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Mathematical Biosciences



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# Effects of fish–human transmission and different life stages of fish on Clonorchiasis: A novel mathematical model

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## ARTICLE INFO

**Original Research Article** 

Keywords: Clonorchiasis Fish–human transmission Larval fish Basic reproduction number Global stability Sensitivity analysis

## ABSTRACT

Clonorchiasis is a zoonotic disease mainly caused by eating raw fish and shrimp, and there is no vaccine to prevent it. More than 30 million people are infected worldwide, of which China alone accounts for about half, and is one of the countries most seriously affected by Clonorchiasis. In this work, we formulate a novel Ordinary Differential Equation (ODE) model to discuss the biological attributes of fish within authentic ecosystems and the complex lifecycle of Clonorchis sinensis. This model includes larval fish, adult fish, infected fish, humans, and cercariae. We derive the basic reproduction number and perform a rigorous stability analysis of the proposed model. Numerically, we use data from 2016 to 2021 in Guangxi, China, to discuss outbreaks of Clonorchiasis and obtain the basic reproduction number  $R_0 = 1.4764$ . The fitted curve appropriately reflects the overall trend and replicates a low peak in the case number of Clonorchiasis. By reducing the release rate of cercariae in 2018, the fitted values of Clonorchiasis cases dropped rapidly and almost disappeared. If we decrease the transmission rate from infected fish to humans, Clonorchiasis can be controlled. Our studies also suggest that strengthening publicity education and cleaning water quality can effectively control the transmission of Clonorchiasis in Guangxi, China.

## 1. Introduction

Clonorchiasis is a highly neglected global foodborne disease, with a high incidence in East Asia [1]. Initially, the infection only causes digestive discomfort. As it prolongs or worsens the infection, it may lead to disorders such as biliary tract disease and bile duct lesions [2,3], which can sometimes lead to death. Patients with Clonorchiasis are 4.47 times more likely to develop cholangiocarcinoma than the general population [4]. Nonetheless, the disease has received limited attention in the medical community. In 2010, the World Health Organization (WHO) incorporated it into the category of neglected tropical diseases. Currently, the standard tests used to diagnose Clonorchiasis include hematology, immunology, parasitology, ultrasound, and Computed Tomography (CT) [5]. Detection of liver fluke eggs or specific DNA fragments in stool or bile samples is a definitive diagnostic sign [6]. Eggs can usually be detected in the feces about four weeks after infection [1]. However, even experts have difficulty differentiating the diagnosis of liver fluke eggs from other micro flukes [6,7]. Therefore, it is essential to take precautions, detect early, and seek medical advice.

The hosts of Clonorchis sinensis commonly comprise humans, cats, and dogs [3]. The life cycle of Clonorchis sinensis mainly includes four stages: egg, cercaria, metacercaria, and adult. Every time it enters a new host, it needs a development period to continue transmission. The complex life cycle further exacerbates the complexity of the disease transmission cycle. The eggs enter the water with the feces of the infected individual and are ingested by the first intermediate host (the freshwater snail). After a development period, the eggs become cercaria and escape from the snail into the water. Cercariae in the water encounters a second intermediate host (freshwater fish) and invades the muscle tissue of the fish, where they develop into metacercariae [1,8,9]. The main cause of Clonorchiasis infection is consuming raw fish and shrimp with metacercariae or drinking infected water [10]. In Guangxi and Guangdong, sashimi is a delicacy for guests of honor [4,11]. From the local Chaoshan raw marinade to the raw fish slices in Japanese cuisine, these raw foods are on the table of every household and spread parasitic diseases to more families.

Since the first discovery of Clonorchis sinensis in an ancient corpse in Hubei Province, China, in 1975 [12], scholars in various fields have studied Clonorchiasis from multiple perspectives [1,4,13–18]. In order

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https://doi.org/10.1016/j.mbs.2024.109209

Received 3 December 2023; Received in revised form 5 February 2024; Accepted 12 May 2024 Available online 15 May 2024 0025-5564/© 2024 The Author(s). Published by Elsevier Inc. This is an open access article under the CC BY license (http://creativecommons.org/licenses/by/4.0/). to consider the impact of death and disability with different symptoms on population health, the WHO has initiated a global burden of disease assessment for foodborne diseases, including Clonorchiasis [19] . The burden of disease is often measured by disability-adjusted life vears (DALY) [20]. Sun Yat-sen University and Guangdong Provincial Center for Disease Control and Prevention have estimated the burden of Clonorchiasis in various provinces in China, and the three provinces with the largest DALY are Guangxi Zhuang Autonomous Region, Guangdong Province and Heilongjiang Province, showing a continuous upward trend [21]. This is closely related to the local climate, geography, dietary habits, other factors, and the need for more awareness of the disease and the imperfect disease prevention and control measures [21-23]. Moreover, the infection of Clonorchiasis also has gender and age differences. It was found that men are more likely to be infected with Clonorchiasis than women [1,4,18], which may be related to the fact that men favor raw freshwater fish more than women [24,25]. Qian et al. [1] found that the prevalence of the disease was positively proportional to age, and the prevalence of infection was highest in the age group of 50-59 years old. Qian et al. [26] conducted a cross-sectional survey in two secondary schools in Qiyang County, Hunan Province, They found that children's knowledge of Clonorchiasis is relatively blank, and their families strongly influence their raw food habits. Knowledge education should be strengthened to increase children's alertness to Clonorchiasis.

The development of dynamics based on mathematical modeling has provided a more comprehensive range of ideas for studying infectious diseases [9,27-31]. Dai et al. [30] established an ODE model to study the dynamics of Clonorchiasis transmission. Yuan et al. [29] constructed a model for the transmission of Clonorchiasis in humans, snails, and fishes, then proposed that it would be highly feasible to break the cercariae-fish transmission cycle. Zhang et al. [9] considered the dynamic behavior during the development of Clonorchis sinensis to construct a Partial Differential Equation (PDE) model, and predicted the future development trend of Clonorchiasis in Guangxi through numerical simulations. Mainul et al. [27] proposed four mathematical models to study the dynamics of Clonorchiasis with human treatment and fish vaccination with snail control, demonstrating that fish fry control is an effective control method. Vaccination of fish can largely protect fish from Clonorchis to a great extent and cut off the transmission of the disease from fish to humans. The rapid development of the genomics of Clonorchis sinensis provides a new opportunity for the research and development of vaccines [32]. Fish of different ages may have different behaviors and living conditions in natural ecosystems. Larval fish may be more susceptible to infection by Clonorchis sinensis because they typically live in shallow waters and are more likely to come into contact with environments infected with Clonorchis sinensis eggs [33,34]. Most existing models assumed that the fish was homogeneous, and the stage structure was not considered, which may lead to an overestimation of disease transmission rates. The main cause of human disease is the consumption of sashimi, but the production of sashimi will have some requirements on the weight of the fish, so we assume that only adult fish are involved in the spread of disease. Our subdivision of fish into larval and adult fish can better model these transmission dynamics, help to study the transmission patterns more accurately, and provide a scientific basis for preventing and treating Clonorchiasis.

The paper is structured as follows. Considering the biology of fish in natural ecosystems and the life cycle of Clonorchis sinensis, we divide the fish into two different life stages (larval and adult). In Section 2, we propose and study an ODE model that covers the critical factors of larval fish, adult fish, infected adult fish, humans, and caecilians. The model describes the linked dynamics between the cercaria-fish interaction and the fish-human interaction. We calculate the basic reproduction number of the system and analyze the stability of the disease-free and vector-free equilibrium, and disease-free equilibrium in Section 3. We discuss the stability of the endemic equilibrium in Section 4. In Section 5, we present a case study of the transmission of

Clonorchiasis in Guangxi, China, by numerical studies, and the fitting curve is consistent with the development trend of the actual data. We also perform some sensitivity analysis on  $R_0$  according to the model parameters and observe the change of the fitting curve by changing the values of some parameters. Some concluding remarks are presented in Section 6.

## 2. Model formulation

Clonorchiasis is a multi-host parasitic disease, which increases the difficulty of disease control and poses a great challenge to public health planning. Mathematical modeling has become an effective tool to transform complex systems into mathematical structures, improve understanding of Clonorchiasis, and help establish better long-term effective disease prevention and control systems and rational allocation of available resources.

We establish a mathematical model to describe the transmission dynamics of Clonorchiasis between human and fish hosts, using cercariae as vectors. Fig. 2.1 depicts the transmission of Clonorchiasis between different hosts. We divide the total population  $N_h$  into the following epidemiological categories, reflecting the immune response to infection: susceptible humans  $S_h$  which are free of Clonorchiasis and are at risk of contracting it from cercariae in the environment, infected humans  $I_h$  who have been infected with Clonorchiasis and can shed eggs into the environment, people who have been cured of Clonorchiasis  $R_h$ . The total population is given as

## $N_h = S_h + I_h + R_h.$

G is the concentration of cercariae in water that survived and was infective. We divide the life cycle of fish into larval stage  $L_f$  and adult stage  $N_f$ .

From an epidemiological view, fry control is a vital vector control measure. We consider that infectivity does not affect vector fecundity b and mortality  $\mu_f$ . The natural mortality and maturity rates of larval fish are  $\sigma$  and  $\lambda_f$ , respectively. As noted by [34,35], larval crowding or competition has a general effect on population development. We use  $\alpha$  to denote the density dependence of larval developmental mortality. Based on the modeling idea of the classic Ross-MacDonald model [36], adult fish were divided into susceptible fish  $S_f$  and infected fish  $I_f$ , then we have

 $N_f = S_f + I_f.$ 

Susceptible humans are recruited at a positive constant rate  $\Lambda$ , and  $\mu_h$  is the natural mortality rate. Since infected humans may die from the disease,  $\delta_h$  is set to be the disease-induced mortality rate for humans. The infected population recovers and gains immunity at a rate of  $\gamma_h$ . Susceptible humans are infected by eating fish infected with Clonorchiasis, and we use  $\beta_h$  to represent the transmission rate between susceptible people and fish multiplied by the probability of transmission of infected fish to susceptible people. The infection rate per unit of susceptible population is given by

$$\frac{\beta_h S_h I_f}{N_h}.$$

Infected humans excrete eggs at a rate of  $s_1$ , which enters the first host (freshwater snail) at a rate of  $s_2$  per unit of time, develop and survive in the snail at a rate of  $s_3$ , and are ultimately released into the aquatic environment as cercariae. Thus, we define

 $\lambda_g = s_1 s_2 s_3$ 

as the concentration of cercariae that are produced through the human population, in which these cercariae survive, and are released into the aquatic environment and removed from the water at a rate of  $\mu_g$ .  $\beta_f$  is the transmission rate from cercaria to fish.



Fig. 2.1. Flowchart of the transmission of Clonorchiasis in system (2.1). Solid lines indicate direct transmission between the same species and dashed lines indicate transmission between different species. Different colors represent different meanings: blue for humans, orange for freshwater fish and green for cercariae.

These assumptions together with the schematic diagram for Clonorchiasis transmission (Fig. 2.1) lead to the following ODE model:

$$\begin{cases} \frac{dL_f(t)}{dt} = bN_f - \lambda_f L_f - \sigma L_f - \alpha L_f^2, \\ \frac{dS_f(t)}{dt} = \lambda_f L_f - \beta_f S_f G - (\mu_f + P)S_f, \\ \frac{dI_f(t)}{dt} = \beta_f S_f G - (\mu_f + P)I_f, \\ \frac{dG(t)}{dt} = \lambda_g I_h - \mu_g G, \\ \frac{dS_h(t)}{dt} = \Lambda - \frac{\beta_h S_h I_f}{N_h} - \mu_h S_h, \\ \frac{dI_h(t)}{dt} = \frac{\beta_h S_h I_f}{N_h} - (\mu_h + \delta_h + \gamma_h)I_h, \\ \frac{dR_h(t)}{dt} = \gamma_h I_h - \mu_h R_h. \end{cases}$$

$$(2.1)$$

Using  $N_f = S_f + I_f$ , system (2.1) can be described by the following system:

$$\begin{cases} \frac{dL_f(t)}{dt} = bN_f - \lambda_f L_f - \sigma L_f - \alpha L_f^2, \\ \frac{dN_f(t)}{dt} = \lambda_f L_f - (\mu_f + P)N_f, \\ \frac{dI_f(t)}{dt} = \beta_f (N_f - I_f)G - (\mu_f + P)I_f, \\ \frac{dG(t)}{dt} = \lambda_g I_h - \mu_g G, \\ \frac{dS_h(t)}{dt} = \Lambda - \frac{\beta_h S_h I_f}{N_h} - \mu_h S_h, \\ \frac{dI_h(t)}{dt} = \frac{\beta_h S_h I_f}{N_h} - (\mu_h + \delta_h + \gamma_h)I_h, \\ \frac{dR_h(t)}{dt} = \gamma_h I_h - \mu_h R_h, \end{cases}$$

$$(2.2)$$

with the following initial conditions:

$$\begin{split} &L_f(0) = L_f^0 \geq 0, \ N_f(0) = N_f^0 \geq 0, \ I_f(0) = I_f^0 \geq 0, \ G(0) = G^0 \geq 0, \\ &S_h(0) = S_h^0 \geq 0, \ I_h(0) = I_h^0 \geq 0, \ R_h(0) = R_h^0 \geq 0. \end{split}$$

The detailed biological considerations and experimental values of all parameters are given in Table 1.

**Theorem 2.1.** *System* (2.2) *has a unique and bounded solution with the initial value* 

$$\begin{split} & \left[L_{f}^{0}, N_{f}^{0}, I_{f}^{0}, G^{0}, S_{h}^{0}, I_{h}^{0}, R_{h}^{0}\right] \in K \\ & := \left\{ \left(L_{f}, N_{f}, I_{f}, G, S_{h}, I_{h}, R_{h}\right) \in \mathbb{R}_{+}^{7} : S_{h} + I_{h} + R_{h} > 0, \ I_{f} \leq N_{f} \right\}. \end{split}$$

Moreover, the compact set

$$\begin{split} \Gamma &:= \left\{ (L_f, N_f, I_f, G, S_h, I_h, R_h) \in K : L_f \leq \frac{b\lambda_f}{\alpha(\mu_f + P)}, \\ N_f &\leq \frac{b\lambda_f^2}{\alpha(\mu_f + P)^2}, \, S_h + I_h + R_h \leq \frac{\Lambda}{\mu_h}, \, G \leq \frac{\Lambda\lambda_g}{\mu_h\mu_g} \right\} \end{split}$$

attracts all positive solutions in K.

**Proof.** It follows from [38, Theorem 5.2.1] that system (2.2) admits a unique nonnegative solution  $(L_f(t), N_f(t), I_f(t), G(t), S_h(t), I_h(t), R_h(t))$  through an initial value  $(L_f^0, N_f^0, I_f^0, G^0, S_h^0, I_h^0, R_h^0) \in K$  with the maximum interval of existence  $[0, \iota)$  for  $0 < \iota \leq \infty$ .

Adding the last three equations in system (2.2), we obtain

$$\begin{split} \frac{d(S_h(t)+I_h(t)+R_h(t))}{dt} &= \frac{dN_h(t)}{dt} \\ &= \Lambda - \mu_h N_h - \delta_h I_h \\ &\geq \Lambda - (\mu_h + \delta_h) N_h, \end{split}$$

and thus

$$N_h(t) \ge \frac{\Lambda}{\mu_h + \delta_h} \left( 1 - e^{-(\mu_h + \delta_h)t} \right) + N_h(0)e^{-(\mu_h + \delta_h)t} > 0$$

if  $N_h(0) > 0$  and  $t \in [0, \iota)$ . For  $t \in [0, \iota)$ , we have

$$\begin{split} \Lambda - \left(\mu_h + \delta_h\right) N_h &\leq \frac{dN_h(t)}{dt} \\ &\leq \Lambda - \mu_h N_h. \end{split}$$

Then

$$\begin{split} \frac{\Lambda}{\mu_h + \delta_h} + \left( N_h(0) - \frac{\Lambda}{\mu_h + \delta_h} \right) e^{-(\mu_h + \delta_h)t} &\leq N_h(t) \\ &\leq \frac{\Lambda}{\mu_h} + \left( N_h(0) - \frac{\Lambda}{\mu_h} \right) e^{-\mu_h t}. \end{split}$$

We can see that  $N_h(t)$  is bounded for  $t \in [0, i)$ . Now we introduce

$$\frac{dG(t)}{dt} \leq \lambda_g N_h - \mu_g G.$$

Then according to the comparison principle, G(t) is bounded on [0, i).

Parameter values of system (2.2).						
Symbol	Description		Unit	Value		
b	Birth rate of larval fish	45	year <sup>-1</sup>	Fitting		
$\lambda_f$	Natural maturity rate of adult fish	0.3	year <sup>-1</sup>	Fitting		
σ	Larval fish mortality	0.3	year <sup>-1</sup>	Fitting		
α	Density-dependent development mortality of larval fish	0.0014	year <sup>-1</sup>	Fitting		
$\mu_f + P$	Death rate and predation rate of fish	0.2846	year <sup>-1</sup>	[9]		
$\lambda_{e}$	Rate of release of cercariae into the water	1014	year <sup>-1</sup>	[37]		
$\mu_{g}$	Clearance rate of cercariae in the water	2.607	year <sup>-1</sup>	[9]		
Ň	Recruitment rate of human	2126468	year <sup>-1</sup>	Fitting		
$\beta_H$	Transmission rate from infected fish to human	$4 \times 10^{-6}$	year <sup>-1</sup>	Fitting		
$\mu_h$	Natural mortality rate of human	1/77	year <sup>-1</sup>	[9]		
$\delta_h$	Disease-induced mortality rate of human	0.00505	year <sup>-1</sup>	[21]		
$\gamma_h$	Recovery rate of human	0.73	year <sup>-1</sup>	[9]		
$\beta_f$	Transmission rate from cercaria to fish	$3.59 \times 10^{-10}$	year <sup>-1</sup>	[9]		

By [39, Corollary 3.2], we have that

Table 1

$$\begin{cases} \frac{dV_1(t)}{dt} = bV_2 - \alpha V_1^2, \\ \frac{dV_2(t)}{dt} = \lambda_f V_1 - (\mu_f + P)V_2, \end{cases}$$

exist a globally asymptotically stable equilibrium  $\left(\frac{b\lambda_f}{\alpha(\mu_f+P)}, \frac{b\lambda_f^2}{\alpha(\mu_f+P)^2}\right)$  with respect to all initial values in  $\mathbb{R}^2_+ \setminus \{(0,0)\}$ . By system (2.2), we obtain

$$\begin{cases} \frac{dL_f(t)}{dt} \leq bN_f - \alpha L_f^2, \\ \frac{dN_f(t)}{dt} = \lambda_f L_f - (\mu_f + P)N_f \end{cases}$$

According to the comparison principle, there exist  $M_1$  and  $M_2$  such that

$$L_f(t) \le M_1, \ N_f(t) \le M_2, \ \forall t \in [0, i).$$

Thus, we see that  $i = \infty$  and the solution of system (2.2) exists globally. From the previous arguments, we can see that

$$\lim_{t\to\infty}\sup(L_f(t),N_f(t))\leq\left(\frac{b\lambda_f}{\alpha(\mu_f+P)},\frac{b\lambda_f^2}{\alpha(\mu_f+P)^2}\right),$$

which completes the proof.

From system (2.2), we have the following system:

$$\begin{cases} \frac{dL_f(t)}{dt} = bN_f - \lambda_f L_f - \sigma L_f - \alpha L_f^2, \\ \frac{dN_f(t)}{dt} = \lambda_f L_f - (\mu_f + P)N_f. \end{cases}$$
(2.3)

By [31,40], we define the vector reproduction number as

$$R_v = \frac{b\lambda_f}{(\sigma + \lambda_f)(\mu_f + P)}.$$

System (2.3) has always one trivial equilibrium (0, 0). The positive equilibrium  $(L_f^*, N_f^*)$  of system (2.3) exists when  $R_v > 1$ , where

$$(L_f^*, N_f^*) = \left(\frac{\lambda_f + \sigma}{\alpha}(R_v - 1), \frac{\lambda_f(\lambda_f + \sigma)}{\alpha(\mu_f + P)}(R_v - 1)\right).$$
(2.4)

From [31, Lemma 2.1], we have the following result.

## Lemma 2.2. The following statements are valid:

- (i) If  $R_v \leq 1$ , the trivial equilibrium (0,0) of system (2.3) is globally asymptotically stable in  $\mathbb{R}^2_+$ ;
- (ii) If R<sub>v</sub> > 1, the positive equilibrium (L<sup>\*</sup><sub>f</sub>, N<sup>\*</sup><sub>f</sub>) of system (2.3) is globally asymptotically stable in ℝ<sup>2</sup><sub>+</sub> \ {(0,0)}.

**Remark 2.3.** Lemma 2.2 shows that if the vector reproduction number is less than or equal to one, the vector population will become extinct, while if the vector reproduction number is greater than one, the vector population will eventually stabilize at a positive equilibrium  $(L_f^*, N_f^*)$ .

### 3. Stability analysis of $E_{00}$ and $E_0$

System (2.2) always exists one disease-free and vector-free equilibrium  $E_{00} = (0, 0, 0, 0, S_h^0, 0, 0)$  with  $S_h^0 = \frac{\Lambda}{\mu_h}$ . And system (2.2) admits one disease-free equilibrium  $E_0 = (L_f^*, N_f^*, 0, 0, S_h^0, 0, 0)$  when  $R_v > 1$ .

Following [41,42], when  $R_v > 1$ , the basic reproduction number of system (2.2) is given by

$$R_0 = \rho(F_1 V_1^{-1}) = \frac{\beta_h \beta_f \lambda_g N_f^*}{\mu_g(\mu_f + P)(\mu_h + \delta_h + \gamma_h)},$$

where

$$F_{1} = \begin{pmatrix} 0 & \beta_{f} N_{f}^{*} & 0 \\ 0 & 0 & 0 \\ \frac{\beta_{h} S_{h}^{0}}{N_{h}} & 0 & 0 \end{pmatrix}$$

and

1

$$V_1 = \begin{pmatrix} \mu_f + P & 0 & 0 \\ 0 & \mu_g & -\lambda_g \\ 0 & 0 & \mu_h + \delta_h + \gamma_h \end{pmatrix}.$$

Here,  $\frac{1}{\mu_h + \delta_h + \gamma_h}$  is the average life span of a human,  $\frac{1}{\mu_g}$  represents the average life span of a cercaria,  $\frac{\lambda_g \beta_f N_f^*}{\mu_g}$  denotes the rate at which the cercariae infect the fish,  $\frac{1}{\mu_f + P}$  represents the average life span of an adult fish, and  $\frac{\beta_h}{\mu_f + P}$  is the rate at which the infected fish infect the susceptible people.

#### 3.1. Local asymptotic stability

The Jacobian matrix taken at  $E_{00} = (0, 0, 0, 0, S_h^0, 0, 0)$  is

$$J_{00} = \begin{pmatrix} -(\lambda_f + \sigma) & b & 0 & 0 & 0 & 0 & 0 \\ \lambda_f & -(\mu_f + P) & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & -(\mu_f + P) & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & -\mu_g & 0 & \lambda_g & 0 \\ 0 & 0 & 0 & 0 & -\mu_h & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & -(\mu_h + \gamma_h + \delta_h) & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & \gamma_h & -\mu_h \end{pmatrix}.$$

Let  $-J_{00} = (a_{ij})$ , where i, j = 1, 2, 3, 4, 5, 6, 7. Clearly,  $a_{ii} > 0$  and  $a_{ij} \leq 0$ . The leading principal minors of  $-J_{00}$  are  $\lambda_f + \sigma$ ,  $(\lambda_f + \sigma)(\mu_f + P)(1 - R_v)$ ,  $(\lambda_f + \sigma)(\mu_f + P)^2(1 - R_v)$ ,  $\mu_g(\lambda_f + \sigma)(\mu_f + P)^2(1 - R_v)$ ,  $\mu_\mu\mu_g(\lambda_f + \sigma)(\mu_f + P)^2(1 - R_v)$ ,  $\mu_\mu\mu_g(\mu_f + P)^2(\mu_h + \gamma_h + \delta_h)(\lambda_f + \sigma)(1 - R_v)$ ,  $\mu_\mu^2\mu_g(\mu_f + P)^2(\mu_h + \gamma_h + \delta_h)(\lambda_f + \sigma)(1 - R_v)$ . We easily find that they are all positive if and only if  $R_v < 1$ . By the M-matrix theory [43], we find that  $-J_{00}$  is an M-matrix when  $R_v < 1$ , implies that all eigenvalues of  $-J_{00}$  have positive real parts. Accordingly, all eigenvalues of  $J_{00}$  have negative real parts when  $R_v < 1$ . We then conclude that  $E_{00}$  is locally asymptotically stable when  $R_v < 1$ .

**Theorem 3.1.** If  $R_v < 1$ , the disease-free and vector-free equilibrium  $E_{00} = (0, 0, 0, 0, S_h^0, 0, 0)$  of system (2.2) is locally asymptotically stable. If  $R_v > 1$ ,  $E_{00}$  is unstable.

As the proof in Section 2, system (2.2) has one disease-free equilibrium  $E_0 = (L_f^*, N_f^*, 0, 0, S_h^0, 0, 0)$  when  $R_v > 1$ . The characteristic polynomial of  $E_0$  is

$$(\lambda + \mu_h)^2 \det(\lambda I - J_{01}) = 0,$$

where

$$J_{01} = \begin{pmatrix} -(\lambda_f + \sigma + 2\alpha L_f^*) & b & 0 & 0 & 0 \\ \lambda_f & -(\mu_f + P) & 0 & 0 & 0 \\ 0 & 0 & -(\mu_f + P) & \beta_f N_f^* & 0 \\ 0 & 0 & 0 & -\mu_g & \lambda_g \\ 0 & 0 & \frac{\beta_h S_h^0}{N_h} & 0 & -(\mu_h + \gamma_h + \delta_h) \end{pmatrix}$$

Let  $-J_{01} = (b_{ij})$ , where i, j = 1, 2, 3, 4, 5. Clearly,  $b_{ii} > 0$  and  $b_{ij} \leq 0$ . The leading principal minors of  $-J_{01}$  are  $\lambda_f + \sigma + 2\alpha L_f^*$ ,  $(\lambda_f + \sigma)(\mu_f + P)(R_v - 1), (\lambda_f + \sigma)(\mu_f + P)^2(R_v - 1), \mu_g(\lambda_f + \sigma)(\mu_f + P)^2(R_v - 1), \mu_g(\lambda_f + \sigma)(\mu_f + P)^2(\mu_h + \gamma_h + \delta_h)(R_v - 1)(1 - R_0)$ . Hence, they are all positive if and only if  $R_0 < 1$ . Clearly,  $-J_{01}$  is an M-matrix when  $R_v > 1$  and  $R_0 < 1$ , which means that all eigenvalues of  $-J_{01}$  have positive real parts. Accordingly, all eigenvalues of  $J_{01}$  have negative real parts when  $R_v > 1$  and  $R_0 < 1$ . We then conclude that  $E_0$  is locally asymptotically stable when  $R_v > 1$  and  $R_0 < 1$ .

**Theorem 3.2.** If  $R_v > 1$  and  $R_0 < 1$ , the disease-free equilibrium  $E_0 = (L_t^*, N_t^*, 0, 0, S_b^0, 0, 0)$  of system (2.2) is locally asymptotically stable.

3.2. Global stability

**Theorem 3.3.** If  $R_v < 1$ , the disease-free and vector-free equilibrium  $E_{00} = (0, 0, 0, 0, S_h^0, 0, 0)$  of system (2.2) is globally asymptotically stable in *K*.

**Proof.** As the conclusion in Theorem 3.1, when  $R_v < 1$ ,  $E_{00}$  is locally asymptotically stable. It is necessary to prove that  $u(t) = (L_f(t), N_f(t), I_f(t), G(t), S_h(t), I_h(t), R_h(t)) \rightarrow E_{00}$ , as  $t \rightarrow \infty$ , for  $u(0) = (L_f(0), N_f(0), I_f(0), G(0), S_h(0), I_h(0), R_h(0)) \in K$ . As Lemma 2.2, when  $R_v < 1$ , we have  $(L_f(t), N_f(t)) \rightarrow (0, 0)$ , then  $I_f(t) \rightarrow 0$ , where  $t \rightarrow \infty$ . Then, we have

$$\lim_{t\to\infty} (G(t), S_h(t), I_h(t), R_h(t)) = (0, \frac{\Lambda}{\mu_h}, 0, 0).$$

By [31, Theorem 2.6 ], we know that  $(0, \frac{\Lambda}{\mu_h}, 0, 0)$  is globally attractive in  $\mathbb{R}^4$ . Thus,  $E_{00}$  is globally attractive in *K*.

**Theorem 3.4.** If  $R_v > 1$  and  $R_0 < R_1 := \frac{\mu_h}{\mu_h + \delta_h}$ , the disease-free equilibrium  $E_0 = (L_f^*, N_f^*, 0, 0, S_h^0, 0, 0)$  of system (2.2) is globally asymptotically stable in K with  $L_f(0) > 0$  and  $N_f(0) > 0$ .

**Proof.** As Lemma 2.2 and  $R_v > 1$ , we have  $\lim_{t\to\infty}(L_f(t), N_f(t)) = (L_f^*, N_f^*)$ , where  $L_f(0) > 0$  and  $N_f(0) > 0$ , then system (2.2) has the following limiting system:

$$\begin{cases} \frac{dI_f(t)}{dt} = \beta_f (N_f^* - I_f)G - (\mu_f + P)I_f, \\ \frac{dG(t)}{dt} = \lambda_g I_h - \mu_g G, \\ \frac{dS_h(t)}{dt} = \Lambda - \frac{\beta_h S_h}{N_h} I_f - \mu_h S_h, \\ \frac{dI_h(t)}{dt} = \frac{\beta_h S_h}{N_h} I_f - (\mu_h + \delta_h + \gamma_h)I_h, \\ \frac{dR_h(t)}{dt} = \gamma_h I_h - \mu_h R_h. \end{cases}$$
(3.1)

Adding the last three equations of system (3.1) gives

$$\begin{split} \Lambda - (\mu_h + \delta_h) N_h &\leq \frac{dN_h(t)}{dt} \\ &= \Lambda - \mu_h N_h - \delta_h I_h \\ &\leq \Lambda - \mu_h N_h. \end{split}$$

This implies that  $\phi_1 \leq \lim_{t \to \infty} \inf N_h(t) \leq \lim_{t \to \infty} \sup N_h(t) \leq S_h^0$ , where  $\phi_1 = \frac{\Lambda}{\mu_1 + \delta_1}$ .

By system (3.1), for sufficiently large t,

$$\frac{dI_{f}(t)}{dt} \leq \beta_{f} N_{f}^{*} G - (\mu_{f} + P) I_{f},$$

$$\frac{dG(t)}{dt} = \lambda_{g} I_{h} - \mu_{g} G,$$

$$\frac{dI_{h}(t)}{dt} \leq \frac{\beta_{h} S_{h}^{0}}{\phi_{1}} I_{f} - (\mu_{h} + \delta_{h} + \gamma_{h}) I_{h}.$$
(3.2)

Define the following auxiliary linear system by (3.2):

$$\begin{cases} \frac{d\widetilde{I_f}(t)}{dt} = \beta_f N_f^* \widetilde{G} - (\mu_f + P) \widetilde{I_f}, \\ \frac{d\widetilde{G}(t)}{dt} = \lambda_g \widetilde{I_h} - \mu_g \widetilde{G}, \\ \frac{d\widetilde{I_h}(t)}{dt} = \frac{\beta_h S_h^0}{\phi_1} \widetilde{I_f} - (\mu_h + \delta_h + \gamma_h) \widetilde{I_h}. \end{cases}$$
(3.3)

The right-hand side of system (3.3) has coefficient matrix J given by

$$\begin{pmatrix} -(\mu_f+P) & \beta_f N_f^* & 0 \\ 0 & -\mu_g & \lambda_g \\ \frac{\beta_h S_h^0}{\phi_1} & 0 & -(\mu_h+\delta_h+\gamma_h) \end{pmatrix}$$

The leading principal minors of -J are  $\mu_f + P$ ,  $\mu_g(\mu_f + P)$  and  $\beta_h \beta_f N_f^* \lambda_g \left(\frac{1}{R_0} - \frac{\mu_h + \delta_h}{\mu_h}\right)$ . Hence, they are all positive if and only if

$$R_0 \frac{\mu_h + \delta_h}{\mu_h} < 1.$$

Namely,  $R_0 < R_1 := \frac{\mu_h}{\mu_h + \delta_h}$ . Obviously, we obtain -J is an Mmatrix when  $R_0 < R_1$ , which means that all eigenvalues of -J have positive real parts. Consequently, any eigenvalue of J lies in the left half plane. Thus, any nonnegative solution of system (3.3) satisfies  $\lim_{t\to\infty} (\widetilde{I_f}, \widetilde{G}, \widetilde{I_h}) = (0, 0, 0)$ . Since system (3.3) is a linear system, the zero solution (0,0,0) of system (3.3) is globally asymptotically stable. As a consequence of the comparison principle, we obtain that any nonnegative solution of system (3.2) satisfies  $\lim_{t\to\infty} (I_f, G, I_h) =$ (0, 0, 0).  $S_h$  and  $R_h$  in system (3.1) satisfy the following limiting system:

$$\begin{cases} \frac{dS_h(t)}{dt} = \Lambda - \mu_h S_h \\ \frac{dR_h(t)}{dt} = -\mu_h R_h. \end{cases}$$

It then follows that  $\lim_{t\to\infty} (S_h(t), R_h(t)) = (S_h^0, 0)$  and  $E_0 = (L_f^*, N_f^*, 0, 0, S_h^0, 0, 0)$  is globally asymptotically stable when  $R_v > 1$  and  $R_0 < R_1$ .

## 4. Global stability of the endemic equilibrium

#### 4.1. The endemic equilibrium

Let

$$\lambda_1 = \frac{\beta_h I_f}{N_h}, \ \lambda_2 = \beta_h G. \tag{4.1}$$

We consider the case when  $R_v > 1$ , in which case  $(L_f, N_f) = (L_f^*, N_f^*)$ . The other components of the endemic equilibrium require to satisfy the

## following conditions:

$$A = \lambda_1 S_h + \mu_h S_h,$$

$$\lambda_1 S_h = (\mu_h + \delta_h + \gamma_h) I_h,$$

$$\gamma_h I_h = \mu_h R_h,$$

$$\lambda_g I_h = \mu_g G,$$

$$\beta_f (N_f^* - L_f) G = (\mu_f + P) I_f.$$
(4.2)

Solving Eq. (4.2) in terms of  $\lambda_1$  and  $\lambda_2$ , we have

$$I_{f} = \frac{\lambda_{2}N_{f}^{*}}{\mu_{f} + P + \lambda_{2}}, \ G = \frac{\lambda_{g}\Lambda\lambda_{1}}{\mu_{g}(\lambda_{1} + \mu_{h})(\mu_{h} + \delta_{h} + \gamma_{h})}, \ S_{h} = \frac{\Lambda}{\lambda_{1} + \mu_{h}},$$

$$I_{h} = \frac{\Lambda\lambda_{1}}{(\lambda_{1} + \mu_{h})(\mu_{h} + \delta_{h} + \gamma_{h})}, \ R_{h} = \frac{\gamma_{h}\Lambda\lambda_{1}}{\mu_{h}(\lambda_{1} + \mu_{h})(\mu_{h} + \delta_{h} + \gamma_{h})}.$$
(4.3)

Then

$$N_{h} = \frac{\Lambda}{\lambda_{1} + \mu_{h}} \left( 1 + \frac{\lambda_{1}}{\mu_{h} + \delta_{h} + \gamma_{h}} + \frac{\gamma_{h}\lambda_{1}}{\mu_{h}(\mu_{h} + \delta_{h} + \gamma_{h})} \right).$$
(4.4)

Substituting Eqs. (4.3) and (4.4) into Eq. (4.1), we obtain

$$\lambda_{1} = \frac{\lambda_{1}\lambda_{g}\beta_{h}N_{f}^{*}K_{3}}{\lambda_{1}K_{3}\lambda_{g} + K_{2}(\lambda_{1} + \mu_{h})} \frac{\lambda_{1} + \mu_{h}}{K_{1}\left[\mu_{h}(\mu_{h} + \delta_{h} + \gamma_{h}) + \lambda_{1}\mu_{h} + \gamma_{h}\lambda_{1}\right]}, \quad (4.5)$$
$$\lambda_{2} = \frac{K_{3}\lambda_{1}\lambda_{g}}{\lambda_{1}}, \quad (4.6)$$

 $\mu_2 = \mu_g(\lambda_1 + \mu_h)$ 

where

$$K_1 = \frac{\Lambda}{\mu_h(\mu_h + \delta_h + \gamma_h)}, \ K_2 = \mu_g(\mu_f + P), \ K_3 = \frac{\Lambda\beta_f}{\mu_h + \delta_h + \gamma_h}.$$

Substituting Eqs. (4.6) into (4.5) and dividing by  $\lambda_1$ , we let

$$1 = \frac{\lambda_g \beta_h N_f^* K_3}{\lambda_1 K_3 \lambda_g + K_2 (\lambda_1 + \mu_h)} \frac{\lambda_1 + \mu_h}{K_1 \left[ \mu_h (\mu_h + \delta_h + \gamma_h) + \lambda_1 \mu_h + \gamma_h \lambda_1 \right]}.$$
 (4.7)

From Eqs. (4.3) and (4.6), we can show that for  $\lambda_1 > 0$ , and system (2.2) has an endemic equilibrium.

We begin by discussing the case when  $\delta_h=0,$  in which case Eq. (4.7) can be written as

$$1 = \frac{\lambda_g \beta_h N_f^* K_3}{\lambda_1 K_3 \lambda_g + K_2 (\lambda_1 + \mu_h)} \frac{\mu_h (\lambda_1 + \mu_h) (\mu_h + \gamma_h)}{\Lambda \left[ \mu_h (\mu_h + \gamma_h) + \lambda_1 \mu_h + \gamma_h \lambda_1 \right]}$$
  
= 
$$\frac{\lambda_g \beta_h N_f^* K_3}{\frac{\lambda_1}{\lambda_1 + \mu_h} K_3 \lambda_g + K_2} \frac{\mu_h (\mu_h + \gamma_h) + \lambda_1 \mu_h + \gamma_h \lambda_1}{\Lambda \left[ \mu_h (\mu_h + \gamma_h) + \lambda_1 \mu_h + \gamma_h \lambda_1 \right]}.$$
(4.8)

Let  $M(\lambda_1)$  represent the right side of Eq. (4.8), it is clearly to see that  $M(\lambda_1)$  is a decreasing function for  $\lambda_1 \in (0, \infty)$ , which implies that  $M(\lambda_1)$  approaches zero when  $\lambda_1 \to \infty$ . Hence, if M(0) > 1, or, equivalently,  $R_0 > 1$ , (4.8) has a unique positive root such that system (2.2) has a unique endemic equilibrium. In contrast, there is no endemic equilibrium if  $M(0) \le 1$  or, equivalently,  $R_0 \le 1$ . Then we obtain the following result.

**Theorem 4.1.** Assume that  $\delta_h = 0$ . If  $R_v > 1$  and  $R_0 > 1$ , then there is a unique endemic equilibrium of system (2.2). Otherwise, there is no endemic equilibrium for system (2.2).

For 
$$\delta_h \ge 0$$
, we reorganize the terms in Eq. (4.7) gives

$$D_1 \lambda^2 + D_2 \lambda + D_3 = 0, (4.9)$$

where

$$D_{1} = (\mu_{h} + \gamma_{h})(K_{1}K_{3}\lambda_{g} + K_{1}K_{2}),$$

$$D_{2} = K_{1}K_{2}\mu_{h}(2\mu_{h} + \delta_{h} + \gamma_{h}) + K_{1}K_{3}\lambda_{g}\mu_{h}(\mu_{h} + \delta_{h} + \gamma_{h}) - K_{3}\lambda_{g}\beta_{f}N_{f}^{*},$$

$$D_{3} = K_{1}K_{2}\mu_{h}^{2}(\mu_{h} + \delta_{h} + \gamma_{h})(1 - R_{0}).$$
(4.10)

It is clear that  $D_1 > 0$  and  $D_3$  has the same sign as  $1 - R_0$ . Define  $h(z) = D_1 z^2 + D_2 z + D_3$ , (4.11) then the roots of Eq. (4.11) can be expressed as

$$z_{1,2} = \frac{-D_2 \pm \sqrt{D_2^2 - 4D_1D_3}}{2D_1} = \frac{-D_2 \pm \sqrt{\Delta}}{2D_1}.$$

If  $\Delta \leq 0$ , then Eq. (4.11) does not have a positive solution, and Eq. (4.10) has no positive real roots. If  $\Delta > 0$  and  $h(0) = D_3 < 0$ , then Eq. (4.11) has a unique positive root, and Eq. (4.10) exists a unique positive solution.

Summarizing the above discussions, we have the following theorem.

**Theorem 4.2.** Assume that  $R_v > 1$ , let  $\Delta$  be defined by (4.11). (i) If  $\Delta > 0$  and  $R_0 > 1$ , then system (2.2) has a unique endemic equilibrium; (ii) If  $\Delta \le 0$ , then system (2.2) has no endemic equilibrium.

#### 4.2. Global stability of the endemic equilibrium

**Theorem 4.3.** If  $\Delta > 0$ ,  $R_v > 1$  and  $R_0 > 1$ , the endemic equilibrium  $E^* = (L_f^*, N_f^*, I_f^*, G^*, S_h^*, I_h^*, R_h^*)$  of system (2.2) is globally asymptotically stable in Int(K).

**Proof.** By Lemma 2.2, the vector population will eventually stabilize at a positive equilibrium  $(L_f^*, N_f^*)$  if  $R_v > 1$ . Hence, the variables  $I_f$ , G,  $I_h$  and  $R_h$  satisfy the following limiting system:

$$\begin{cases} \frac{dI_{f}(t)}{dt} = \beta_{f}(N_{f}^{*} - I_{f}^{*})G - (\mu_{h} + P)I_{f}, \\ \frac{dG(t)}{dt} = \lambda_{g}I_{h} - \mu_{g}G, \\ \frac{dI_{h}(t)}{dt} = \frac{\beta_{h}(S_{h}^{0} - I_{h} - R_{h})I_{f}}{S_{h}^{0}} - (\mu_{h} + \delta_{h} + \gamma_{h})I_{h}, \\ \frac{dR_{h}(t)}{dt} = \gamma_{h}I_{h} - \mu_{h}R_{h}. \end{cases}$$
(4.12)

Let  $V = [0, N_f^*] \times \left[0, \frac{\lambda_g}{\mu_g} S_h^0\right] \times [0, S_h^0] \times \left[0, \frac{\gamma_h}{\mu_h} S_h^0\right]$ , it then follows that  $\omega(I_f(0), G(0), I_h(0), R_h(0)) \in V$ , where  $\omega(I_f(0), G(0), I_h(0), R_h(0))$  is the omega limit set of  $(I_f(0), G(0), I_h(0), R_h(0)) \in \mathbb{R}_+^4$  for the solution semiflow of system (4.12). It is easy to verify that V is positively invariant for system (4.12).

Let

$$j(u) = \begin{pmatrix} \beta_f(N_f^* - u_1)u_2 - (\mu_f + P)u_1 \\ \lambda_g u_3 - \mu_g u_2 \\ \frac{\beta_h(S_h^0 - u_3 - u_4)u_1}{S_h^0} - (\mu_h + \delta_h + \gamma_h)u_3 \\ \gamma_h u_3 - \mu_h u_4 \end{pmatrix},$$

then  $j : \mathbb{R}^4_+ \to \mathbb{R}^4$  is a continuously differentiable map. Clearly, j(0) = 0 and  $j_m(u) \ge 0$  for all  $u \in V$  with  $u_m = 0$ , m = 1, 2, 3, 4. Since

$$D_{j}(u) = \frac{\partial j_{m}}{\partial u_{n}} = \begin{pmatrix} -\beta_{f}u_{2} - (\mu_{f} + P) & \beta_{f}(N_{f}^{*} - u_{1}) & 0 & 0\\ 0 & -\mu_{g} & \lambda_{g} & 0\\ \frac{\beta_{h}(S_{h}^{0} - u_{5} - u_{4})}{S_{h}^{0}} & 0 & -\frac{\beta_{h}u_{1}}{S_{h}^{0}} - (\mu_{h} + \delta_{h} + \gamma_{h}) & -\frac{\beta_{h}u_{1}}{S_{h}^{0}}\\ 0 & 0 & \gamma_{h} & -\mu_{h} \end{pmatrix},$$

then  $\frac{\partial j_m}{\partial u_n} \ge 0$ ,  $(m \ne n)$  for  $u \in V$ , thus *j* is cooperative on *V*. Clearly,  $D_j(u)$  is irreducible for every  $u \in V$ . For any  $\rho \in (0, 1)$  and  $(u_1, u_2, u_3, u_4) \in \operatorname{Int}(\mathbb{R}^4_+)$ , we have

$$\begin{aligned} j_1(\rho u_1, \rho u_2, \rho u_3, \rho u_4) = & \beta_f(N_f^* - \rho u_1)\rho u_2 - (\mu_f + P)\rho u_1 \\ > & \beta_f(N_f^* - u_1)\rho u_2 - (\mu_f + P)\rho u_1 \\ = & \rho j_1(u_1, u_2, u_3, u_4). \end{aligned}$$

Similarly, we can show that  $j_m(\rho u_1, \rho u_2, \rho u_3, \rho u_4) = \rho j_m(u_1, u_2, u_3, u_4)$ , m = 2, 4 and  $j_3(\rho u_1, \rho u_2, \rho u_3, \rho u_4) > \rho j_3(u_1, u_2, u_3, u_4)$ . Thus, *j* is strictly

## Table 2

Summary of the existence and stability for  $E_{00}$ ,  $E_0$  and  $E_0$ .

Equilibrium	Existence	LAS	GAS
$E_{00} = (0, 0, 0, 0, S_h^0, 0, 0)$	always	$R_v < 1$	<i>R<sub>v</sub></i> < 1
$E_0 = (L_f^*, N_f^*, 0, 0, S_h^0, 0, 0)$	$R_v > 1$	$R_v > 1, R_0 < 1$	$R_v > 1, R_0 < R_1$
$E^* = (L_f^*, N_f^*, I_f^*, G^*, S_h^*, I_h^*, R_h^*)$	$\Delta > 0, R_v > 1, R_0 > 1$	$\Delta > 0, R_v > 1, R_0 > 1$	$\Delta > 0, R_v > 1, R_0 > 1$



Fig. 5.1. The simulation result of Clonorchiasis cases from 2016 to 2021 in Guangxi, China.

sublinear on V. Since

$$D_{j}(0) = \begin{pmatrix} -(\mu_{f} + P) & \beta_{f} N_{f}^{*} & 0 & 0\\ 0 & -\mu_{g} & \lambda_{g} & 0\\ \beta_{h} & 0 & -(\mu_{h} + \delta_{h} + \gamma_{h}) & 0\\ 0 & 0 & \gamma_{h} & -\mu_{h} \end{pmatrix},$$
(4.13)

the leading principal minors of  $-D_j(0)$  are  $\mu_f + P$ ,  $\mu_g(\mu_f + P)$  and  $\beta_h\beta_f N_f^*\lambda_g\left(\frac{1}{R_0}-1\right)$ . Obviously,  $\beta_h\beta_f N_f^*\lambda_g\left(\frac{1}{R_0}-1\right) < 0$  when  $R_0 > 1$ . By the M-matrix theory [43], we get  $-D_j(0)$  has at least one eigenvalue with negative real part, which means  $D_j(0)$  has at least one eigenvalue with positive real part. Then the spectral bound of  $D_j(0)$ ,  $s(D_j(0)) := \max\{\text{Re}\lambda : \det(\lambda I - D_j(0)) = 0\} > 0$ . It then follows from [39, Corollary 3.2], the positive equilibrium  $(I_f^*, S_h^*, I_h^*, R_h^*)$  for system (4.12) is globally asymptotically stable in  $\mathbb{R}_+^4 \setminus \{(0, 0, 0, 0)\}$ .

$$\lim_{t \to \infty} S_h(t) = \lim_{t \to \infty} (N_h(t) - I_h(t) - R_h(t)) = S_h^0 - I_h^* - R_h^* = S_h^*$$

We finally obtain that the endemic equilibrium  $E^* = (L_f^*, N_f^*, I_f^*, G^*, S_h^*, I_h^*, R_h^*)$  of system (2.2) is globally asymptotically stable in Int(*K*).

Table 2 summarizes the existence and stability conditions for equilibria  $E_{00}$ ,  $E_0$  and  $E^*$ .

#### 5. A case study

In this section, we perform some numerical simulations based on a real case of Clonorchiasis transmission in Guangxi, China.

#### 5.1. Model validation

In 2016, the region with the highest burden of Clonorchiasis disease in China was the Guangxi, China. We simulate the Clonorchiasis transmission case in Guangxi, China based on the data from the Table 3.

According to the analysis of the disease burden of Clonorchiasis [21], the DALY in Guangxi, China, is 5.05. We assume the diseaseinduced mortality rate for humans  $\delta_h = 0.00505$ . For convenience, we define  $\beta_H = \frac{\beta_h}{N_h}$ . Over 100 freshwater fish species have been identified as intermediate hosts of clonorchis sinensis [45]. Different fish have different effects on disease transmission in the ecological environment. Usually, it takes 3–4 years for fish to become reproductively capable from birth, so we chose  $\lambda_f = 0.3$ . The lifespan of the larvae fish is about three years, and we choose  $\sigma = 0.3$ . Table 1 summarizes these parameters. We select the initial values:  $G^0 = 3 \times 10^5$ ,  $S_h^0 = 35000$ ,  $I_h^0 = 2791332$ ,  $I_f^0 = 12080$ ,  $N_h = 23261104$  from [9], and assume that  $L_f^0 = 30800$ ,  $N_f^0 = 20800$ .

According to the above-estimated parameter values and the initial values, we fitted the Guangxi, China Clonorchiasis cases by system (2.2). The reported data and the simulation result based on our model, are shown in Fig. 5.1. As we can see, they match very well. We compute the basic reproduction number in this case,  $R_v = 79.0583$  and  $R_0 = 1.4764 > 1$ . To explore the impact of control measures on long-term trends in Clonorchiasis, we adjust the values of  $\beta_H$  and  $\lambda_g$ . In Figs. 5.2 and 5.3, we observe that reducing  $\lambda_g$  is the most effective method, and just a single control of  $\beta_H$  cannot eradicate the disease. Reducing the concentration of cercariae in the water environment can effectively decrease the rate of disease in fish, then control the spread of Clonorchiasis in the population.

### 5.2. Sensitivity analysis

The basic reproduction number reflects the outbreak potential and severity of the disease. In order to take more targeted and effective measures to control the prevalence of Clonorchiasis, we evaluate the influence of parameters on  $R_0$  by sensitivity analysis of the model. Therefore, we use the Latin Hypercube Sampling (LHS) method to evaluate partial rank correlation coefficients (PRCC) for various input parameters against output variables, to determine which parameters can be adjusted to more effectively intervene in the transmission of Clonorchiasis [46]. Parameters with larger absolute values of PRCC have a more significant effect on disease, where the positive PRCC value represents a positive effect on the basic reproduction number and

## Table 3

Population data related to clonorchiasis in	Guangxi, China from	2016 to 2021 [44]
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Year	2016	2017	2018	2019	2020	2021
Population size $(* 10^4)$	4838	4926	4960	5019	5037	5013
Number of infectious humans	2791332	1 442 345	2 721 221	2534064	2165916	2435271



Fig. 5.2. The simulation result of Clonorchiasis cases from 2016 to 2021 in Guangxi, China. The black curve is the system simulation. Colored lines is the system simulation when we take control of  $\beta_H$ .



Fig. 5.3. The simulation result of Clonorchiasis cases from 2016 to 2021 in Guangxi, China. The black curve is the system simulation. Colored lines is the system simulation when we take control of  $\lambda_g$ .

a negative impact on disease control, and the negative PRCC value is the opposite.

We obtain the sensitivity index of  $R_0$  to all the parameters of the system in Table 4 and plot the sensitivity analysis of  $R_0$ . Fig. 5.4 shows the global sensitivity analysis of  $R_0$ .  $\mu_f + P$ ,  $\lambda_f$  and  $\beta_f$  are the parameters that affect  $R_0$  to a large extent, suggesting that fish play a vital role in the propagation of Clonorchiasis. Obviously,  $R_0$  is an increasing function of parameters  $b, \lambda_f, \lambda_g, \beta_H, \beta_f$ , respectively. As  $\mu_f + P, \mu_g, \mu_h, \gamma_h, \delta_h$  increase,  $R_0$  is reducing but insensitive to the parameters  $\mu_h$  and  $\delta_h$ . It means that when controlling the transmission

of Clonorchiasis, effective measures need to be taken, such as purifying water quality, cultural education, or strengthening treatment, to reduce  $\lambda_g$ ,  $\beta_H$ ,  $\beta_f$ , and increase  $\mu_g$ ,  $\gamma_h$ ,  $\delta_h$ .

#### 5.3. $R_0$ and long-term behaviors

To simulate the long-term behavior of Clonorchiasis, we change some parameters to satisfy the theorem in Section 3. In Fig. 5.5, we change *b*,  $\lambda_f$  and  $\mu_f + P$ , then we get  $R_v < 1$  and  $R_0 < 1$ . By Theorem 3.3, the disease-free and vector-free equilibrium  $E_{00}$  is

#### Table 4

Sensitivity index of  $R_0$ .

Input parameter	PRCC	Input parameter	PRCC
b	0.193729	$\beta_H$	0.272050
$\lambda_f$	0.356660	$\mu_h$	-0.016188
$\mu_f + P$	-0.552336	$\gamma_h$	-0.194397
$\lambda_{g}$	0.217804	$\delta_h$	-0.013166
$\mu_g$	-0.212093	$\beta_f$	0.311500



Fig. 5.4. Sensitivity analysis diagram of  $R_0$ .

globally asymptotically stable. In Fig. 5.7, we change  $\lambda_g$ ,  $\mu_g$ ,  $\beta_H$ ,  $\beta_f$  and  $\mu_f + P$ , then we get  $R_v > 1$  and  $R_0 < 1$ . Theorem 3.4 implies that the solution converges to the disease-free equilibrium  $E_0$ , and the disease eventually becomes extinct.

Then, we explore the effect of different parameters on the basic reproduction number  $R_0$ . From Fig. 5.6(a), we have  $R_0 = 1.2655$  when the disease recovery rate  $\gamma_h = 1$ . This means that the disease is still prevalent even if all patients can recover. It can be seen that prevention is better than cure for Clonorchiasis. Larval fish have relatively weak immune mechanisms and are more susceptible to infection by caecilians in the water. In fact, regular water system cleaning is feasible. To introduce water system cleaning in system (2.2), we replace  $\lambda_g$  with  $(1 - \pi)\lambda_g$ , where  $\pi \in [0.2, 1]$  denotes the degree of completion of water system cleaning. Keeping other parameter values the same as in Table 1, Fig. 5.6(b) shows that  $R_0$  decreases with increasing  $\pi$ . Therefore, to improve the water quality monitoring system, regular clean water, and dismantle toilets at the edge of ponds may effectively control Clonorchiasis.

The infection rate of the host by the parasite plays an important role in disease transmission. Fig. 5.8 simulates the effect of  $\beta_H$  and  $\beta_f$  on  $R_0$ , which is an increasing function with respect to  $\beta_H$  and  $\beta_f$ , respectively. A sufficiently small basic reproduction number can be achieved by controlling either  $\beta_H$  or  $\beta_f$ . Although there is no commercially produced vaccine against Clonorchiasis, the possibility of developing a fish vaccine has been proposed [32]. To reduce the basic reproduction number to less than one, we need to reduce the prevalence of Clonorchiasis infection in fish by at least 56%. In addition to enhancing cultural education to reduce the consumption of raw fish, using non-polluted water for fish farming may also be a reasonable measure to reduce  $\beta_f$  and  $\beta_H$  effectively.

In Fig. 5.9, we demonstrate the relationship between  $R_0$  and  $(\beta_H, \beta_f)$  in three-dimensional space. We can observe the value of  $R_0$  by changing the value of the parameter  $\beta_H$  and  $\beta_f$  with the other parameters unchanged. We can see that even if  $\beta_H$  is large,  $R_0$  can be smaller than one as long as the value of  $\beta_f$  is small enough. Meanwhile, when  $\beta_f$  is large,  $R_0$  can be smaller than one as long as  $\beta_H$  is small enough. The parameters  $\beta_H$  and  $\beta_f$  are important in Clonorchiasis transmission, and they determine the trend of  $R_0$  together. Fish health is a key factor



Fig. 5.5. Long-term behavior of  $L_f$ ,  $I_f$ , G and  $I_h$  when  $R_v < 1$  and  $R_0 < 1$ . Black line: b = 0.45,  $R_v = 0.7906$ ,  $R_0 = 0.1476$ . Blue line:  $\mu_f + P = 28.46$ ,  $R_v = 0.7906$ ,  $R_0 = 0.0015$ . Red line:  $\lambda_f = 0.0015$ ,  $R_v = 0.7867$ ,  $R_0 = 0.0074$ . Yellow line: b = 30,  $\lambda_f = 0.0025$ ,  $R_v = 0.8712$ ,  $R_0 = 0.1$ . Green line: b = 25,  $\lambda_f = 12.846$ ,  $R_v = 0.9731$ ,  $R_0 = 0.0036$ .







Fig. 5.7. Long-term behavior of  $L_f$ ,  $I_f$ , G and  $I_h$  when  $R_v > 1$  and  $R_0 < 1$ . Black line:  $\lambda_g = 60$ ,  $R_0 = 0.3591$ . Blue line:  $\mu_g = 26.07$ ,  $\beta_H = 10^{-6}$ ,  $R_0 = 0.2334$ . Red line:  $\mu_g = 26.07$ ,  $\beta_f = 3.59 \times 10^{-11}$ ,  $R_0 = 0.1476$ . Yellow line:  $\lambda_g = 200$ ,  $\mu_f + P = 0.5$ ,  $R_0 = 0.2816$ . Green line:  $\lambda_g = 500$ ,  $\beta_H = 4 \times 10^{-7}$ ,  $R_0 = 0.1674$ .

affecting the prevalence of Clonorchiasis, and cultural education should be strengthened to avoid eating raw freshwater fish.

#### 6. Concluding remarks

Based on the mechanisms of transmission and related studies of Clonorchiasis, fish is the most essential link in transmitting the disease to humans. In this work, we propose a novel mathematical model to study the dynamics of Clonorchiasis around humans and vectors. Due to their biological considerations, fish are divided into two stages: larval and adult. Larval fish may be more susceptible to Clonorchiasis because they usually live in shallow waters and are more likely to come into contact with environments contaminated with Clonorchis sinensis eggs. The mathematical results show that (i) the disease-free and vector-free equilibrium  $E_{00}$  is GAS if  $R_v < 1$  holds; (ii) if  $R_v > 1$ , the disease-free

equilibrium  $E_0$  exists and is GAS for  $R_0 < R_1 := \frac{\mu_h}{\mu_h + \delta_h}$ ; (iii) the endemic equilibrium  $E^*$  exists and is GAS if  $\Delta > 0$ ,  $R_v > 1$  and  $R_0 > 1$ .

To demonstrate the practical value of the parameters of system (2.2), Fig. 5.1 conducted a case study on the transmission dynamics of Clonorchiasis from 2016–2021 using data from Guangxi, China. We obtain the basic reproduction number as  $R_0 = 1.4764$ , which means that the disease will persist if no action is taken. Figs. 5.2 and 5.3 predict the development of Clonorchiasis in Guangxi, China under the control of  $\beta_H$  and  $\lambda_g$ , respectively.

The basic reproduction number plays a decisive role in the spread of infectious diseases, and only by figuring out which factors affect the basic reproduction number can we better provide reasonable control measures. We conduct a sensitivity analysis of the parameters that may affect  $R_0$  in Fig. 5.4. The results show that fish is a key factor influencing the spread of disease and  $\lambda_g$ ,  $\beta_H$ , and  $\beta_f$  are important



**Fig. 5.8.** The effect of parameters  $\beta_H$  and  $\beta_f$  on  $R_0$ .



**Fig. 5.9.** The contour plot of the  $R_0$  as a function of  $\beta_H$  and  $\beta_f$ . (a) The red plane represents  $R_0 = 1$ , above the red lane  $R_0 > 1$ , below the red plane  $R_0 < 1$ . (b) The dashed line represents the value of  $R_0$ .

in influencing the spread of disease. After that, we explore the effect of these parameters on  $R_0$  in Figs. 5.6(b), 5.8 and 5.9. Fig. 5.6(a) implies that improving medical care is not enough and that integrated prevention and treatment measures must be taken. Clonorchis sinensis cannot attack humans directly to cause infection. They must invade the human body through food such as fish and shrimp, and the only way for organisms such as fish and shrimp to become infected is through caecilians in the water. Therefore, to eliminate Clonorchiasis, attention must be paid to controlling the concentration of cercariae in the water to reduce the probability of fish becoming infected with Clonorchiasis. This can be achieved through water purification and vaccination of fish.

The current treatment of Clonorchiasis is mainly praziquantel, but praziquantel is associated with serious adverse effects [47]. Studies have shown that Clonorchiasis can be prevented to a large extent by controlling the health status of fish. Raw fish consumption and cutting boards that do not distinguish between raw and cooked food can lead to disease infection, which requires the health sector to strengthen the culture and education of the people to raise their awareness and vigilance against Clonorchiasis. Preventive chemotherapy can be administered for risk groups such as schoolchildren and fishermen [4]. The entire life cycle of Clonorchis sinensis is limited by temperature and rainfall, and the growth state of the cercariae also heavily depends on temperature [46]. For future study, it would be interesting to incorporate these environmental drivers into the model and study their impact on Clonorchiasis transmission.

## CRediT authorship contribution statement

Wei Wang: Writing – original draft, Visualization, Validation, Methodology, Investigation, Funding acquisition, Formal analysis, Data curation, Conceptualization. Xiaohui Huang: Writing – original draft, Visualization, Software, Formal analysis. Hao Wang: Writing – review & editing, Validation, Resources, Project administration, Methodology, Investigation, Conceptualization.

## Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

### Data availability

Data were collected from public sources, which have been cited as references in the paper.

#### Acknowledgments

We thank the anonymous referees and the handling editor for their valuable comments which have led to a substantial improvement in the revision. Wei Wang gratefully acknowledges support from the National Natural Science Foundation of China (No. 12271308, 11901360). Hao Wang gratefully acknowledges support from the Natural Sciences and Engineering Research Council of Canada (Discovery Grant RGPIN-2020-03911 and Accelerator Grant RGPAS-2020-00090) and the CRC program (Tier 1 Canada Research Chair Award).

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