First, we present the local stability condition for \mathbf{Y}_0 .

Theorem 3.1. If $\frac{\varphi}{w_5} < 1$, then \mathbf{Y}_0 is locally asymptotically stable. If $\frac{\varphi}{w_5} > 1$, then \mathbf{Y}_0 is unstable. *Proof.* The Jacobian matrix of system (2.1) at \mathbf{Y}_0 is

$$\mathbf{J}_{\mathbf{Y}_{0}} = \begin{pmatrix} -d_{h} & 0 & 0 & \theta_{ts} & 0 & 0 & -w_{1}\beta_{ch}S_{h}^{*0} & 0 & 0 \\ 0 & -w_{2} & 0 & 0 & 0 & w_{1}\beta_{ch}S_{h}^{*0} & 0 & 0 \\ 0 & \theta_{ei} & -w_{3} & 0 & 0 & 0 & 0 & 0 \\ 0 & \theta_{et} & \theta_{it} & -w_{4} & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & \varphi - w_{5} & \varphi & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & -w_{5} & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & \sigma & -d_{c} & 0 & 0 \\ 0 & 0 & w_{6} & 0 & 0 & 0 & 0 & -d_{m} & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & -d_{p} \end{pmatrix}.$$
(3.1)

The eigenvalues of matrix (3.1) are the roots of polynomial (3.2).

$$\mathbf{L}_0(\lambda) = (\lambda + d_p)(\lambda + d_m)(\lambda + d_h)(\lambda + d_c)(\lambda + w_5)(\lambda - (\varphi - w_5))(\lambda + w_2)(\lambda + w_3)(\lambda + w_4).$$
(3.2)

It easy to see that all eigenvalues of matrix (3.1) are negative if $\frac{\varphi}{w_5} < 1$. Moreover, if $\frac{\varphi}{w_5} > 1$, then one eigenvalue of matrix (3.1) is positive. The proof is completed.

Now, we investigate the local stability condition of \mathbf{Y}_0^a . The Jacobian matrix of system (2.1) at \mathbf{Y}_0^a is given as

$$\mathbf{J}_{\mathbf{Y}_{0}^{a}} = \begin{pmatrix} -d_{h} & 0 & 0 & \theta_{ts} & 0 & 0 & -w_{1}\beta_{ch}S_{h}^{a*} & 0 & 0 \\ 0 & -w_{2} & 0 & 0 & 0 & 0 & w_{1}\beta_{ch}S_{h}^{a*} & 0 & 0 \\ 0 & \theta_{ei} & -w_{3} & 0 & 0 & 0 & 0 & 0 \\ 0 & \theta_{ei} & \theta_{it} & -w_{4} & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & w_{5} - \varphi & \varphi - \rho S_{v}^{a*} & 0 & -\beta_{mv}S_{v}^{a*} & -\xi S_{v}^{a*} \\ 0 & 0 & 0 & 0 & 0 & -(w_{5} + \rho S_{v}^{a*}) & 0 & \beta_{mv}S_{v}^{a*} & 0 \\ 0 & 0 & 0 & 0 & 0 & \sigma & -d_{c} & 0 & 0 \\ 0 & 0 & w_{6} & 0 & 0 & 0 & 0 & -d_{m} & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & (\tau S_{v}^{a*} - d_{p}) \end{pmatrix}.$$

$$(3.3)$$

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Therefore, we get the following characteristic polynomial

$$\mathbf{L}(\lambda) = (\lambda - (\tau S_v^{a*} - d_p))(\lambda + d_h)(\lambda - (w_5 - \varphi))(\lambda + w_4)\mathbf{L}_1(\lambda),$$
(3.4)

where $\mathbf{L}_1(\lambda) = \lambda^5 + l_1\lambda^4 + l_2\lambda^3 + l_3\lambda^2 + l_4\lambda + l_5$ and

$$l_{1} = \sum_{i=1}^{5} K_{i}, \qquad l_{4} = \sum_{1 \le i < ... < m}^{5} K_{i}K_{j}K_{k}K_{m},$$

$$l_{2} = \sum_{1 \le i < j}^{5} K_{i}K_{j}, \qquad l_{5} = \left(1 - (R_{0}^{a})^{2}\right)w_{2}w_{3}(w_{5} + \rho S_{v}^{a*})d_{c}d_{m},$$

$$l_{3} = \sum_{1 \le i < j < k}^{5} K_{i}K_{j}K_{k}.$$

Here $K_1 = w_2, K_2 = w_3, K_3 = w_5 + \rho S_v^{a*}, K_4 = d_c$, and $K_5 = d_m$. It is clear that matrix (3.3) has three negative eigenvalues, i.e., $\lambda_1 = -d_h, \lambda_2 = -w_3$, and $\lambda_3 = -(\varphi - w_5)$. Moreover, $\lambda_4 = \tau S_v^{a*} - d_p < 0$ if $\frac{\tau S_v^{a*}}{d_p} = \frac{\tau(\varphi - w_5)}{\rho d_p} < 1$. The other eigenvalues of matrix (3.3) are zeros of $\mathbf{L}_1(\lambda)$. It is easy to see that $l_i > 0$ for i = 1, 2, 3, 4. Further, if $R_0^a < 1$, then $l_5 > 0$. Notice that $l_5 < 0$ if $R_0^a > 1$. Based on Descartes' rule of signs [35], matrix (3.3) has exactly one positive eigenvalue if $R_0^a > 1$. Thus \mathbf{Y}_0^a is unstable if $R_0^a > 1$. We observe that $l_5 = 0$ if $R_0^a = 1$. It indicates that one root of $\mathbf{L}_1(\lambda)$ is zero if $R_0^a = 1$. As a result, if $\frac{\tau S_v^{a*}}{d_p} = \frac{\tau(\varphi - w_5)}{\rho d_p} < 1$ and $R_0^a = 1$ then one eigenvalue of matrix (3.3) is zero. We now use Routh-Hurwitz test [36] to investigate the local stability condition of \mathbf{Y}_0^a . Firstly we establish Routh's array.

Table 2. Routh's array associated to characteristic polynomial $L_1(\lambda)$.

	Column 1	Column 2	Column 3	Column 4
λ^5	1	l_2	l_4	0
λ^4	l_1	l_3	l_5	0
λ^3	$hz_1 = \frac{l_1 l_2 - l_3}{l_1}$	$hz_2 = \frac{l_1 l_4 - l_5}{l_1}$	0	0
λ^2	$hz_3 = \frac{hz_1l_3 - l_1r_2}{hz_1}$	l_5	0	0
λ^1	$hz_4 = \frac{hz_2hz_3 - l_5hz_1}{hz_3}$	0	0	0
λ^0	l_5	0	0	0

Obviously, $l_1 > 0$ always holds. According to the Routh-Hurwitz condition [36, 37], all roots of $\mathbf{L}_1(\lambda)$ have negative real part if all entries in column 1 of Table 2 have the same sign. We observe that $hz_1 > 0$ always holds. Clearly, l_5 is positive if $R_0^a < 1$. Hence, all eigenvalues of matrix (3.3) have negative real parts if $hz_3 > 0$, $hz_4 > 0$, $R_0^a < 1$, and $\frac{\tau(\varphi - w_5)}{d_p \rho} < 1$.

Theorem 3.2. \mathbf{Y}_0^a is locally asymptotically stable if $hz_3 > 0$, $hz_4 > 0$, $R_0^a < 1$, and $\frac{\tau(\varphi - w_5)}{d_p \rho} < 1$. If $R_0^a > 1$ then \mathbf{Y}_0^a is unstable.

The method developed in [38] is used to study the stability condition of \mathbf{Y}_1^a . Similar approach is used in [20, 21]. Firstly, we investigate the conditions so that assumption A1 of Theorem 4.1 in [38] is fulfilled. Let us choose β_{mv} as the bifurcation parameter. We determine the bifurcation point when $R_0^a = 1$ and obtain $\beta_{mv}^{a*} = \frac{d_m d_h d_c w_3 w_2 (w_5 + \rho N_v^a)}{N_v^a w_6 \theta_{ei} w_1 \beta_{ch} \sigma \Omega_h}$. Substituting β_{mv}^{a*} into (3.4) produces the following characteristic

polynomial of $\mathbf{J}_{\mathbf{Y}_{0}^{a}}(\beta_{mv}^{a*})$.

$$\mathbf{L}^{(1a)}(\lambda) = (\lambda - (\tau S_{\nu}^{a*} - d_p))(\lambda + d_h)(\lambda - (w_5 - \varphi))(\lambda + w_4)\lambda \mathbf{L}_1^{(1a)}(\lambda),$$
(3.5)

where $\mathbf{L}_{1}^{(1a)}(\lambda) = \lambda^{4} + l_{1}^{(1a)}\lambda^{3} + l_{2}^{(1a)}\lambda^{2} + l_{3}^{(1a)}\lambda^{1} + l_{4}^{(1a)}$ and

$$l_1^{(1a)} = \sum_{i=1}^5 K_i^{(1a)}, \qquad l_2^{(1a)} = \sum_{1 \le i < j}^5 K_i^{(1a)} K_j^{(1a)}, \\ l_3^{(1a)} = \sum_{1 \le i < j < k}^5 K_i^{(1a)} K_j^{(1a)} K_k^{(1a)}, \qquad l_4^{(1a)} = \sum_{1 \le i < \dots < m}^5 K_i^{(1a)} K_j^{(1a)} K_k^{(1a)} K_m^{(1a)},$$

 $K_1^{(1a)} = w_2, K_2^{(1a)} = w_3, K_3^{(1a)} = w_5 + \rho S_v^{a*}, K_4^{(1a)} = d_c$, and $K_5^{(1a)} = d_m$. Clearly, polynomial (3.5) has three negative roots and a simple zero root. Furthermore, the fourth root is also negative, i.e., $\lambda_4 = \tau S_v^{a*} - d_p < 0$ if $\frac{\tau(\varphi - w_5)}{d_p \rho} < 1$. The other roots of (3.5) are roots of $\mathbf{L}_1^{(1a)}(\lambda)$. The Routh's array associated with $\mathbf{L}_1^{(1a)}(\lambda)$ is shown in Table 3.

	Column 1	Column 2	Column 3
λ^4	1	$l_{2}^{(1a)}$	$l_{\Delta}^{(1a)}$
λ^3	$l_1^{(1a)}$	$l_3^{(1a)}$	0
λ^2	$hr_1 = \frac{l_1^{(1a)} l_2^{(1a)} - l_3^{(1a)}}{l_1^{(1a)}}$	$l_{4}^{(1a)}$	0
λ^1	$hr_2 = \frac{hz_1 l_3^{(1a)} - l_1^{(1a)} l_4^{(1a)}}{hz_1}$	0	0
λ^0	$l_4^{(1a)}$	0	0

Table 3. Routh's array associated to $L_1^{(1a)}(\lambda)$.

Based on Routh-Hurwitz condition [36, 37], all roots of $\mathbf{L}_{1}^{(1a)}(\lambda)$ have negative real part if $hr_2 > 0$, since the other entries in column 1 are always positive. Thus, if $hr_2 > 0$ and $\frac{\tau(\varphi - w_3)}{d_p \rho} < 1$ then $\mathbf{J}_{\mathbf{Y}_0^a}(\beta_{mv}^{a*})$ has one zero eigenvalue while the other eigenvalues have negative real part, which implies that assumption A1 of Theorem 4.1 [38] is met. The right eigenvector of $\mathbf{J}_{\mathbf{Y}_0^a}(\beta_{mv}^{a*})$ corresponding to zero eigenvalue is



where v_8 is arbitrarily positive. Obviously, $v_1 < 0$ and $v_i > 0$ for i = 2, 3, 4, 5, 6, 7. The left eigenvector of $\mathbf{J}_{\mathbf{Y}_a^a}(\beta_{mv}^{a*})$ corresponding to zero eigenvalue is $\vec{k} = (k_1, k_2, k_3, k_4, k_5, k_6, k_7, k_8, k_9)$, where $k_1 = k_4 = k_5 = k_4$

 $\overline{k_{9} = 0, k_{2} = \frac{\theta_{ei}w_{6}\beta_{mv}^{a*}S_{v}^{a*}}{w_{2}w_{3}d_{m}}k_{6}, k_{3} = \frac{w_{6}\beta_{mv}^{a*}S_{v}^{a*}}{w_{3}d_{m}}k_{6}, k_{7} = \frac{w_{1}\beta_{ch}S_{h}^{a*}\theta_{ei}w_{6}\beta_{mv}^{a*}S_{v}^{a*}}{d_{c}w_{2}w_{3}d_{m}}k_{6}, k_{8} = \frac{\beta_{mv}^{a*}S_{v}^{a*}}{d_{m}}k_{6}, \text{ and } k_{6} \text{ is chosen, such that } \vec{k}.\vec{v} = 1. \text{ It is easy to show that } k_{6} > 0. \text{ Hence, } k_{j} > 0 \text{ for } j = 2, 3, 7, 8.$

Set $x_1 = S_h, x_2 = E_h, x_3 = I_h, x_4 = T_h, x_5 = S_v, x_6 = I_v, x_7 = C, x_8 = M, x_9 = P$ and $f_i = \frac{dx_i}{dt}$ for i = 1...9. Now, we calculate *a* and *b*, where

$$a = \sum_{l,i,j=1}^{9} k_l v_i v_j \frac{\partial^2 f_l(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_i \partial x_j} \quad , \quad b = \sum_{l,i=1}^{9} k_l v_i \frac{\partial^2 f_l(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_i \partial \beta_{mv}}.$$

The only non-zero terms of a and b obtained are

$$\begin{split} k_{2}v_{1}v_{6}\frac{\partial^{2}f_{2}(\mathbf{Y}_{0}^{a}\beta_{mv}^{a*})}{\partial x_{1}\partial x_{6}} &= k_{2}v_{1}v_{6}\beta_{ch} < 0, \qquad k_{2}v_{6}v_{1}\frac{\partial^{2}f_{2}(\mathbf{Y}_{0}^{a}\beta_{mv}^{a*})}{\partial x_{6}\partial x_{1}} &= k_{2}v_{6}v_{1}\beta_{ch} < 0, \\ k_{6}v_{5}v_{8}\frac{\partial^{2}f_{6}(\mathbf{Y}_{0}^{a}\beta_{mv}^{a*})}{\partial x_{5}\partial x_{8}} &= v_{5}v_{8}k_{6}\beta_{mv}^{a*} > 0, \qquad k_{6}v_{8}v_{5}\frac{\partial^{2}f_{6}(\mathbf{Y}_{0}^{a}\beta_{mv}^{a*})}{\partial x_{8}\partial x_{5}} &= k_{6}v_{8}v_{5}\beta_{mv}^{a*} > 0, \\ k_{6}v_{5}v_{6}\frac{\partial^{2}f_{6}(\mathbf{Y}_{0}^{a}\beta_{mv}^{a*})}{\partial x_{5}\partial x_{6}} &= -k_{6}v_{5}v_{6}\rho < 0, \qquad k_{6}v_{6}v_{5}\frac{\partial^{2}f_{6}(\mathbf{Y}_{0}^{a}\beta_{mv}^{a*})}{\partial x_{6}\partial x_{5}} &= -k_{6}v_{6}v_{5}\rho < 0, \\ k_{6}v_{6}v_{6}\frac{\partial^{2}f_{6}(\mathbf{Y}_{0}^{a}\beta_{mv}^{a*})}{\partial x_{6}\partial x_{6}} &= -k_{6}v_{6}v_{6}\rho < 0, \qquad k_{6}v_{8}\frac{\partial^{2}f_{6}(\mathbf{Y}_{0}^{a}\beta_{mv}^{a*})}{\partial x_{8}\partial \beta_{mv}} &= k_{6}v_{8}S_{v}^{a*} > 0. \end{split}$$

Since $v_1 < 0$, we have

$$\begin{array}{lll} a & = & k_2 v_1 v_6 \frac{\partial^2 f_2(\mathbf{Y}_0^a \beta_{mv}^a)}{\partial x_1 \partial x_6} + k_2 v_6 v_1 \frac{\partial^2 f_2(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_6 \partial x_1} + k_6 v_5 v_8 \frac{\partial^2 f_6(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_5 \partial x_8} + k_6 v_8 v_5 \frac{\partial^2 f_6(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_8 \partial x_5} + k_6 v_5 v_6 \frac{\partial^2 f_6(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_5 \partial x_6} \\ & & + k_6 v_6 v_5 \frac{\partial^2 f_6(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_6 \partial x_5} + 2k_6 v_6 v_6 \frac{\partial^2 f_6(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_6 \partial x_6} \\ & = & 2 \left(\beta_{ch} k_2 v_1 v_6 - \frac{k_6 v_8^2 \beta_{mv}^{a*} S_v^{a*} (w_5 + S_v^{a*})}{(w_5 + \beta_v^{a*})^2} \right) < 0, \\ b & = & k_6 v_8 \frac{\partial^2 f_6(\mathbf{Y}_0^a \beta_{mv}^{a*})}{\partial x_8 \partial \beta_{mv}} = k_6 v_8 S_v^{a*} > 0. \end{array}$$

Based on Theorem 4.1 in [38], forward bifurcation occurs at $R_0^a = 1$. Hence, the predator extinctionendemic equilibrium point \mathbf{Y}_1^a is locally asymptotically stable if $R_0^a > 1$, $hr_2 > 0$, and $\frac{\tau(\varphi - w_5)}{d_p \rho} < 1$.

Theorem 3.3. \mathbf{Y}_1^a is locally asymptotically stable if $hr_2 > 0$, $\frac{\tau(\varphi - w_5)}{d_p \rho} < 1$, and $R_0^a > 1$ (near 1).

Now, we investigate the stability condition of \mathbf{Y}_0^b . The Jacobian matrix of system (2.1) at \mathbf{Y}_0^b is

$$\mathbf{J}_{\mathbf{Y}_{0}^{b}} = \begin{pmatrix} -d_{h} & 0 & 0 & \theta_{ts} & 0 & 0 & -w_{1}\beta_{ch}S_{h}^{b*} & 0 & 0 \\ 0 & -w_{2} & 0 & 0 & 0 & 0 & w_{1}\beta_{ch}S_{h}^{b*} & 0 & 0 \\ 0 & \theta_{ei} & -w_{3} & 0 & 0 & 0 & 0 & 0 \\ 0 & \theta_{ei} & \theta_{ii} & -w_{4} & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & -\frac{d_{p}\rho}{\tau} & \varphi - \rho S_{v}^{b*} & 0 & -\beta_{mv}S_{v}^{b*} & -\xi S_{v}^{b*} \\ 0 & 0 & 0 & 0 & 0 & -(w_{5} + \rho S_{v}^{b*} + \xi P^{b*}) & 0 & \beta_{mv}S_{v}^{b*} & 0 \\ 0 & 0 & 0 & 0 & \sigma & -d_{c} & 0 & 0 \\ 0 & 0 & w_{6} & 0 & 0 & 0 & 0 & -d_{m} & 0 \\ 0 & 0 & 0 & 0 & \tau P^{b*} & \tau P^{b*} & 0 & 0 & 0 \end{pmatrix}.$$
 (3.6)

The eigenvalues of (3.6) are the zeros of $\mathbf{B}(\lambda)$.

$$\mathbf{B}(\lambda) = (\lambda + d_h)(\lambda + w_4)\mathbf{B}_1(\lambda), \tag{3.7}$$

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where $\mathbf{B}_1(\lambda) = \lambda^7 + b_1\lambda^6 + b_2\lambda^5 + b_3\lambda^4 + b_4\lambda^3 + b_5\lambda^2 + b_6\lambda + b_7$ and

$$\begin{split} b_{1} &= \frac{d_{p}\rho}{\tau} + q_{1}, \\ b_{2} &= \tau P^{b*}\xi S_{v}^{b*} + q_{1}\frac{d_{p}\rho}{\tau} + q_{2}, \\ b_{3} &= \tau P^{b*}\xi S_{v}^{b*}q_{1} + \frac{d_{p}\rho}{\tau}q_{2} + q_{3}, \\ b_{4} &= \tau P^{b*}\xi S_{v}^{b*}q_{2} + \frac{d_{p}\rho}{\tau}q_{3} + q_{4}, \\ b_{5} &= q_{3}\tau P^{b*}\xi S_{v}^{b*} + \frac{d_{p}\rho}{\tau}q_{4} + q_{5} - w_{1}\beta_{ch}S_{h}^{b*}\theta_{ei}w_{6}\sigma\beta_{mv}S_{v}^{b*} \\ &= q_{3}\tau P^{b*}\xi S_{v}^{b*} + \frac{d_{p}\rho}{\tau}q_{4} + \left(1 - \left(R_{0}^{b}\right)^{2}\right)q_{5}, \\ b_{6} &= q_{4}\tau P^{b*}\xi S_{v}^{b*} + q_{5}\frac{d_{p}\rho}{\tau} - w_{1}\beta_{ch}S_{h}^{b*}\theta_{ei}w_{6}\sigma\beta_{mv}S_{v}^{b*}\frac{d_{p}\rho}{\tau} \\ &= q_{4}\tau P^{b*}\xi S_{v}^{b*} + \left(1 - \left(R_{0}^{b}\right)^{2}\right)q_{5}\frac{d_{p}\rho}{\tau}, \\ b_{7} &= q_{5}\tau P^{b*}\xi S_{v}^{b*} - w_{1}\beta_{ch}S_{h}^{b*}\theta_{ei}w_{6}\sigma\beta_{mv}S_{v}^{b*}\tau P^{b*}\xi S_{v}^{b*} \\ &= \left(1 - \left(R_{0}^{b}\right)^{2}\right)q_{5}\tau P^{b*}\xi S_{v}^{b*}, \end{split}$$

$$q_{1} = \sum_{i=1}^{5} K_{i}^{(0b)}, \qquad q_{4} = \sum_{1 \le i ... < m}^{5} K_{i}^{(0b)} K_{j}^{(0b)} K_{m}^{(0b)}, \qquad q_{5} = w_{2} w_{3} d_{m} (w_{5} + \rho S_{v}^{b*} + \xi P^{b*}) d_{c}.$$

$$q_{3} = \sum_{1 \le i < j < k}^{5} K_{i}^{(0b)} K_{j}^{(0b)} K_{k}^{(0b)}, \qquad q_{5} = w_{2} w_{3} d_{m} (w_{5} + \rho S_{v}^{b*} + \xi P^{b*}) d_{c}.$$

 $K_1^{(0b)} = w_2, K_2^{(0b)} = w_3, K_3^{(0b)} = (w_5 + \rho S_v^{b*} + \xi P^{b*}), K_4^{(0b)} = d_c$, and $K_5^{(0b)} = d_m$. Noticeably, polynomial (3.7) has two negative roots, i.e., $\lambda_1 = -d_h$ and $\lambda_2 = -w_4$. The other roots of polynomial (3.7) are zeros of **B**₁(λ). It is easy to see that $b_i > 0$ for i = 1, 2, 3, 4. Additionally, $b_5 > 0, b_6 > 0$ and $b_7 > \text{if } R_0^b < 1$. It is easily recognized that $b_7 = 0$ if $R_0^b = 1$. Hence, if $R_0^b = 1$ then one eigenvalue of matrix (3.6) is zero. Now, we use Routh's Table [36] to analyze the local stability condition of **Y**_0^b.

	Column 1	Column 2	Column 3	Column 4	Column 5
λ^7	1	b_2	b_4	b_6	0
λ^6	b_1	b_3	b_5	b_7	0
λ^5	$hs_1 = \frac{b_1 b_2 - b_3}{h_1}$	$hs_2 = \frac{b_1 b_4 - b_5}{b_1}$	$hs_3 = \frac{b_6b_1 - b_7}{b_1}$	0	0
λ^4	$hs_4 = \frac{hs_1b_3 - b_1hs_2}{hs_1}$	$hs_5 = \frac{hs_1b_5 - b_1hs_3}{hs_1}$	b_7	0	0
λ^3	$hs_6 = \frac{hs_2hs_4 - hs_1hs_5}{hs_4}$	$hs_7 = \frac{hs_3hs_4 - hs_1b_7}{hs_4}$	0	0	0
λ^2	$hs_8 = \frac{hs_5hs_6 - hs_4hs_7}{hs_6}$	b_7	0	0	0
λ^1	$hs_9 = \frac{hs_7hs_8 - hs_6b_7}{hs_8}$	0	0	0	0
λ^0	b_7	0	0	0	0

Table 4. Routh's array associated to $\mathbf{B}_1(\lambda)$.

It is clear that $b_1 > 0$. According to the Routh-Hurwitz criteria [36, 37], all roots of $\mathbf{B}_1(\lambda)$ have negative real parts if all entries in column 1 of Table 4 have the same sign. We recognize that hs_1 is always positive. It is obvious that $b_7 > 0$ if $R_0^b < 1$. Hence, all eigenvalues of (3.6) have negative real

part if $hs_4 > 0$, $hs_6 > 0$, $hs_8 > 0$, $hs_9 > 0$, and $R_0^b < 1$. Notice that $b_7 < 0$ if $R_0^b > 1$. Hence, if $R_0^b > 1$ then \mathbf{Y}_0^b is ustable.

Theorem 3.4. \mathbf{Y}_0^b is locally asymptotically stable if $hs_4 > 0$, $hs_6 > 0$, $hs_8 > 0$, $hs_9 > 0$, and $R_0^b < 1$. If $R_0^b > 1$, then \mathbf{Y}_0^b is unstable.

We utilize center manifold theory [38] to investigate the local stability condition of \mathbf{Y}_{1}^{b} . It is shown that if $R_{0}^{b} = 1$ then matrix (3.6) has one zero eigenvalue. Let us pick β_{mv} as bifurcation parameter. We determine the bifurcation point when $R_{0}^{b} = 1$ and obtain $\beta_{mv}^{b*} = \frac{d_{m}d_{h}d_{c}w_{3}w_{2}(w_{5}+\rho N_{v}^{b}+\xi P^{b*})}{N_{v}^{b}w_{6}\theta_{ei}w_{1}\beta_{ch}\sigma\Omega_{h}}$. Clearly, the characteristic polynomial of $\mathbf{J}_{\mathbf{Y}_{0}^{b}}(\beta_{mv}^{b*})$ is obtained by substituting β_{mv}^{b*} into (3.7). Hence we get (3.8) as a characteristic polynomial of $\mathbf{J}_{\mathbf{Y}_{0}^{b}}(\beta_{mv}^{b*})$.

$$\mathbf{B}^{(1b)}(\lambda) = (\lambda + d_h)(\lambda + w_4)\lambda \mathbf{B}_1^{(1b)}(\lambda), \qquad (3.8)$$

where $\mathbf{B}_{1}^{(1b)}(\lambda) = \lambda^{6} + b_{1}^{(1b)}\lambda^{5} + b_{2}^{(1b)}\lambda^{4} + b_{3}^{(1b)}\lambda^{3} + b_{4}^{(1b)}\lambda^{2} + b_{5}^{(1b)}\lambda + b_{6}^{(1b)}$ and

$$\begin{split} b_1^{(1b)} &= \frac{d_{p\rho}}{\tau} + q_1^{(1b)}, & b_2^{(1b)} &= \tau P^{b*}\xi S_v^{b*} + q_1^{(1b)}\frac{d_{p\rho}}{\tau} + q_2^{(1b)}, \\ b_3^{(1b)} &= \tau P^{b*}\xi S_v^{b*}q_1^{(1b)} + \frac{d_{p\rho}}{\tau}q_2^{(1b)} + q_3^{(1b)}, & b_4^{(1b)} &= \tau P^{b*}\xi S_v^{b*}q_2^{(1b)} + \frac{d_{p\rho}}{\tau}q_3^{(1b)} + q_4^{(1b)}, \\ b_5^{(1b)} &= \tau P^{b*}\xi S_v^{b*}q_3^{(1b)} + \frac{d_{p\rho}}{\tau}q_4^{(1b)}, & b_6^{(1b)} &= \tau P^{b*}\xi S_v^{b*}q_4^{(1b)}, \\ q_1^{(1b)} &= \sum_{i=1}^5 K_i^{(1b)}, & q_2^{(1b)} &= \sum_{1 \le i < j}^5 K_i^{(1b)}K_j^{(1b)}, \\ q_3^{(1b)} &= \sum_{1 \le i < j < k}^5 K_i^{(1b)}K_j^{(1b)}K_k^{(1b)}, & q_4^{(1b)} &= \sum_{1 \le i < \dots < m}^5 K_i^{(1b)}K_j^{(1b)}K_k^{(1b)}K_m^{(1b)}, \end{split}$$

 $K_1^{(1b)} = w_2, K_2^{(1b)} = w_3, K_3^{(1b)} = w_5 + \rho S_v^{b*} + \tau P^{b*}, K_4^{(1b)} = d_c$, and $K_5^{(1b)} = d_m$. Thus, $\mathbf{J}_{\mathbf{Y}_0^b}(\beta_{mv}^{b*})$ has one zero eigenvalue and two negative eigenvalues, i.e., $-d_h$ and $-w_4$. The sign of the other roots of (3.8) is studied by investigating the sign of the roots of polynomial $\mathbf{B}_1^{(1b)}(\lambda)$. Hence, we utilize Routh-Hurwitz test.

	Column 1	Column 2	Column 3	Column 4	Column 5
λ^6	1	$b_2^{(1b)}$	$b_4^{(1b)}$	$b_{6}^{(1b)}$	0
λ^5	$b_1^{(1b)}$	$b_{3}^{(1b)}$	$b_{5}^{(1b)}$	0	0
λ^4	$ht_1 = \frac{b_1^{(1b)}b_2^{(1b)} - b_3^{(1b)}}{b_1^{(1b)}}$	$ht_2 = \frac{b_1^{(1b)}b_4^{(1b)} - b_5^{(1b)}}{b_1^{(1b)}}$	$b_{6}^{(1b)}$	0	0
λ^3	$ht_3 = \frac{ht_1b_3^{(1b)} - b_1^{(1b)}ht_2}{ht_1}$	$ht_4 = \frac{ht_1b_5^{(1b)} - b_1^{(1b)}b_6^{(1b)}}{ht_1}$	0	0	0
λ^2	$ht_5 = \frac{ht_2ht_3 - ht_1ht_4}{ht_3}$	$b_{6}^{(1b)}$	0	0	0
λ^1	$ht_6 = \frac{ht_4ht_5 - ht_3b_6^{(1b)}}{ht_5}$	0	0	0	0
λ^0	$b_{6}^{(1b)}$	0	0	0	0

Table 5. Routh's array associated to $\mathbf{B}_{1}^{(1b)}(\lambda)$.

According to Routh-Hurwitz condition [36, 37], all roots of $\mathbf{B}_1^{(1b)}(\lambda)$ have negative real part if ht_1 , ht_3 , ht_5 and ht_6 are positive, since the other entries in column 1 of Table 5 are positive, i.e., $b_1^{(1b)} > 0$

and $b_6^{(1b)} > 0$. Consequently, $\mathbf{J}_{\mathbf{Y}_0^b}(\boldsymbol{\beta}_{mv}^{b*})$ has a zero eigenvalue and the other eigenvalues have negative real part if $ht_1 > 0$, $ht_3 > 0$, $ht_5 > 0$ and $ht_6 > 0$. Therefore, we now investigate the left and the right eigenvector of $\mathbf{J}_{\mathbf{Y}_0^b}(\boldsymbol{\beta}_{mv}^{b*})$ corresponding to zero eigenvalue. The right eigenvector of $\mathbf{J}_{\mathbf{Y}_0^b}(\boldsymbol{\beta}_{mv}^{b*})$ corresponding to zero eigenvalue is as follows:



where m_8 is arbitrarily positive. It is not difficult to show that $m_1 < 0$, $m_5 < 0$ and $m_i > 0$ for i = 2, 3, 4, 6, 7. The left eigenvector of $\mathbf{J}_{\mathbf{Y}_0^b}(\beta_{mv}^{b*})$ corresponding to zero eigenvalue is $\vec{h} = (h_1, h_2, h_3, h_4, h_5, h_6, h_7, h_8, h_9)$, where $h_1 = h_4 = h_5 = h_9 = 0$, $h_2 = \frac{\theta_{ei}w_6\beta_{mv}^{b*}S_v^{b*}}{w_2w_3d_m}h_6$, $h_3 = \frac{w_6\beta_{mv}^{b*}S_v^{b*}}{w_3d_m}h_6$, $h_7 = \frac{w_1\beta_{ch}S_h^{b*}\theta_{ei}w_6\beta_{mv}^{b*}S_v^{b*}}{d_cw_2w_3d_m}h_6$, $h_8 = \frac{\beta_{mv}^{b*}S_v^{b*}}{d_m}h_6$, and h_6 is chosen, such that $\vec{h}.\vec{m} = 1$. It is easy to show that $h_6 > 0$. Obviously, $h_j > 0$ for j = 2, 3, 7, 8.

Suppose $x_1 = S_h$, $x_2 = E_h$, $x_3 = I_h$, $x_4 = T_h$, $x_5 = S_v$, $x_6 = I_v$, $x_7 = C$, $x_8 = M$, $x_9 = P$ and $f_i = \frac{dx_i}{dt}$ for i = 1...9. Now, we compute *a* and *b*, where

$$a = \sum_{l,i,j=1}^{9} h_l m_i m_j \frac{\partial^2 f_l(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_i \partial x_j} \quad , \quad b = \sum_{l,i=1}^{9} h_l m_i \frac{\partial^2 f_l(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_i \partial \beta_{mv}}.$$

The only non-zero terms of a and b are

$$\begin{split} h_2 m_1 m_6 & \frac{\partial^2 f_2(\mathbf{Y}_0^b \beta_{mv}^{b*})}{\partial x_1 \partial x_6} = h_2 m_1 m_6 \beta_{ch} < 0, \qquad h_2 m_6 m_1 \frac{\partial^2 f_2(\mathbf{Y}_0^b \beta_{mv}^{b*})}{\partial x_6 \partial x_1} = h_2 m_6 m_1 \beta_{ch} < 0, \\ h_6 m_5 m_8 \frac{\partial^2 f_6(\mathbf{Y}_0^b \beta_{mv}^{b*})}{\partial x_5 \partial x_8} = h_6 m_5 m_8 \beta_{mv}^{b*} > 0, \qquad h_6 m_8 m_5 \frac{\partial^2 f_6(\mathbf{Y}_0^b \beta_{mv}^{b*})}{\partial x_8 \partial x_5} = h_6 m_8 m_5 \beta_{mv}^{b*} > 0, \\ h_6 m_5 m_6 \frac{\partial^2 f_6(\mathbf{Y}_0^b \beta_{mv}^{b*})}{\partial x_5 \partial x_6} = -h_6 m_5 m_6 \rho < 0, \qquad h_6 m_8 m_5 \frac{\partial^2 f_6(\mathbf{Y}_0^b \beta_{mv}^{b*})}{\partial x_8 \partial x_5} = -h_6 m_8 m_5 \rho < 0, \\ h_6 m_6 m_6 \frac{\partial^2 f_6(\mathbf{Y}_0^b \beta_{mv}^{b*})}{\partial x_6 \partial x_6} = -h_6 m_6 m_6 \rho < 0, \qquad h_6 m_8 \frac{\partial^2 f_6(\mathbf{Y}_0^b \beta_{mv}^{b*})}{\partial x_8 \partial \beta_{mv}} = h_6 m_8 S_v^{b*} > 0. \end{split}$$

Since $m_1 < 0$, we obtain

$$a = h_2 m_1 m_6 \frac{\partial^2 f_2(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_1 \partial x_6} + h_2 m_6 m_1 \frac{\partial^2 f_2(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_6 \partial x_1} + h_6 m_5 m_8 \frac{\partial^2 f_6(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_5 \partial x_8} + h_6 m_8 m_5 \frac{\partial^2 f_6(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_8 \partial x_5} + h_6 m_5 m_6 \frac{\partial^2 f_6(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_5 \partial x_6} \\ + h_6 m_6 m_5 \frac{\partial^2 f_6(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_6 \partial x_5} + 2 h_6 m_6 m_6 \frac{\partial^2 f_6(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_6 \partial x_6} \\ = 2 \left(\beta_{ch} h_2 m_1 m_6 - \frac{h_6 m_8^2 \beta_{mv}^{b*} S_v^{b*} \beta_{mv}^{b*} (w_5 + \rho S_v^{b*} + \xi P^{b*})^2}{(w_5 + \rho S_v^{b*} + \xi P^{b*})^2} \right) < 0, \\ b = h_6 m_8 \frac{\partial^2 f_6(\mathbf{Y}_0^b, \beta_{mv}^{b*})}{\partial x_8 \partial g_{mv}} = h_6 m_8 S_v^{b*} > 0.$$

Based on Theorem 4.1 in [38], forward bifurcation occurs at $R_0^b = 1$. Hence, the endemic equilibrium point \mathbf{Y}_1^b is locally asymptotically stable if $R_0^b > 1$, $ht_1 > 0$, $ht_3 > 0$, $ht_5 > 0$ and $ht_6 > 0$.

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Theorem 3.5. \mathbf{Y}_1^b is locally asymptotically stable if $ht_1 > 0$, $ht_3 > 0$, $ht_5 > 0$, $ht_6 > 0$ and $R_0^b > 1$ (near 1).

3.2. Global stability

To investigate the global behavior of system (2.1), we use the theory of asymptotically autonomous systems [39, 40], i.e., by studying the limiting systems related to system (2.1). Diaby et al. [41] used this theory to study the dynamics of schistosomiasis model. Since $S_v + I_v = N_v$, system (2.1) is equivalent to the following system.

$$\frac{dS_{h}}{dt} = \Omega_{h} - w_{1}\beta_{ch}CS_{h} + \theta_{ts}T_{h} - d_{h}S_{h}$$

$$\frac{dE_{h}}{dt} = w_{1}\beta_{ch}CS_{h} - w_{2}E_{h}$$

$$\frac{dI_{h}}{dt} = \theta_{ei}E_{h} - w_{3}I_{h}$$

$$\frac{dI_{h}}{dt} = \theta_{ei}E_{h} + \theta_{it}I_{h} - w_{4}T_{h}$$

$$\frac{dI_{v}}{dt} = \beta_{mv}M(N_{v} - I_{v}) - w_{5}I_{v} - \rho I_{v}N_{v} - \xi PI_{v}$$

$$\frac{dC}{dt} = \sigma I_{v} - d_{c}C$$

$$\frac{dM}{dt} = w_{6}I_{h} - d_{m}M$$

$$\frac{dN_{v}}{dt} = (\varphi - w_{5})N_{v} - \rho N_{v}^{2} - \xi N_{v}P,$$

$$\frac{dP}{dt} = \tau PN_{v} - d_{p}P.$$
(3.9)

As mentioned in [42], global stability can be used as the basis for formulating a limiting system. Nakata [43] studied the global dynamics of the total populations, though before constructing a limiting system. Here, we consider studying the global dynamics of snail and snail predator populations before we construct the limiting systems of (3.9), we examine the global behavior of the system formed by the last two equations of (3.9), i.e., system (3.10).

$$\frac{dN_v}{dt} = (\varphi - w_5)N_v - \rho N_v^2 - \xi N_v P,$$

$$\frac{dP}{dt} = \tau P N_v - d_p P.$$
(3.10)

The equilibrium points of system (3.10) are $\mathscr{P}_0 = (0,0)$, which always exists, $\mathscr{P}_1 = (N_v^a, 0) = \left(\frac{\varphi - w_5}{\rho}, 0\right)$ which exists if $\frac{\varphi}{w_5} > 1$, and $\mathscr{P}_1 = (N_v^b, P^{b*}) = \left(\frac{d_p}{\tau}, \frac{\tau(\varphi - w_5) - \rho d_p}{\xi \tau}\right)$, which exists if $\frac{\tau(\varphi - w_5)}{d_p \rho} > 1$. Obviously, \mathscr{P}_0 and \mathscr{P}_2 represent the situation in which both snail and snail predator populations

Obviously, \mathscr{P}_0 and \mathscr{P}_2 represent the situation in which both snail and snail predator populations become extinct and survive, respectively. In addition, the condition in which the snail population survives but the snail predator population becomes extinct is denoted by \mathscr{P}_1 . We now present the stability condition of the equilibrium points of the system (3.10).

Theorem 3.6. (i). \mathscr{P}_0 is global asymptotically stable if $\frac{\varphi}{w_5} < 1$.

(ii). \mathscr{P}_1 is global asymptotically stable if $\frac{\tau(\varphi - w_5)}{d_p \rho} < 1$.

(iii). \mathscr{P}_2 is global asymptotically stable if it exists.

Proof. (i). First, we prove the stability condition of \mathcal{P}_0 . We define the following Lyapunov function.

$$\mathscr{V} = \frac{\tau}{\xi} N_v + P.$$

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The derivative of \mathscr{V} with respect to *t* is given by

$$\begin{aligned} \frac{d\mathcal{V}}{dt} &= \frac{\tau}{\xi} \left((\varphi - w_5) N_v - \rho N_v^2 - \xi N_v P \right) + \left(\tau P N_v - d_p P \right) \\ &= \frac{\tau}{\xi} (\varphi - w_5) N_v - \frac{\tau}{\xi} \rho N_v^2 - d_p P \\ &\leq \frac{\tau}{\xi} (\varphi - w_5) N_v - d_p P. \end{aligned}$$

It is easy to see that $\frac{d\psi}{dt} \le 0$ if $\varphi - w_5 < 0$. Furthermore, $\frac{d\psi}{dt} = 0$ if and only if $N_v = 0$ and P = 0. Thus, \mathscr{P}_0 is global asymptotically stable if $\frac{\varphi}{w_5} < 1$.

(ii). Now, we prove the stability condition of \mathcal{P}_1 . Consider the following Lyapunov function.

$$\mathscr{L} = \frac{\tau}{\xi} \left(N_v - N_v^a - N_v^a \ln \frac{N_v}{N_v^a} \right) + P$$

The time derivative of \mathscr{L} is given by

$$\begin{aligned} \frac{d\mathscr{L}}{dt} &= \frac{\tau}{\xi} \frac{\left(N_{\nu} - N_{\nu}^{a}\right)}{N_{\nu}} \left((\varphi - w_{5})N_{\nu} - \rho N_{\nu}^{2} - \xi N_{\nu}P \right) + \left(\tau P N_{\nu} - d_{p}P \right) \\ &= \frac{\tau}{\xi} \left(N_{\nu} - N_{\nu}^{a}\right) \left(\rho \left(N_{\nu}^{a} - N_{\nu}\right) - \xi P \right) + \left(\tau P N_{\nu} - d_{p}P \right) \\ &= -\frac{\rho \tau}{\xi} \left(N_{\nu} - N_{\nu}^{a}\right)^{2} + d_{p} \left(\frac{\tau N_{\nu}^{a}}{dp} - 1\right) P. \end{aligned}$$

Clearly, $\frac{d\mathscr{L}}{dt} \leq 0$ if $\frac{\tau N_v^a}{d_p} < 1$. Furthermore, $\frac{d\mathscr{L}}{dt} = 0$ if and only if $(N_v, P) = (N_v^a, 0)$. Thus, \mathscr{P}_1 is global asymptotically stable if $\frac{\tau N_v^a}{d_p} = \frac{\tau(\varphi - w_5)}{d_p \rho} < 1$.

(iii). We now prove the stability condition of \mathscr{P}_2 . Consider the following Lyapunov function.

$$\mathscr{M} = \frac{\tau}{\xi} \left(N_{\nu} - N_{\nu}^{b} - N_{\nu}^{b} \ln \frac{N_{\nu}}{N_{\nu}^{b}} \right) + \left(P - P^{b*} - P^{b*} \ln \frac{P}{P^{b*}} \right).$$

The time derivative of \mathcal{M} is given by

$$\frac{d\mathcal{M}}{dt} = \frac{\tau}{\xi} \frac{(N_v - N_v^b)}{N_v} \left((\varphi - w_5) N_v - \rho N_v^2 - \xi N_v P \right) + \frac{(P - P^{b*})}{P} \left(\tau P N_v - d_p P \right)$$

$$= \frac{\tau}{\xi} (N_v - N_v^b) \left((\varphi - w_5) - \rho N_v - \xi P \right) + (P - P^{b*}) \left(\tau N_v - d_p \right).$$

From the equilibrium \mathscr{P}_2 , we have $\tau N_v^b = d_p$ and $\varphi - w_5 = \xi P^{b*} + \frac{\rho d_p}{\tau}$. Hence, we obtain

$$\begin{split} \frac{d\mathcal{M}}{dt} &= \frac{\tau}{\xi} (N_v - N_v^b) \left(\xi P^{b*} + \frac{\rho d_p}{\tau} - \rho N_v - \xi P \right) + (P - P^{b*}) \left(\tau N_v - \tau N_v^b \right) \\ &= \frac{\tau}{\xi} (N_v - N_v^b) \left(\rho \left(N_v^b - N_v \right) - \xi \left(P - P^{b*} \right) \right) + (P - P^{b*}) \tau \left(u N_v - N_v^b \right) \\ &= -\frac{\tau \rho}{\xi} (N_v - N_v^b)^2 \le 0. \end{split}$$

It is clear that $\frac{d\mathcal{M}}{dt} = 0$ if and only if $N_v = N_v^b$. Substituting N_v^b into system (3.10) shows that $P = P^{b*}$. Thus, \mathscr{P}_2 is global asymptotically stable if it exists.

According to [40], we can investigate the dynamics of Systems (2.1) and (3.9) by studying the dynamics of the limiting systems. Based on the global behavior of the equilibrium points of System (3.10), we divide the process into three parts. In this paper, System (3.9) is called an

asymptotically autonomous system with limit Systems (3.11)–(3.13). Theorem 3.6 (i) guarantees that as $t \to \infty$, $(N_v, P) \to (0, 0)$ if $\frac{\varphi}{w_5} < 1$. Notice that $S_v + I_v = N_v = 0$ implies that $S_v = 0$ and $I_v = 0$. Hence, we obtain the following limiting system of (3.9).

$$\frac{dS_{h}}{dt} = \Omega_{h} - w_{1}\beta_{ch}CS_{h} + \theta_{ts}T_{h} - d_{h}S_{h}$$

$$\frac{dE_{h}}{dt} = w_{1}\beta_{ch}CS_{h} - w_{2}E_{h}$$

$$\frac{dI_{h}}{dt} = \theta_{et}E_{h} - w_{3}I_{h}$$

$$\frac{dT_{h}}{dt} = \theta_{et}E_{h} + \theta_{it}I_{h} - w_{4}T_{h}$$

$$\frac{dC_{h}}{dt} = -d_{c}C$$

$$\frac{dM}{dt} = w_{6}I_{h} - d_{m}M.$$
(3.11)

We now investigate the stability condition of the equilibrium point of limiting system (3.11), namely $\mathscr{Y}_0 = \left(S_h^{*0}, E_h^{*0}, I_h^{*0}, T_h^{*0}, C^{*0}, M^{*0}\right) = \left(\frac{\Omega_h}{d_h}, 0, 0, 0, 0, 0\right).$

Theorem 3.7. \mathscr{Y}_0 is global asymptotically stable.

Proof. We define the following Lyapunov function.

$$\mathscr{U} = \left(S_h - S_h^{*0} - S_h^{*0} \ln \frac{S_h}{S_h^{*0}}\right) + E_h + \frac{w_2}{\theta_{ei}}I_h + \frac{w_1\beta_{ch}S_h^{*0}}{d_c}C + \frac{w_3w_2}{\theta_{ei}w_6}M.$$

The derivative of \mathscr{U} with respect to *t* is given as follows:

$$\frac{d\mathscr{U}}{dt} = \left(\frac{S_h - S_h^{*0}}{S_h}\right) (\Omega_h - w_1 \beta_{ch} CS_h + \theta_{ts} T_h - d_h S_h) + (w_1 \beta_{ch} CS_h - w_2 E_h) + \frac{w_2}{\theta_{ei}} (\theta_{ei} E_h - w_3 I_h) + \frac{w_1 \beta_{ch} S_h^{*0}}{d_c} (-d_c C) + \frac{w_3 w_2}{\theta_{ei} w_6} (w_6 I_h - d_m M), = \left(\frac{S_h - S_h^{*0}}{S_h}\right) (\Omega_h + \theta_{ts} T_h - d_h S_h).$$

From the equilibrium point \mathscr{Y}_0 , we have $\Omega_h = d_h S_h^{*0}$. Thus, we obtain

$$\frac{d\mathscr{U}}{dt} = -\frac{d_h}{S_h} \left(S_h - S_h^{*0} \right)^2 + \left(\frac{S_h - S_h^{*0}}{S_h} \right) \theta_{ts} T_h.$$

Since $S_h \leq \frac{\Omega_h}{d_h} = S_h^{*0}$, we conclude that $\frac{d\mathcal{U}}{dt} \leq 0$. Further, $\frac{d\mathcal{U}}{dt} = 0$ if and only if $\frac{d\mathcal{U}}{dt}$ is evaluated at \mathscr{Y}_0 . Therefore, \mathscr{Y}_0 is global asymptotically stable.

Theorem 3.6 (ii) assures that $(N_v, P) \to (N_v^a, 0)$ as $t \to \infty$ if $\frac{\tau(\varphi - w_5)}{d_p \rho} < 1$. Thus, we have the following limiting system of (3.9).

$$\frac{dS_{h}}{dt} = \Omega_{h} - w_{1}\beta_{ch}CS_{h} + \theta_{ts}T_{h} - d_{h}S_{h}$$

$$\frac{dE_{h}}{dt} = w_{1}\beta_{ch}CS_{h} - w_{2}E_{h}$$

$$\frac{dI_{h}}{dt} = \theta_{et}E_{h} - w_{3}I_{h}$$

$$\frac{dT_{h}}{dt} = \theta_{et}E_{h} + \theta_{tt}I_{h} - w_{4}T_{h}$$

$$\frac{dI_{v}}{dt} = \beta_{mv}M(N_{v}^{a} - I_{v}) - w_{5}I_{v} - \rho I_{v}N_{v}^{a}$$

$$\frac{dC}{dt} = \sigma I_{v} - d_{c}C$$

$$\frac{dM}{dt} = w_{6}I_{h} - d_{m}M.$$
(3.12)

We now investigate the global stability condition of the equilibrium point of System (3.12), i.e., $\mathscr{Y}_0^a = (S_h^{a*}, E_h^{a*}, I_h^{a*}, T_h^{a*}, I_v^{a*}, C^{a*}, M^{a*}) = (\frac{\Omega_h}{d_h}, 0, 0, 0, 0, 0, 0).$

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Theorem 3.8. \mathscr{Y}_0^a is global asymptotically stable if $R_0^a \leq 1$.

Proof. Consider the following Lyapunov function:

$$Z = z_1 \left(S_h - S_h^{a*} - S_h^{a*} \ln \frac{S_h}{S_h^{a*}} \right) + z_2 E_h + z_3 I_h + z_4 T_h + z_5 I_v + z_6 C + z_7 M_h$$

where

$$\begin{aligned} z_1 &= z_2 = \frac{d_c(w_5 + \rho N_v^a)}{w_1\beta_{ch}S_h^{a*}\sigma}, & z_3 = (1 - (R_0^a)^2) \left(\frac{(w_5 + \rho N_v^a)w_3w_2\theta_{it}}{w_1\beta_{ch}S_h^{a*}\sigma(w_3\theta_{et} + \theta_{it}\theta_{ei})} + \frac{\beta_{mv}N_v^{a*}w_6}{d_m}\right), \\ z_4 &= (1 - (R_0^a)^2) \frac{(w_5 + \rho N_v^a)w_3w_2}{w_1\beta_{ch}S_h^{a*}\sigma(w_3\theta_{et} + \theta_{it}\theta_{ei})}, & z_5 = 1, \\ z_6 &= \frac{\rho N_v^a + w_5}{\sigma}, & z_7 = \frac{\beta_{mv}N_v^a}{d_m}. \end{aligned}$$

The time derivative of Z is

$$\frac{dZ}{dt} \leq \frac{z_1}{S_h} \left(S_h - S_h^{a*} \right) \left(\Omega_h - d_h S_h \right) + z_1 \left(\frac{S_h - S_h^{a*}}{S_h} \right) \theta_{ts} T_h - z_4 w_4 T_h.$$

From the equilibrium point \mathscr{Y}_0^a , we have $\Omega_h = d_h S_h^{a*}$. Hence, we obtain

$$\frac{dZ}{dt} \leq -\frac{z_1 d_h}{S_h} \left(S_h - S_h^{a*} \right)^2 + z_1 \left(\frac{S_h - S_h^{a*}}{S_h} \right) \theta_{ts} T_h - z_4 w_4 T_h.$$

Certainly, $S_h \leq S_h^{a*}$. Hence, it is easy to see that $\frac{dZ}{dt} \leq 0$ if $R_0^a \leq 1$. Notice that $\frac{dZ}{dt} = 0$ if and only if $S_h = S_h^{a*}$ which implies that $E_h = I_h = T_h = 0$. Hence, the largest invariant set contained in $\{(S_h, E_h, I_h, T_h, I_v, C, M) | \frac{dZ}{dt} = 0\}$ is a singleton set $\{\mathscr{Y}_0^a\}$. Therefore, \mathscr{Y}_0^a is global asymptotically stable if $R_0^a \leq 1$.

Theorem 3.6 (iii) ensures that $(N_{\nu}, P) \to (N_{\nu}^{b}, P^{b*})$ as $t \to \infty$ if $\frac{\tau(\varphi - w_{5})}{d_{p}\rho} > 1$. Thus, we have the following limiting system of (3.9).

$$\frac{dS_{h}}{dt} = \Omega_{h} - w_{1}\beta_{ch}CS_{h} + \theta_{ts}T_{h} - d_{h}S_{h}$$

$$\frac{dE_{h}}{dt} = w_{1}\beta_{ch}CS_{h} - w_{2}E_{h}$$

$$\frac{dI_{h}}{dt} = \theta_{ei}E_{h} - w_{3}I_{h}$$

$$\frac{dI_{h}}{dt} = \theta_{ei}E_{h} + \theta_{it}I_{h} - w_{4}T_{h}$$

$$\frac{dI_{v}}{dt} = \beta_{mv}M\left(N_{v}^{b} - I_{v}\right) - w_{5}I_{v} - \rho I_{v}N_{v}^{b} - \xi I_{v}P^{b*}$$

$$\frac{dC}{dt} = \sigma I_{v} - d_{c}C$$

$$\frac{dM}{dt} = w_{6}I_{h} - d_{m}M.$$
(3.13)

Next, we determine the global stability condition of the equilibrium point of system (3.13), i.e., $\mathscr{Y}_0^b = (S_h^{b*}, E_h^{b*}, I_h^{b*}, T_h^{b*}, I_v^{b*}, C^{b*}, M^{b*}) = (\frac{\Omega_h}{d_h}, 0, 0, 0, 0, 0, 0).$

Theorem 3.9. \mathscr{Y}_0^b is global asymptotically stable if $R_0^b \leq 1$.

Proof. Consider the Lyapunov function as follows:

$$\mathscr{Z} = \mathscr{Z}_1 \left(S_h - S_h^{b*} - S_h^{b*} \ln \frac{S_h}{S_h^{b*}} \right) + \mathscr{Z}_2 E_h + \mathscr{Z}_3 I_h + \mathscr{Z}_4 T_h + \mathscr{Z}_5 I_v + \mathscr{Z}_6 C + z_7 M,$$

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where

$$\begin{aligned} \mathscr{Z}_{1} &= \mathscr{Z}_{2} = \frac{d_{c}(w_{5}+\rho N_{v}^{b}+\xi P^{b*})}{w_{1}\beta_{ch}S_{h}^{b*}\sigma}, & \mathscr{Z}_{3} = (1-(R_{0}^{b})^{2}) \left(\frac{(w_{5}+\rho N_{v}^{b}+\xi P^{b*})w_{3}w_{2}\theta_{it}}{w_{1}\beta_{ch}S_{h}^{b*}\sigma(w_{3}\theta_{et}+\theta_{it}\theta_{ei})} + \frac{\beta_{mv}N_{v}^{b}w_{6}}{d_{m}}\right), \\ \mathscr{Z}_{4} &= (1-(R_{0}^{b})^{2})\frac{(w_{5}+\rho N_{v}^{b}+\xi P^{b*})w_{3}w_{2}}{w_{1}\beta_{ch}S_{h}^{b*}\sigma(w_{3}\theta_{et}+\theta_{it}\theta_{ei})}, & \mathscr{Z}_{5} = 1, \\ \mathscr{Z}_{6} &= \frac{w_{5}+\rho N_{v}^{b}+\xi P^{b*}}{\sigma}, & \mathscr{Z}_{7} = \frac{\beta_{mv}N_{v}^{b}}{d_{m}}. \end{aligned}$$

The time derivative of \mathscr{Z} is

$$\frac{d\mathscr{Z}}{dt} \leq \frac{\mathscr{Z}_1}{S_h} \left(S_h - S_h^{b*} \right) \left(\Omega_h - d_h S_h \right) + \mathscr{Z}_1 \left(\frac{S_h - S_h^{b*}}{S_h} \right) \theta_{ts} T_h - \mathscr{Z}_4 w_4 T_h.$$

From the equilibrium point \mathscr{Y}_0^b , we have $\Omega_h = d_h S_h^{b*}$. Hence, we obtain

$$\frac{d\mathscr{Z}}{dt} \leq -\frac{\mathscr{Z}_1 d_h}{S_h} \left(S_h - S_h^{b*} \right)^2 + \mathscr{Z}_1 \left(\frac{S_h - S_h^{b*}}{S_h} \right) \theta_{ts} T_h - \mathscr{Z}_4 w_4 T_h.$$

Obviously, $S_h \leq S_h^{b*}$. Hence, it is easy to see that $\frac{d\mathscr{X}}{dt} \leq 0$ if $R_0^b \leq 1$. Notice that $\frac{d\mathscr{X}}{dt} = 0$ if and only if $S_h = S_h^{b*}$ which implies that $E_h = I_h = T_h = 0$. Hence, the largest invariant set contained in $\{(S_h, E_h, I_h, T_h, I_v, C, M) | \frac{d\mathscr{X}}{dt} = 0\}$ is a singleton set $\{\mathscr{Y}_0^b\}$. Hence, \mathscr{Y}_0^b is global asymptotically stable if $R_0^b \leq 1$.

Remark \mathscr{Y}_0 of system (3.11) is qualitatively equivalent to the equilibrium \mathbf{Y}_0 of system (2.1) when $\frac{\varphi}{w_5} < 1$. Hence, we conclude that \mathbf{Y}_0 is globally asymptotically stable if $\frac{\varphi}{w_5} < 1$, since \mathscr{Y}_0 is globally asymptotically stable if $\frac{\varphi}{w_5} < 1$. This result suggests that when the snail and the snail predator populations become extinct, provided by $\frac{\varphi}{w_5} < 1$, schistosomiasis will be eradicated.

 \mathscr{Y}_0^a is qualitatively equivalent to the equilibrium \mathbf{Y}_0^a of system (2.1) if $\frac{\tau(\varphi-w_5)}{d_p\rho} < 1$. Thus, \mathbf{Y}_0^a is globally asymptotically stable if $R_0^a \leq 1$ and $\frac{\tau(\varphi-w_5)}{d_p\rho} < 1$, since \mathscr{Y}_0^a of is globally asymptotically stable if $R_0^a \leq 1$. This result indicates that when the snail population survives and the population of snail predators becomes extinct, provided by $\frac{\tau(\varphi-w_5)}{\rho d_p} < 1$, schistosomiasis will be successfully eradicated if $R_0^a \leq 1$.

 \mathscr{Y}_0^b is qualitatively equivalent to the equilibrium point of system (2.1), i.e., \mathbf{Y}_0^b if $\frac{\tau(\varphi - w_5)}{d_p \rho} > 1$. Thus, \mathbf{Y}_0^b is globally asymptotically stable if $R_0^b \leq 1$ and $\frac{\tau(\varphi - w_5)}{d_p \rho} > 1$, since \mathscr{Y}_0^b is globally asymptotically stable if $R_0^b \leq 1$. This result implies that when both snail and snail predator populations survive, provided by $\frac{\tau(\varphi - w_5)}{\rho d_p} > 1$, the disease will die out if $R_0^b \leq 1$.

4. Numerical simulations

We conduct numerical simulations for system (2.1) to validate and support the previous theoretical results. The numerical simulations are performed by using parameter values given in Table 6.

Par.	Definition	Value	Units	Source
Ω_h	Recruitment rate of humans	$\frac{1000 \times 1\%}{365}$	human× day ⁻¹	Assumed
β_{ch}	Cercariae infection rate on humans	1.914×10^{-5}	cercariae ⁻¹ ×day ⁻¹	[22]
ϕ_e	The effectiveness of health education	0.9		Assumed
ϕ	Education coverage	0.9		Assumed
θ_{it}	Average waiting time for infectious humans to receive treatment ⁻¹	$\frac{1}{1\times 4\times 7}$	day ⁻¹	Assumed
θ_{et}	Average waiting time of exposed humans to receive treatment ⁻¹	$\frac{1}{3\times4\times7}$	day ⁻¹	Assumed
θ_{ts}	Average treatment time ⁻¹	1	day ⁻¹	Assumed
d_h	Natural death rate of humans	$\frac{1}{365\times65}$	day ⁻¹	[23]
θ_{ei}	Average latent period ⁻¹	$\frac{1}{6\times7}$	day ⁻¹	[23]
φ	Birth rate of snail	0.4	day ⁻¹	Assumed
d_v	Natural death rate of snail	5.69×10^{-4}	day ⁻¹	[23]
d_r	Molluscicide induced death rate of snail	0.0001	day ⁻¹	Assumed
ξ	Predation rate	0.01	$predator^{-1} \times day^{-1}$	Assumed
ρ	Competition rate of snail	0.01	snail ^{−1} ×day ^{−1}	Assumed
σ	Cercariae production rate	200	$cercariae \times snail^{-1} \times day^{-1}$	[7]
α	Schistosoma egg hatch rate	0.01	miracidia× egg ^{−1}	[22]
h_h	The number of Schistosoma eggs per ml urine	5	$eggs \times ml^{-1}$	Assumed
g_h	The average volume of human urine per day	800	$ml \times human^{-1} \times day^{-1}$	Assumed
d_p	Natural death rate of snail predator	$\frac{1}{2 \times 12 \times 4 \times 7}$	day ⁻¹	Assumed
$\dot{d_c}$	Natural death rate of cercariae	1	day ⁻¹	[22]
d_m	Natural death rate of miracidia	2	day ⁻¹	[22]

Table 6.Parameter values.

4.1. Snail and snail predator go to extinction



Figure 2. (a) Solution curve and (b) phase portrait (I_h, I_v, P) for $\frac{\varphi}{d_v+d_r} < 1$.

We perform numerical simulation using parameter values given in Table 6 and $\tau = 0.00001$. We set $d_r = 0.8$, which indicates the high use of molluscicides. These parameter values yield $\frac{\varphi}{d_v+d_r} = 0.4996 < 1$. Based on Theorem 3.1, \mathbf{Y}_0 is locally asymptotically stable. This result is the same as the numerical simulation result shown in Figure 2.

Figure 2(a) shows that snail and snail predator populations lead to zero. It suggests that both populations go to extinction. Furthermore, the infectious human population tends to zero. It indicates

that the disease will be eradicated. This result is also supported by the phase portrait displayed in Figure 2(b). Notice that all solution trajectories are towards \mathbf{Y}_0 .

Therefore, the use of molluscicides on a large scale has both positive and negative impacts. This kind of intervention can suppress the spread of schistosomiasis, but it can disrupt the environmental balance, e.g., snail predators may become extinct.

4.2. Snail survives but snail predator goes to extinction

Now, we present the second numerical simulation results. The parameter values used are presented in Table 6 with $\tau = 0.00001$, which gives $\frac{\varphi}{d_v+d_r} = 6.9084 \times 10^2 > 1$ and $\frac{\tau(\varphi-w_5)}{\rho d_p} = 0.2684 < 1$. Thus, the snail population survives, but the snail predator goes to extinction. Moreover, we get a bifurcation point for β_{mv} , i.e., $\beta_{mv}^{a*} = 5.6887 \times 10^{-5}$ which corresponds to $R_0^a = 1$ and $hr_2 = 1.8276 > 0$. Using $\beta_{mv} = 5.6887 \times 10^{-6} < \beta_{mv}^{a*}$ implies $R_0^a = 0.3162 < 1$, $hz_3 = 1.8280 > 0$, and $hz_4 = 0.1143 > 0$. Based on Theorem 3.2, \mathbf{Y}_0^a is asymptotically stable. The numerical simulation result is shown in Figure 3(a). If we set $\beta_{mv} = 5.6887 \times 10^{-4} > \beta_{mv}^{a*}$, we get $R_0^a = 3.1626 > 1$. Based on Theorem 3.3, \mathbf{Y}_1^a is asymptotically stable. The numerical simulation result (b). Figure 3(a)



Figure 3. Solution curve for (a) $R_0^a < 1$ and (b) $R_0^a > 1$. Three dimensional phase portrait (I_h, I_v, P) for (c) $\frac{\tau(\varphi - w_5)}{\rho d_p} < 1$, $R_0^a < 1$ and (d) $\frac{\tau(\varphi - w_5)}{\rho d_p} < 1$, $R_0^a > 1$.

shows that the disease will be eradicated. On the other hand, Figure 3(b) shows that the disease will be endemic. Furthermore, the snail population does not tend to zero, while the predator population tends

to zero. These results tell us that we can eradicate schistosomiasis even though the snail population is extant.

Figure 3(c) displays a phase portrait of system (2.1) when $\frac{\varphi}{d_v+d_r} = 6.9084 \times 10^2 > 1$, $\frac{\tau(\varphi-w_5)}{\rho d_p} = 0.2684 < 1$, and $R_0^a = 0.3162 < 1$. It is clear that all solution trajectories tend to the equilibrium point \mathbf{Y}_0^a . Notice that \mathbf{Y}_1^a does not exist. On the other hand, when $R_0^a = 3.1626 > 1$, the equilibrium point \mathbf{Y}_1^a exists, and all solution trajectories go to \mathbf{Y}_1^a as illustrated in Figure 3(d).

4.3. Snail and snail predator survive

Now, we present the third numerical simulation result. The parameter values used are presented in Table 6, and $\tau = 0.0001$, which gives $\frac{\tau(\varphi - w_5)}{\rho d_p} = 2.6841 > 1$, $\beta_{mv}^{b*} = 1.5265 \times 10^{-4}$. Substituting $\beta_{mv} = 1.5265 \times 10^{-5} < \beta_{mv}^{b*}$ gives $R_0^b = 0.3162 < 1$, $hs_4 = 2.2101 > 0$, $hs_6 = 0.4636 > 0$, $hs_8 = 0.0263 >$, and $hs_9 = 0.0003 > 0$. Based on Theorem 3.4, \mathbf{Y}_0^b is asymptotically stable. This result is confirmed by Figure 4(a). If we use $\beta_{mv} = 1.5265 \times 10^{-4} = \beta_{mv}^{b*}$, we get $R_0^b = 1$, $ht_1 = 5.1379 > 0$, $ht_3 = 1.1616 > 0$, $ht_5 = 0.1338 > 0$, and $ht_6 = 0.0071 > 0$. Furthermore, $R_0^b = 3.1622 > 1$ is obtained if we set $\beta_{mv} = 1.5265 \times 10^{-3} > \beta_{mv}^{b*}$. Theorem 3.5 states that \mathbf{Y}_1^b is locally asymptotically stable. This result is similar to the numerical simulation result shown in Figure 4(b). Figure 4(a) shows



Figure 4. Numerical solution for (a) $R_0^b < 1$ and (b) $R_0^b > 1$. Three dimensional phase portrait (I_h, I_v, P) for (c) $\frac{\tau(\varphi - w_5)}{\rho d_p} > 1$, $R_0^b < 1$ and (d) $\frac{\tau(\varphi - w_5)}{\rho d_p} > 1$, $R_0^b > 1$.

that the disease will be eradicated. Nevertheless, Figure 4(b) shows that the disease will be endemic. Furthermore, the snail and its predator population do not tend to zero. These results tell us that we can

eradicate schistosomiasis even though the snail and snail predator populations are extant.

Figure 4(c) illustrates the phase portrait of the system (1) when $\frac{\tau(\varphi-w_5)}{\rho d_p} = 2.6841 > 1$ and $R_0^b = 0.3162 < 1$. We can notice that the equilibrium points \mathbf{Y}_0^a , \mathbf{Y}_0^b , and \mathbf{Y}_1^a exist while \mathbf{Y}_1^b does not exist. It can be seen that all solution trajectories tend to the equilibrium point \mathbf{Y}_0^b . Otherwise, when $R_0^b = 3.1622 > 1$, \mathbf{Y}_0^a , \mathbf{Y}_0^b , \mathbf{Y}_1^a and \mathbf{Y}_1^b exist. However, it can be witnessed, from Figure 4(d), that all solution trajectories go to \mathbf{Y}_1^b .

4.4. Effect of control measures

In this section, we investigate the effect of control measures, i.e., θ_{et} , ρ , τ and ξ , on schistosomiasis spread dynamics. The parameter values used are listed in Table 6 with $\tau = 0.0001$ and $\beta_{mv} = 1.5265 \times 10^{-3}$. After performing numerical simulations, we obtained the following results.



Figure 5. Solution curves with varying (a) θ_{et} , (b) ρ , (c) τ and (d) ξ .

We firstly studied the impact of the average waiting time of a latent human to get treatment on schistosomiasis prevalence. θ_{et} is varied. The results are shown in Figure 5(a). The figure shows that the prevalence of schistosomiasis decreases as the average waiting time $(1/\theta_{et})$ decreases. In addition, the basic reproduction numbers (R_0^b) obtained are 3.3988, 3.1624 and 2.4300 for $\theta_{et} = 0.0071$, $\theta_{et} = 0.0119$ and $\theta_{et} = 0.0357$, respectively. It is clear that the basic reproduction number decreases as θ_{et} increases. Agbata et al. [44] stated that a screening program could increase the possibility of early detection and treatment. Hence, we can increase θ_{et} by implementing a screening program followed by early treatment can

reduce the prevalence of schistosomiasis.

Figure 5(b) shows the effect of habitat modification on the spread of schistosomiasis. We varied ρ while the other parameter values are fixed as given in Table 6, $\tau = 0.0001$ and $\beta_{mv} = 1.5265 \times 10^{-3}$. The basic reproduction numbers (R_0^b) gained is 3.1622 for $\rho = 0.001$, $\rho = 0.01$ and $\rho = 0.1$. It can be seen that the number of infectious human at the beginning of intervention decreases as ρ increases. Here, ρ is the competition rate of the snails. According to [45], habitat change can affect interspecific and intraspecific competition. It is already known that intraspecific competition is competition of members of the same species for limited resources. Thus, snail habitat modification, which may increase the competition rate in the snail population, can reduce the prevalence of schistosomiasis at the beginning of intervention. Xu et al. [17] also stated that an intervention that can be used to control the spread of schistosomiasis is snail habitat modification.

Figure 5(c) shows the effects of the conversion rate of snail predators on the dynamics of infectious humans. We varied τ . The other parameter values are fixed as given in Table 6, and $\beta_{mv} = 1.5265 \times 10^{-3}$. It is seen that the number of infectious humans decreases as τ increases. Furthermore, the basic reproduction values (R_0^b) obtained are 3.1622, 2.5819 and 2.2360 for $\tau = 0.0001$, $\tau = 0.00015$ and $\tau = 0.0002$, respectively. The conversion rate is related to the birth rate. Therefore, the birth rate of snail predators has an important role in reducing schistosomiasis cases. This result is similar to the result given in [27].

Figure 5(d) shows the dynamics of infectious humans with varying ξ . The other parameter values used are given in Table 6 with $\beta_{mv} = 1.5265 \times 10^{-3}$ and $\tau = 0.0001$. The basic reproduction number (R_0^b) obtained is 3.1622 for $\xi = 0.001$, $\xi = 0.005$ and $\xi = 0.01$. The number of infectious humans at the beginning of the outbreak decreases as ξ increases. Here, ξ is predation rate, which is related to the proportion of snails killed per predator per time. Thus, releasing snail predators at snail habitats can reduce the prevalence of schistosomiasis. We recommend using snail predators which can hunt and kill snails effectively, for example, river prawn [13], to maximize the effect of intervention.

5. Conclusions

A schistosomiasis model with treatment, habitat modification and biological control is discussed in this work. Our results show that the basic reproduction number is inversely proportional to θ_{et} . It means that a screening program followed by early treatment can reduce the basic reproduction number and schistosomiasis prevalence. On the other hand, the basic reproduction number is independent of predation rate and competition rate of the snails. However, modifying the snail habitat and releasing snail predators at the snail habitat can reduce the prevalence of schistosomiasis at the beginning of intervention. To maximize this effect, we should use snail predators which can hunt and kill snails effectively.

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Conflict of interest

The authors declare there is no conflict of interest.

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