

http://www.aimspress.com/journal/MBE

MBE, 16(5): 4339–4358. DOI: 10.3934/mbe.2019216 Received: 02 February 2019 Accepted: 25 April 2019 Published: 16 May 2019

Research article

Transmission dynamics of cholera with hyperinfectious and hypoinfectious vibrios: mathematical modelling and control strategies

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Abstract: Cholera is a common infectious disease caused by Vibrio cholerae, which has different infectivity. In this paper, we propose a cholera model with hyperinfectious and hypoinfectious vibrios, in which both human-to-human and environment-to-human transmissions are considered. By analyzing the characteristic equations, the local stability of disease-free and endemic equilibria is established. By using Lyapunov functionals and LaSalle's invariance principle, it is verified that the global threshold dynamics of the model can be completely determined by the basic reproduction number. Numerical simulations are carried out to illustrate the corresponding theoretical results and describe the cholera outbreak in Haiti. The study of optimal control helps us seek cost-effective solutions of time-dependent control strategies against cholera outbreaks, which shows that control strategies, such as vaccination and sanitation, should be taken at the very beginning of the outbreak and become less necessary after a certain period.

Keywords: Cholera model; hyperinfectious vibrios; human-to-human transmission; environment-to -human transmission; global dynamics; optimal control

1. Introduction

Cholera is an acute disease caused by Vibrio cholerae O-group 1 or O-group 139, which can give rise to diarrhea and vomit. The World Health Organization (WHO) estimates that there are 1.3 to 4 million cholera cases per year with about 21,000 to 143,000 deaths all over the world [1]. Beginning in April 2017, a major cholera epidemic occurred in Yemen, with about 500,000 reported cases and 2,000 deaths. Due to the deterioration of health systems and associated infrastructures, cholera spreads more seriously. The WHO has announced that as many as 30,000 health care workers are devoted to the treatment in Yemen. At present, the epidemic situation of major cities including Sanaa, Hajja, and

Amran, is under control [1]. In order to better understand the transmission dynamics of cholera and provide some valuable insights on the prevention and control, some cholera models have been proposed (see, for example, [2, 3, 4, 5, 6]). In [2], Tien and Earn considered the following cholera model with both human-to-human and environment-to-human transmissions:

$$\frac{dS(t)}{dt} = \mu N - (\beta_h I(t) + \beta B(t)) S(t) - \mu S(t),$$

$$\frac{dI(t)}{dt} = (\beta_h I(t) + \beta B(t)) S(t) - (\gamma + \mu) I(t),$$

$$\frac{dR(t)}{dt} = \gamma I(t) - \mu R(t),$$

$$\frac{dB(t)}{dt} = \xi I(t) - \delta B(t),$$
(1.1)

where S(t), I(t) and R(t) denote the densities of susceptible, infected and recovered individuals, respectively; B(t) denotes the concentration of V. cholerae in contaminated water; μ is the birth and natural death rate; β_h and β are the human-to-human transmission rate and environment-to-human transmission rate, respectively; γ is the recovery rate of infected individuals; ξ is the contribution rate of each infected individual to the concentration of V. cholerae; δ is the net death rate of V. cholerae.

In [7], Hartley et al. found that short-lived, hyperinfectious state of vibrios decays in a matter of hours into a state of lower infectiousness, and incorporated this hyperinfectious state into a cholera model to provide a much better fit with the observed epidemic pattern. Besides, Neilan et al. [8] formulated a mathematical model to include two classes of bacterial concentrations, one is hyperinfectious and another is less-infectious. Furthermore, in [9], Mukandavire et al. indicated that human-to-human transmission is a very fast transmission process with a lower infectious dose as a result of immediate water or food contamination by hyperinfectious vibrios from freshly passed human stool. Besides, Mukandavire et al. pointed out that if vibrios have been in the environment for a sufficiently long period (anywhere from 5 to 18 hours), they are no longer hyperinfectious, namely, in a hypoinfectious state.

Note that the incidence rate in system (1.1) is bilinear, which regards the infection rate per density of infected individuals or per concentration of vibrios as a constant. Actually, the incidence rate is influenced by the inhibition effect from behavioral change of susceptible individuals and the crowding effect of vibrios. In [10], experimental studies indicated that the probability of infection depends on the concentration of vibrios in the contaminated water and only enough inoculum of vibrios can lead to cholera. In [11], Codeço introduced a new form $\beta B/(k + B)$ to measure the effect of saturation, where β measures the infection force of vibrios and B/(k + B) measures the inhibition effect and crowding effect. There have been several works on cholera models with saturation incidence in the literature (see, for example, [3, 4, 6, 12]).

Motivated by the works of Tien and Earn [2], Mukandavire et al. [9], and Codeço [11], in the present paper, we are concerned with the effects of hyperinfectious and hypoinfectious vibrios, both human-to-human and environment-to-human transmissions on the global dynamics of cholera. Besides, vaccination is widely used to prevent and control cholera, thus some susceptible individuals become vaccinated individuals. In [13], A.P. Lemos-Paião et al. proposed a cholera model including vaccination and illustrated the importance of vaccination campaigns on the control and eradication of

a cholera outbreak. To this end, we consider the following differential equations:

$$\begin{aligned} \frac{dS(t)}{dt} &= \mu N - \left(\beta_h I(t) + \frac{\beta_H B_H(t)}{k_H + B_H(t)} + \frac{\beta_L B_L(t)}{k_L + B_L(t)}\right) S(t) - (\mu + \phi) S(t), \\ \frac{dV(t)}{dt} &= \phi S(t) - \sigma \left(\beta_h I(t) + \frac{\beta_H B_H(t)}{k_H + B_H(t)} + \frac{\beta_L B_L(t)}{k_L + B_L(t)}\right) V(t) - \mu V(t), \\ \frac{dI(t)}{dt} &= \left(\beta_h I(t) + \frac{\beta_H B_H(t)}{k_H + B_H(t)} + \frac{\beta_L B_L(t)}{k_L + B_L(t)}\right) (S(t) + \sigma V(t)) - (\gamma + \mu) I(t), \\ \frac{dR(t)}{dt} &= \gamma I(t) - \mu R(t), \\ \frac{dB_H(t)}{dt} &= \xi I(t) - \chi B_H(t), \\ \frac{dB_L(t)}{dt} &= \chi B_H(t) - \delta_L B_L(t), \end{aligned}$$
(1.2)

where S(t), V(t), I(t), R(t), $B_H(t)$, $B_L(t)$ and other parameters are described in Table 1. Corresponding flowchart of cholera transmission in system (1.2) is depicted in Figure 1.



Figure 1. Flowchart of cholera transmission in system (1.2).

The initial condition for system (1.2) takes the form

$$S(0) = S^{0}, V(0) = V^{0}, I(0) = I^{0}, R(0) = R^{0}, B_{H}(0) = B_{H}^{0}, B_{L}(0) = B_{L}^{0},$$
(1.3)

where constants S^0 , V^0 , I^0 , R^0 , B^0_H , B^0_L are defined on $[0, \infty)$. It can be proved by the fundamental theory of functional differential equations [14] that system (1.2) has a unique solution $(S(t), V(t), I(t), R(t), B_H(t), B_L(t))$ satisfying the initial condition (1.3). It is easy to show that all solutions of system (1.2) with initial condition (1.3) are defined on $[0, +\infty)$ and remain positive for all $t \ge 0$, while, Ω is a positively invariant set for system (1.2), where

$$\Omega = \{(S, V, I, R, B_H, B_L) | S, V, I, R < N, B_H < \xi N / \chi, B_L < \xi N / \delta_L \},\$$

which implies that S(t), V(t), I(t), R(t), $B_H(t)$ and $B_L(t)$ are bounded in the invariant set Ω .

This paper is organized as follows. In Section 2, we calculate the basic reproduction number to system (1.2) and establish the existence of disease-free and endemic equilibria. In Section 3, the

local asymptotic stability of each of the equilibria is studied. In Section 4, we investigate the global asymptotic stability of each of the equilibria. In Section 5, we carry out a study of optimal control to seek cost-effective solutions of control strategies for cholera. In Section 6, we present numerical simulations to illustrate the theoretical results and describe the cholera outbreak in Haiti. Furthermore, we obtain the optimal control solution by the Forward-Backward Sweep Method. The paper ends with a conclusion in Section 7.

Variables	Description
S(t)	the density of susceptible individuals at time <i>t</i>
V(t)	the density of vaccinated individuals at time t
I(t)	the density of infected individuals at time t
R(t)	the density of recovered individuals at time t
$B_H(t)$	the concentration of hyperinfectious vibrios at time t
$B_L(t)$	the concentration of hypoinfectious vibrios at time t
Ν	the constant of total population
μ	birth and natural death rate
eta_h	transmission rate of infected individuals
eta_H	transmission rate of hyperinfectious vibrios
β_L	transmission rate of hypoinfectious vibrios
k_H	the concentration of hyperinfectious vibrios in contaminated water
	that yields 50% chance of catching cholera
k_L	the concentration of hypoinfectious vibrios in contaminated water
	that yields 50% chance of catching cholera
ϕ	vaccination rate of susceptible individuals
σ	reduction rate of vaccination due to the infection (less than one)
γ	recovery rate of infected individuals
ξ	the contribution rate of each infected individual to the concentration
	of V. cholerae
χ	the decay rate from hyperinfectious state to reduced infectiousness
δ_L	net death rate of V. cholerae

Table 1.	Definitions	of free	quently	used	variables.

2. Basic reproduction number and feasible equilibria

System (1.2) always has a disease-free equilibrium $E_0(S_0, V_0, 0, 0, 0, 0) = (\mu N/(\mu + \phi), \phi N/(\mu + \phi), 0, 0, 0, 0)$. By the method of van den Driessche and Watmough [15], the associated next generation matrices are given by

	$\frac{\beta_h(\mu N + \sigma \phi N)}{\mu + \phi}$	$\frac{\beta_H(\mu N + \sigma \phi N)}{k_H(\mu + \phi)}$	$\frac{\beta_L(\mu N + \sigma \phi N)}{k_L(\mu + \phi)}$	
F =	0	0	0	,
	0	0	0	

and

$$V = \left[\begin{array}{ccc} \gamma + \mu & 0 & 0 \\ -\xi & \chi & 0 \\ 0 & -\chi & \delta_L \end{array} \right].$$

Then, the basic reproduction number is determined by the spectral radius of FV^{-1} , namely,

$$\mathcal{R}_0 = \rho(FV^{-1}) = \left(\beta_h + \frac{\beta_H \xi}{\chi k_H} + \frac{\beta_L \xi}{\delta_L k_L}\right) \frac{(\mu + \sigma \phi) N}{(\mu + \phi) (\mu + \gamma)},$$

which represents the average number of new infections generated by a single newly infectious individual during the full infectious period. It can be shown that system (1.2) has an endemic equilibrium $E^*(S^*, V^*, I^*, R^*, B^*_H, B^*_L)$ if $\mathcal{R}_0 > 1$, which satisfies that

$$S^{*} = \mu N \left(\beta_{h} I^{*} + \frac{\beta_{H} \xi I^{*}}{\chi k_{H} + \xi I^{*}} + \frac{\beta_{L} \xi I^{*}}{\delta_{L} k_{L} + \xi I^{*}} + \mu + \phi \right)^{-1},$$

$$V^{*} = \phi S^{*} \left(\beta_{h} \sigma I^{*} + \frac{\beta_{H} \xi \sigma I^{*}}{\chi k_{H} + \xi I^{*}} + \frac{\beta_{L} \sigma \xi I^{*}}{\delta_{L} k_{L} + \xi I^{*}} + \mu \right)^{-1},$$

$$R^{*} = \frac{\gamma I^{*}}{\mu}, \quad B^{*}_{H} = \frac{\xi I^{*}}{\chi}, \quad B^{*}_{L} = \frac{\xi I^{*}}{\delta_{L}},$$
(2.1)

and I^* is the positive real root of the following equation:

$$s_1 I^{*6} + s_2 I^{*5} + s_3 I^{*4} + s_4 I^{*3} + s_5 I^{*2} + s_6 I^* + s_7 = 0,$$
(2.2)

where

$$\begin{split} s_{1} = &\sigma\xi^{4}(\mu + \gamma), \\ s_{2} = &2\sigma\xi^{3}(\mu + \gamma)\left(\beta_{H}\xi + \beta_{L}\xi + k_{H}\chi + k_{L}\delta_{L}\right) + \beta_{h}\sigma\xi^{4}(\mu + \gamma)(\mu + \phi) + \beta_{h}\mu\xi^{4}(\mu + \gamma) - \mu\sigma\xi^{4}N, \\ s_{3} = &2\sigma\xi^{2}(\mu + \gamma)\left(\beta_{L}\xi\chi k_{H} + \beta_{H}\xi\delta_{L}k_{L} + \beta_{h}\chi\delta_{L}k_{H}k_{L}\right) + \sigma\xi^{2}(\mu + \gamma)(\beta_{H}\xi + \beta_{L}\xi + \chi k_{H} + \delta_{L}k_{L})^{2} \\ &+ \xi^{3}(\mu + \gamma)\left(\beta_{h}\delta_{L}k_{L} + \beta_{h}\chi k_{H} + \beta_{H}\xi + \beta_{L}\xi\right)\left[\sigma(\mu + \phi) + \mu\right] \\ &+ \beta_{h}\xi^{3}(\mu + \gamma)(\delta_{L}k_{L} + \chi k_{H})\left[\sigma(\mu + \phi) + \mu\right] + \mu\xi^{4}(\mu + \gamma)(\mu + \phi) \\ &- \beta_{h}\mu\xi^{4}N(\mu + \sigma\phi) - 2\mu\sigma\xi^{3}N(\beta_{H}\xi + \beta_{L}\xi + \chi k_{H} + \delta_{L}k_{L}), \\ s_{4} = &2\sigma\xi(\mu + \gamma)(\beta_{H}\xi + \beta_{L}\xi + \chi k_{H} + \delta_{L}k_{L})(\beta_{L}\xi\chi k_{H} + \beta_{H}\xi\delta_{L}k_{L} + \beta_{h}\chi\delta_{L}k_{H}k_{L}) \\ &+ \xi^{2}(\mu + \gamma)(\beta_{h}\chi\delta_{L}k_{H}k_{L} + \beta_{H}\xi\delta_{L}k_{L} + \beta_{L}\xi\chi k_{H})\left[\sigma(\mu + \phi) + \mu\right] \\ &+ \xi^{2}(\mu + \gamma)(\delta_{L}k_{L} + \chi k_{H})(\beta_{h}\delta_{L}k_{L} + \beta_{h}\chi k_{H} + \beta_{H}\xi + \beta_{L}\xi)\left[\sigma(\mu + \phi) + \mu\right] \\ &+ \beta_{h}\delta_{L}\chi\xi^{2}k_{H}k_{L}(\mu + \gamma)\left[\sigma(\mu + \phi) + \mu\right] + 2\mu\xi^{3}(\mu + \gamma)(\mu + \phi)(\chi k_{H} + \delta_{L}k_{L}) \\ &- \beta_{h}\mu\xi^{3}N(\mu + \sigma\phi)(\chi k_{H} + \delta_{L}k_{L})^{2} - \mu\xi^{3}N(\mu + \sigma\phi)(\beta_{h}\delta_{L}k_{L} + \beta_{h}\chi \delta_{L}k_{H}k_{L})^{2} \\ &- \mu\sigma\xi^{2}N(\beta_{H}\xi + \beta_{L}\xi + \chi k_{H} + \delta_{L}k_{L})^{2} - \mu\xi^{3}N(\mu + \sigma\phi)(\beta_{h}\delta_{L}k_{L} + \beta_{h}\chi k_{H} + \beta_{H}\xi + \beta_{L}\xi)\left[\sigma(\mu + \phi) + \mu\right] \\ &+ \xi\delta_{L}\chi k_{H}k_{L}(\mu + \gamma)(\beta_{h}\delta_{L}k_{L} + \beta_{h}\chi \delta_{L}k_{H}k_{L})^{2} \\ &+ \xi\delta_{L}\chi k_{H}k_{L}(\mu + \gamma)(\beta_{h}\delta_{L}k_{L} + \beta_{h}\chi \delta_{L}k_{H}k_{L})^{2} \\ &+ \xi\delta_{L}\chi k_{H}k_{L}(\mu + \gamma)(\beta_{h}\delta_{L}k_{L} + \beta_{h}\chi \delta_{L}k_{H}k_{L})^{2} \\ &+ \xi\delta_{L}\chi k_{H}k_{L}(\mu + \gamma)(\beta_{h}\delta_{L}k_{L} + \beta_{h}\chi \delta_{L}k_{H}k_{L})^{2} \\ &+ \xi\delta_{L}\chi k_{H} + \beta_{H}\xi\xi\delta_{L}k_{L} + \beta_{h}\chi \delta_{L}k_{H}k_{L}) \end{aligned}$$

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$$\times \begin{bmatrix} \xi(\mu+\gamma)(\delta_{L}k_{L}+\chi k_{H}) \left[\sigma(\mu+\phi)+\mu\right] \\ -\mu\xi^{2}N\left(\mu+\sigma\phi\right)-2\mu\sigma\xi N\left(\beta_{H}\xi+\beta_{L}\xi+k_{H}\chi+k_{L}\delta_{L}\right) \end{bmatrix} \\ -\mu\xi^{2}N\left(\mu+\sigma\phi\right)\left(\delta_{L}k_{L}+\chi k_{H}\right)\left(\beta_{h}\delta_{L}k_{L}+\beta_{h}\chi k_{H}+\beta_{H}\xi+\beta_{L}\xi\right) \\ -\beta_{h}\mu\chi\delta_{L}\xi^{2}k_{L}k_{H}N\left(\mu+\sigma\phi\right)+\mu\xi^{2}(\mu+\gamma)(\mu+\phi)\left(\delta_{L}^{2}k_{L}^{2}+4\chi\delta_{L}k_{H}k_{L}+\chi^{2}k_{H}^{2}\right), \\ s_{6} =\delta_{L}\chi k_{H}k_{L}(\mu+\gamma)\left(\beta_{h}\chi k_{H}\delta_{L}k_{L}+\beta_{H}\xi\delta_{L}k_{L}+\beta_{L}\xi\chi k_{H}\right)\left[\sigma(\mu+\phi)+\mu\right] \\ +2\chi\mu\delta_{L}\xi k_{H}k_{L}(\mu+\gamma)(\mu+\phi)\left(\chi k_{H}+\delta_{L}k_{L}\right) \\ -\mu\xi N\left(\mu+\sigma\phi\right)\left(\delta_{L}k_{L}+\chi k_{H}\right)\left(\beta_{h}\chi k_{H}\delta_{L}k_{L}+\beta_{H}\xi\delta_{L}k_{L}+\beta_{L}\xi\chi k_{H}\right) \\ -\mu\chi\delta_{L}\xi k_{H}k_{L}N\left(\mu+\sigma\phi\right)\left(\beta_{h}\delta_{L}k_{L}+\beta_{h}\chi k_{H}+\beta_{H}\xi+\beta_{L}\xi\right) \\ -\mu\sigma N(\beta_{L}k_{H}\xi\chi+\beta_{H}k_{L}\xi\delta_{L}+\beta_{h}k_{H}k_{L}\chi\delta_{L})^{2}, \\ s_{7} = k_{H}^{2}k_{L}^{2}\mu\chi^{2}\delta_{L}^{2}(\mu+\phi)(\mu+\gamma)\left(1-\mathcal{R}_{0}\right).$$

Denote the left side of (2.2) by $P(I^*)$. Note that $\lim_{I^* \to +\infty} P(I^*) = +\infty$ and $P(0) = s_7 < 0$ if $\mathcal{R}_0 > 1$, thus system (1.2) has a positive equilibrium E^* .

3. Local asymptotic stability

In this section, we are concerned with the local asymptotic stability of disease-free and endemic equilibria. The approach of proofs is to analyze the distribution of roots for corresponding characteristic equations.

Theorem 3.1. If $\mathcal{R}_0 < 1$, the disease-free equilibrium E_0 of system (1.2) is locally asymptotically stable; if $\mathcal{R}_0 > 1$, E_0 is unstable.

Proof. The characteristic equation of system (1.2) at E_0 is

$$(\lambda + \chi) (\lambda + \delta_L) (\lambda + \mu)^2 (\lambda + \mu + \phi) (\lambda + \mu + \gamma) = \frac{(\mu + \sigma\phi) N}{\mu + \phi} (\lambda + \mu)^2 (\lambda + \mu + \phi) \left[\beta_h (\lambda + \chi) (\lambda + \delta_L) + \frac{\beta_H \xi}{k_H} (\lambda + \delta_L) + \frac{\beta_L \chi \xi}{k_L} \right].$$
(3.1)

It is clear that (3.1) has negative real roots $\lambda = -\mu$, $\lambda = -(\mu + \phi)$ and other roots are determined by the following equation:

$$(\lambda + \chi)(\lambda + \delta_L)(\lambda + \mu + \gamma) - \frac{(\mu + \sigma\phi)N}{\mu + \phi} \left[\beta_h(\lambda + \chi)(\lambda + \delta_L) + \frac{\beta_H\xi}{k_H}(\lambda + \delta_L) + \frac{\beta_L\chi\xi}{k_L} \right] = 0.$$
(3.2)

Denote $\mathcal{R}_0 = \mathcal{R}_{01} + \mathcal{R}_{02} + \mathcal{R}_{03}$, where

$$\mathcal{R}_{01} = \frac{\beta_h \left(\mu + \sigma\phi\right) N}{\left(\mu + \phi\right) \left(\mu + \gamma\right)}, \quad \mathcal{R}_{02} = \frac{\beta_H \xi \left(\mu + \sigma\phi\right) N}{\chi k_H \left(\mu + \phi\right) \left(\mu + \gamma\right)}, \quad \mathcal{R}_{03} = \frac{\beta_L \xi \left(\mu + \sigma\phi\right) N}{\delta_L k_L \left(\mu + \phi\right) \left(\mu + \gamma\right)}.$$

Substituting $\mathcal{R}_0, \mathcal{R}_{01}, \mathcal{R}_{02}$ into (3.2) yields

$$\left(\frac{\lambda^2}{\delta_L\chi} + \frac{\lambda}{\delta_L} + \frac{\lambda}{\chi} + 1\right) \left(\frac{\lambda}{\mu + \gamma} + 1\right) = \left[\mathcal{R}_{01}\frac{\lambda^2}{\delta_L\chi} + (\mathcal{R}_{01} + \mathcal{R}_{02})\frac{\lambda}{\delta_L} + \mathcal{R}_{01}\frac{\lambda}{\chi} + \mathcal{R}_0\right].$$
(3.3)

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Now, we claim that all roots of (3.3) have negative real parts. Otherwise, system (1.2) has a root $\lambda_1 = x_1 + iy_1$ with $x_1 \ge 0$. In this case, if $\mathcal{R}_0 < 1$, it is easy to see that

$$\left|\frac{\lambda_1^2}{\delta_L \chi} + \frac{\lambda_1}{\delta_L} + \frac{\lambda_1}{\chi} + 1\right| > \left|\mathcal{R}_{01} \frac{\lambda_1^2}{\delta_L \chi} + (\mathcal{R}_{01} + \mathcal{R}_{02}) \frac{\lambda_1}{\delta_L} + \mathcal{R}_{01} \frac{\lambda_1}{\chi} + \mathcal{R}_{0}\right|, \quad \left|\frac{\lambda_1}{\mu + \gamma} + 1\right| > 1$$

It follows that

$$\left| \left(\frac{\lambda_1^2}{\delta_L \chi} + \frac{\lambda_1}{\delta_L} + \frac{\lambda_1}{\chi} + 1 \right) \left(\frac{\lambda_1}{\mu + \gamma} + 1 \right) \right| > \left| \mathcal{R}_{01} \frac{\lambda_1^2}{\delta_L \chi} + \left(\mathcal{R}_{01} + \mathcal{R}_{02} \right) \frac{\lambda_1}{\delta_L} + \mathcal{R}_{01} \frac{\lambda_1}{\chi} + \mathcal{R}_{0} \right|,$$

which contradicts to (3.3). Therefore, if $\mathcal{R}_0 < 1$, all roots of (3.1) have negative real parts and E_0 is locally asymptotically stable. If $\mathcal{R}_0 > 1$, we denote the left side of (3.1) by $G(\lambda)$:

$$G(\lambda) = (\lambda + \chi) (\lambda + \delta_L) (\lambda + \mu)^2 (\lambda + \mu + \phi) (\lambda + \gamma + \mu) - \frac{(\mu + \sigma\phi)N}{\mu + \phi} (\lambda + \mu)^2 (\lambda + \mu + \phi) \left[\beta_h (\lambda + \chi) (\lambda + \delta_L) + \frac{\beta_H \xi}{k_H} (\lambda + \delta_L) + \frac{\beta_L \chi \xi}{k_L} \right],$$

where $G(0) = \delta_L \chi \mu^2 (\mu + \phi)(\gamma + \mu)(1 - \mathcal{R}_0) < 0$ and $G(\lambda) \to \infty$ as $\lambda \to \infty$. Noting that $G(\lambda)$ is a continuous function with respect to λ , if $\mathcal{R}_0 > 1$, Eq. (3.1) has a positive real root, then E_0 is unstable.

Theorem 3.2. If $\mathcal{R}_0 > 1$, the endemic equilibrium E^* of system (1.2) is locally asymptotically stable.

Proof. The characteristic equation of system (1.2) at E^* is

$$(\lambda + \mu) (\lambda + \chi) (\lambda + \delta_L) (\lambda + \mu + \gamma) (\lambda + \mu + \phi + XI^*) (\lambda + \mu + \sigma XI^*)$$

= $(\lambda + \mu)^2 (\lambda + \chi) (\lambda + \delta_L) [S^* (\lambda + \mu + \phi + \sigma XI^*) + \sigma V^* (\lambda + \mu + \phi + XI^*)] Y,$ (3.4)

where

$$\begin{split} X &= \beta_h + \frac{\beta_H \xi}{\chi k_H + \xi I^*} + \frac{\beta_L \xi}{\delta_L k_L + \xi I^*}, \\ Y &= \beta_h + \frac{\beta_H \xi}{\chi k_H + \xi I^*} \frac{\chi k_H}{\chi k_H + \xi I^*} \frac{\chi}{\lambda + \chi} + \frac{\beta_L \xi}{\delta_L k_L + \xi I^*} \frac{\delta_L k_L}{\delta_L k_L + \xi I^*} \frac{\chi}{\lambda + \chi} \frac{\delta_L}{\lambda + \delta_L}. \end{split}$$

It is obvious that (3.4) has a negative real root $\lambda = -\mu$ and other roots are determined by the following equation:

$$(\lambda + \chi) (\lambda + \delta_L) (\lambda + \mu + \gamma) (\lambda + \mu + \phi + XI^*) (\lambda + \mu + \sigma XI^*) = (\lambda + \chi) (\lambda + \delta_L) (\lambda + \mu) [S^* (\lambda + \mu + \phi + \sigma XI^*) + \sigma V^* (\lambda + \mu + \phi + XI^*)] Y.$$
(3.5)

For the sake of contradiction, let $\lambda_2 = x_2 + iy_2$ with $x_2 \ge 0$. Noting that $X(S^* + \sigma V^*) = \mu + \gamma$ and $\sigma < 1$, we have

$$\begin{aligned} &|(\lambda_{2} + \mu + \gamma) (\lambda_{2} + \mu + \phi + XI^{*})| \\ &> |X (S^{*} + \sigma V^{*}) (\lambda_{2} + \mu + \phi + XI^{*})| \\ &> |X [S^{*} (\lambda_{2} + \mu + \phi + \sigma XI^{*}) + \sigma V^{*} (\lambda_{2} + \mu + \phi + XI^{*})]| \\ &> |Y [S^{*} (\lambda_{2} + \mu + \phi + \sigma XI^{*}) + \sigma V^{*} (\lambda_{2} + \mu + \phi + XI^{*})]|. \end{aligned}$$
(3.6)

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Hence, we obtain that

$$\begin{aligned} &|(\lambda + \chi)(\lambda + \delta_L)(\lambda_2 + \mu + \gamma)(\lambda_2 + \mu + \phi + XI^*)(\lambda_2 + \mu + \sigma XI^*)| \\ &> \left|(\lambda + \chi)(\lambda + \delta_L)(\lambda_2 + \mu)\left[S^*(\lambda_2 + \mu + \phi + \sigma XI^*) + \sigma V^*(\lambda_2 + \mu + \phi + XI^*)\right]Y\right|, \end{aligned}$$

which contradicts to (3.5). Therefore, if $\mathcal{R}_0 > 1$, all roots of (3.4) have negative real parts and E^* is locally asymptotically stable.

4. Global asymptotic stability

In this section, we study the global stability of each of the equilibria to system (1.2). The approach of proofs is to use suitable Lyapunov functionals and LaSalle's invariance principle. Since the variable R(t) does not appear explicitly in the first three and last two equations in system (1.2), we don't need to consider it in later analysis.

Theorem 4.1. If $\mathcal{R}_0 < 1$, the disease-free equilibrium E_0 of system (1.2) is globally asymptotically stable.

Proof. Let $(S(t), V(t), I(t), B_H(t), B_L(t))$ be any positive solution of system (1.2) with initial condition (1.3). Define

$$V_1(t) = S_0 \left(\frac{S(t)}{S_0} - 1 - \ln \frac{S(t)}{S_0} \right) + V_0 \left(\frac{V(t)}{V_0} - 1 - \ln \frac{V(t)}{V_0} \right) + I(t) + c_1 B_H(t) + c_2 B_L(t),$$

where constants c_1 and c_2 will be determined later. Calculating the derivative of $V_1(t)$ along positive solutions of system (1.2) yields

$$\begin{split} \dot{V}_{1}(t) &= \left(1 - \frac{S_{0}}{S(t)}\right) \left[\mu N - \left(\beta_{h}I(t) + \frac{\beta_{H}B_{H}(t)}{k_{H} + B_{H}(t)} + \frac{\beta_{L}B_{L}(t)}{k_{L} + B_{L}(t)}\right) S(t) - (\mu + \phi)S(t) \right] \\ &+ \left(1 - \frac{V_{0}}{V(t)}\right) \left[\phi S(t) - \sigma \left(\beta_{h}I(t) + \frac{\beta_{H}B_{H}(t)}{k_{H} + B_{H}(t)} + \frac{\beta_{L}B_{L}(t)}{k_{L} + B_{L}(t)}\right) V(t) - \mu V(t) \right] \\ &+ \left(\beta_{h}I(t) + \frac{\beta_{H}B_{H}(t)}{k_{H} + B_{H}(t)} + \frac{\beta_{L}B_{L}(t)}{k_{L} + B_{L}(t)}\right) (S(t) + \sigma V(t)) - (\gamma + \mu)I(t) \\ &+ c_{1} \left(\xi I(t) - \chi B_{H}(t)\right) + c_{2} \left(\chi B_{H}(t) - \delta_{L}B_{L}(t)\right). \end{split}$$
(4.1)

Substituting $S_0 = \mu N/(\mu + \phi)$, $V_0 = \phi N/(\mu + \phi)$ into (4.1), we obtain that

$$\dot{V}_{1}(t) = \frac{\mu^{2}N}{\mu + \phi} \left(2 - \frac{S(t)}{S_{0}} - \frac{S_{0}}{S(t)} \right) + \frac{\mu\phi N}{\mu + \phi} \left(3 - \frac{S_{0}}{S(t)} - \frac{V(t)}{V_{0}} - \frac{S(t)V_{0}}{S_{0}V(t)} \right) \\ + \left[\frac{\beta_{h}N(\mu + \sigma\phi)}{\mu + \phi} + c_{1}\xi - (\gamma + \mu) \right] I(t) + \frac{\beta_{L}(\mu + \sigma\phi)NB_{L}(t)}{(\mu + \phi)(k_{L} + B_{L}(t))} - c_{2}\delta_{L}B_{L}(t)$$

$$+ \frac{\beta_{H}(\mu + \sigma\phi)NB_{H}(t)}{(\mu + \phi)(k_{H} + B_{H}(t))} + c_{2}\chi B_{H}(t) - c_{1}\chi B_{H}(t).$$

$$(4.2)$$

Choose

$$c_1 = \frac{N(\mu + \sigma\phi)}{\mu + \phi} \left(\frac{\beta_H}{\chi k_H} + \frac{\beta_L}{\delta_L k_L} \right), \quad c_2 = \frac{\beta_L N(\mu + \sigma\phi)}{\delta_L k_L(\mu + \phi)}$$

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From (4.2), we have

$$\begin{split} \dot{V}_{1}(t) &= \frac{\mu^{2}N}{\mu + \phi} \left(2 - \frac{S(t)}{S_{0}} - \frac{S_{0}}{S(t)} \right) + \frac{\mu\phi N}{\mu + \phi} \left(3 - \frac{S_{0}}{S(t)} - \frac{V(t)}{V_{0}} - \frac{S(t)V_{0}}{S_{0}V(t)} \right) \\ &- \frac{\beta_{H}N(\mu + \sigma\phi)B_{H}^{2}(t)}{k_{H}(\mu + \phi)(k_{H} + B_{H}(t))} - \frac{\beta_{L}N(\mu + \sigma\phi)B_{L}^{2}(t)}{k_{L}(\mu + \phi)(k_{L} + B_{L}(t))} \\ &+ (\mathcal{R}_{0} - 1)(\gamma + \mu)I(t). \end{split}$$

It follows from the inequality of arithmetic and geometric means that $\dot{V}_1(t) \leq 0$ with equality holding if and only if $S = S_0$, $V = V_0$, $I = B_H = B_L = 0$. It can be verified that $M_0 = \{E_0\} \subset \Omega$ is the largest invariant subset of $\{(S(t), V(t), I(t), B_H(t), B_L(t)) : \dot{V}_1(t) = 0\}$. Noting that if $\mathcal{R}_0 < 1$, E_0 is locally asymptotically stable, thus we obtain the global asymptotic stability of E_0 from LaSalle's invariance principle.

Theorem 4.2. If $\mathcal{R}_0 > 1$, the endemic equilibrium E^* of system (1.2) is globally asymptotically stable.

Proof. Let $(S(t), V(t), I(t), B_H(t), B_L(t))$ be any positive solution of system (1.2) with initial condition (1.3). Define

$$\begin{split} V_2(t) = S^* \left(\frac{S(t)}{S^*} - 1 - \ln \frac{S(t)}{S^*} \right) + V^* \left(\frac{V(t)}{V^*} - 1 - \ln \frac{V(t)}{V^*} \right) + I^* \left(\frac{I(t)}{I^*} - 1 - \ln \frac{I(t)}{I^*} \right) \\ + l_1 B_H^* \left(\frac{B_H(t)}{B_H^*} - 1 - \ln \frac{B_H(t)}{B_H^*} \right) + l_2 B_L^* \left(\frac{B_L(t)}{B_L^*} - 1 - \ln \frac{B_L(t)}{B_L^*} \right), \end{split}$$

where constants l_1 and l_2 will be determined later. Calculating the derivative of $V_2(t)$ along positive solutions of system (1.2) yields

$$\begin{split} \dot{V}_{2}(t) &= \left(1 - \frac{S^{*}}{S(t)}\right) \left[\mu N - \left(\beta_{h}I(t) + \frac{\beta_{H}B_{H}(t)}{k_{H} + B_{H}(t)} + \frac{\beta_{L}B_{L}(t)}{k_{L} + B_{L}(t)}\right) S(t) - (\mu + \phi)S(t) \right] \\ &+ \left(1 - \frac{V^{*}}{V(t)}\right) \left[\phi S(t) - \sigma \left(\beta_{h}I(t) + \frac{\beta_{H}B_{H}(t)}{k_{H} + B_{H}(t)} + \frac{\beta_{L}B_{L}(t)}{k_{L} + B_{L}(t)}\right) V(t) - \mu V(t) \right] \\ &+ \left(1 - \frac{I^{*}}{I(t)}\right) \left[\left(\beta_{h}I(t) + \frac{\beta_{H}B_{H}(t)}{k_{H} + B_{H}(t)} + \frac{\beta_{L}B_{L}(t)}{k_{L} + B_{L}(t)}\right) (S(t) + \sigma V(t)) - (\gamma + \mu)I(t) \right] \\ &+ l_{1} \left(1 - \frac{B^{*}_{H}}{B_{H}(t)}\right) (\xi I(t) - \chi B_{H}(t)) + l_{2} \left(1 - \frac{B^{*}_{L}}{B_{L}(t)}\right) (\chi B_{H}(t) - \delta_{L}B_{L}(t)) \,. \end{split}$$

$$(4.3)$$

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Substituting (2.1) into (4.3) yields

$$\begin{split} \dot{V}_{2}(t) = S^{*} \left(\mu + \beta_{h}I^{*}\right) \left(2 - \frac{S^{*}}{S(t)} - \frac{S(t)}{S^{*}}\right) + V^{*} \left(\mu + \beta_{h}\sigma I^{*}\right) \left(3 - \frac{S^{*}}{S(t)} - \frac{S(t)V^{*}}{S^{*}V(t)} - \frac{V(t)}{V^{*}}\right) \\ &+ \frac{\beta_{H}S^{*}B_{H}^{*}}{k_{H} + B_{H}^{*}} \left[2 - \frac{S^{*}}{S(t)} - \frac{S(t)I^{*}B_{H}(t)\left(k_{H} + B_{H}^{*}\right)}{S^{*}I(t)B_{H}^{*}\left(k_{H} + B_{H}(t)\right)}\right] \\ &+ \frac{\beta_{L}S^{*}B_{L}^{*}}{k_{L} + B_{L}^{*}} \left[2 - \frac{S^{*}}{S(t)} - \frac{S(t)I^{*}B_{L}(t)\left(k_{L} + B_{L}^{*}\right)}{S^{*}I(t)B_{L}^{*}\left(k_{L} + B_{L}(t)\right)}\right] \\ &+ \frac{\beta_{H}\sigma V^{*}B_{H}^{*}}{k_{H} + B_{H}^{*}} \left[3 - \frac{S^{*}}{S(t)} - \frac{S(t)V^{*}}{S^{*}V(t)} - \frac{V(t)I^{*}B_{H}(t)\left(k_{H} + B_{H}^{*}\right)}{V^{*}I(t)B_{H}^{*}\left(k_{H} + B_{H}(t)\right)}\right] \\ &+ \frac{\beta_{L}\sigma V^{*}B_{L}^{*}}{k_{L} + B_{L}^{*}} \left[3 - \frac{S^{*}}{S(t)} - \frac{S(t)V^{*}}{S^{*}V(t)} - \frac{V(t)I^{*}B_{L}(t)\left(k_{L} + B_{L}^{*}\right)}{V^{*}I(t)B_{L}^{*}\left(k_{L} + B_{L}(t)\right)}\right] \\ &+ \left[l_{1}\chi B_{H}^{*} - (S^{*} + \sigma V^{*})\left(\frac{\beta_{L}B_{L}^{*}}{k_{L} + B_{L}^{*}} + \frac{\beta_{H}B_{H}^{*}}{k_{H} + B_{H}^{*}}\right)\right]\frac{I(t)}{I^{*}} \\ &+ (S^{*} + \sigma V^{*})\frac{\beta_{H}B_{H}(t)}{k_{H} + B_{H}(t)} - l_{1}\chi B_{H}(t) - l_{1}\chi B_{H}^{*}(t) + l_{2}\delta_{L}B_{L}^{*}B_{H}^{*}(t) \\ &+ (S^{*} + \sigma V^{*})\frac{\beta_{L}B_{L}(t)}{k_{L} + B_{L}(t)} - l_{2}\delta_{L}B_{L}(t) + l_{2}\delta_{L}B_{L}^{*}\frac{B_{H}(t)}{B_{H}^{*}} - l_{2}\delta_{L}B_{L}^{*}\frac{B_{L}^{*}B_{H}(t)}{B_{L}(t)B_{H}^{*}}. \end{split}$$

Choose

$$l_{1} = \frac{(S^{*} + \sigma V^{*})}{\chi B_{H}^{*}} \left(\frac{\beta_{L} B_{L}^{*}}{k_{L} + B_{L}^{*}} + \frac{\beta_{H} B_{H}^{*}}{k_{H} + B_{H}^{*}} \right), \quad l_{2} = \frac{(S^{*} + \sigma V^{*})}{\delta_{L} B_{L}^{*}} \frac{\beta_{L} B_{L}^{*}}{k_{L} + B_{L}^{*}}.$$

Direct calculation shows that

$$\begin{split} \dot{V}_{2}(t) &= S^{*} \left(\mu + \beta_{h}I^{*}\right) \left(2 - \frac{S^{*}}{S(t)} - \frac{S(t)}{S^{*}}\right) + V^{*} \left(\mu + \beta_{h}\sigma I^{*}\right) \left(3 - \frac{S^{*}}{S(t)} - \frac{S(t)V^{*}}{S^{*}V(t)} - \frac{V(t)}{V^{*}}\right) \\ &+ \frac{\beta_{H}S^{*}B_{H}^{*}}{k_{H} + B_{H}^{*}} \left[4 - \frac{S^{*}}{S(t)} - \frac{I(t)B_{H}^{*}}{I^{*}B_{H}(t)} - \frac{k_{H} + B_{H}(t)}{k_{H} + B_{H}^{*}} - \frac{S(t)I^{*}B_{H}(t)\left(k_{H} + B_{H}^{*}\right)}{S^{*}I(t)B_{H}^{*}\left(k_{H} + B_{H}(t)\right)}\right] \\ &+ \frac{\beta_{L}S^{*}B_{L}^{*}}{k_{L} + B_{L}^{*}} \left[5 - \frac{S^{*}}{S(t)} - \frac{I(t)B_{H}^{*}}{I^{*}B_{H}(t)} - \frac{B_{L}^{*}B_{H}(t)}{B_{L}(t)B_{H}^{*}} - \frac{k_{L} + B_{L}(t)}{k_{L} + B_{L}^{*}} - \frac{S(t)I^{*}B_{L}(t)\left(k_{L} + B_{L}^{*}\right)}{S^{*}I(t)B_{L}^{*}\left(k_{L} + B_{L}(t)\right)}\right] \\ &+ \frac{\beta_{H}\sigma V^{*}B_{H}^{*}}{k_{H} + B_{H}^{*}} \left[5 - \frac{S^{*}}{S(t)} - \frac{S(t)V^{*}}{S^{*}V(t)} - \frac{I(t)B_{H}^{*}}{I^{*}B_{H}(t)} - \frac{k_{H} + B_{H}(t)}{k_{H} + B_{H}^{*}} - \frac{V(t)I^{*}B_{H}(t)\left(k_{H} + B_{H}^{*}\right)}{V^{*}I(t)B_{H}^{*}\left(k_{H} + B_{H}(t)\right)}\right] \\ &+ \frac{\beta_{L}\sigma V^{*}B_{L}^{*}}{k_{L} + B_{L}^{*}} \left[6 - \frac{S^{*}}{S(t)} - \frac{S(t)V^{*}}{S^{*}V(t)} - \frac{I(t)B_{H}^{*}}{I^{*}B_{H}(t)} - \frac{B_{L}^{*}B_{H}(t)}{B_{L}(t)B_{H}^{*}} - \frac{k_{L} + B_{L}(t)}{k_{L} + B_{L}^{*}} - \frac{V(t)I^{*}B_{L}(t)\left(k_{L} + B_{H}^{*}\right)}{V^{*}I(t)B_{L}^{*}\left(k_{L} + B_{L}^{*}\right)}\right] \\ &- \frac{\beta_{H}k_{H}\left(S^{*} + \sigma V^{*}\right)\left(B_{H}(t) - B_{H}^{*}\right)^{2}}{\left(k_{H} + B_{H}(t)\right)} - \frac{\beta_{L}k_{L}\left(S^{*} + \sigma V^{*}\right)\left(B_{L}(t) - B_{L}^{*}\right)^{2}}{\left(k_{L} + B_{L}^{*}\right)^{2}\left(k_{L} + B_{L}(t)\right)}. \end{split}$$

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Thus, it follows from the inequality of arithmetic and geometric means that $\dot{V}_2(t) \leq 0$ with equality holding if and only if $S = S^*$, $V = V^*$, $I = I^*$, $B_H = B_H^*$, $B_L = B_L^*$. It can be proved that $M^* = \{E^*\} \subset \Omega$ is the largest invariant subset of $\{(S(t), V(t), I(t), B_H(t), B_L(t)) : \dot{V}_2(t) = 0\}$. Noting that if $\mathcal{R}_0 > 1$, E^* is locally asymptotically stable, we therefore obtain the global asymptotic stability of E^* from LaSalle's invariance principle.

5. Optimal control strategies

Optimal control methods help us find cost-effective strategies to control various kinds of diseases. We aim to minimize the number of infected individuals and the corresponding cost of the strategies during the course of an epidemic. Control strategies, such as quarantine, vaccination, treatment, and sanitation, can realize the control of cholera at different cost.

Let $X = (S, V, I, R, B_H, B_L)$. Define a control function set as $U = \{u_i | i = 1, 2, 3, 4\}$, where the meanings of u_i are listed as follows:

(1) u_1 is a quarantine strategy which is used to reduce the contact between susceptible individuals and infected individuals;

(2) u_2 is a vaccination strategy that can improve the immunocompetence of susceptible individuals;

(3) u_3 is a treatment strategy that aims at increasing the recovery rate of infected individuals;

(4) u_4 is a sanitation strategy that aims at killing vibrios in contaminated water.

Due to the limitation of medical technology or cost, each of control strategies u_i has upper bound u_{imax} . The model with control strategies is given by the following system of nonlinear ordinary differential equations:

$$\begin{aligned} \frac{dS(t)}{dt} &= \mu N - \left[(1 - u_1(t))\beta_h I(t) + \frac{\beta_H B_H(t)}{k_H + B_H(t)} + \frac{\beta_L B_L(t)}{k_L + B_L(t)} \right] S(t) - (\mu + u_2(t))S(t), \\ \frac{dV(t)}{dt} &= u_2(t)S(t) - \sigma \left[(1 - u_1(t))\beta_h I(t) + \frac{\beta_H B_H(t)}{k_H + B_H(t)} + \frac{\beta_L B_L(t)}{k_L + B_L(t)} \right] V(t) - \mu V(t), \\ \frac{dI(t)}{dt} &= \left[(1 - u_1(t))\beta_h I(t) + \frac{\beta_H B_H(t)}{k_H + B_H(t)} + \frac{\beta_L B_L(t)}{k_L + B_L(t)} \right] (S(t) + \sigma V(t)) - (u_3(t) + \mu) I(t), \\ \frac{dR(t)}{dt} &= u_3(t)I(t) - \mu R(t), \\ \frac{dB_H(t)}{dt} &= \xi I(t) - (u_4(t) + \chi) B_H(t), \end{aligned}$$
(5.1)

 $\frac{dB_L(t)}{dt} = \chi B_H(t) - (u_4(t) + \delta_L) B_L(t).$ The set X of admissible trajectories is given by

$$\mathcal{X} = \left\{ X(\cdot) \in W^{1,1}\left([0,T]; \mathbb{R}^6\right) | (1.3) \text{ and } (5.1) \text{ are satisfied} \right\},\$$

and the admissible control set \mathcal{U} is given by

$$\mathcal{U} = \left\{ U(\cdot) \in L^{\infty} \left([0,T]; \mathbb{R}^4 \right) | 0 < u_i \le u_{imax} < 1, \forall t \in [0,T] \right\}.$$

From [16, 17], we consider the objective functional

$$J(X(\cdot), U(\cdot)) = \int_0^T \left(AI + \frac{B_1}{2}u_1^2 + \frac{B_2}{2}u_2^2 + \frac{B_3}{2}u_3^2 + \frac{B_4}{2}u_4^2 \right) dt.$$

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Thereinto, *A* denotes the weight constant of the infected individuals, while, B_1 , B_2 , B_3 and B_4 are the weight constants for the control strategies. $B_1u_1^2/2$, $B_2u_2^2/2$, $B_3u_3^2/2$ and $B_4u_4^2/2$ describe the cost associated with quarantine, vaccination, treatment, and sanitation strategies, respectively. The square of the control variables shows the severity of the side effects of the four strategies [17]. Our aim is to minimize the number of infected individuals and corresponding cost of the strategies. The optimal control problem consists of determining the vector function $X^{\circ}(\cdot) = (S^{\circ}(\cdot), V^{\circ}(\cdot), I^{\circ}(\cdot), R^{\circ}(\cdot), B_{H}^{\circ}(\cdot), B_{L}^{\circ}(\cdot)) \in X$ associated with an admissible control $U^{\circ}(\cdot) \in \mathcal{U}$ on the time interval [0, T], minimizing the objective functional *J*, i.e.,

$$J(X^{\diamond}(\cdot), U^{\diamond}(\cdot)) = \min_{X(\cdot), U(\cdot) \in \mathcal{X} \times \mathcal{U}} J(X(\cdot), U(\cdot)).$$
(5.2)

The existence of an optimal control $U(\cdot)$ comes from the convexity of the objective functional J with respect to the controls and the regularity of the system (5.1): see, e.g., [18, 19].

According to the Pontryagin Maximum Principle [20], if $U(\cdot) \in X$ is optimal for problem (5.2) with fixed final time *T*, then there exists a nontrivial absolutely continuous mapping $\lambda : [0, T] \to \mathbb{R}^6$, $\lambda(t) = (\lambda_1(t), \lambda_2(t), \lambda_3(t), \lambda_4(t), \lambda_5(t), \lambda_6(t))$, called the adjoint vector, such that

(1) the control system:

$$S' = \frac{\partial H}{\partial \lambda_1}, \quad V' = \frac{\partial H}{\partial \lambda_2}, \quad I' = \frac{\partial H}{\partial \lambda_3}, \quad R' = \frac{\partial H}{\partial \lambda_4}, \quad B_H' = \frac{\partial H}{\partial \lambda_5}, \quad B_L' = \frac{\partial H}{\partial \lambda_6};$$

(2) the adjoint system:

$$\lambda_1' = -\frac{\partial H}{\partial S}, \quad \lambda_2' = -\frac{\partial H}{\partial V}, \quad \lambda_3' = -\frac{\partial H}{\partial I}, \quad \lambda_4' = -\frac{\partial H}{\partial R}, \quad \lambda_5' = -\frac{\partial H}{\partial B_H}, \quad \lambda_6' = -\frac{\partial H}{\partial B_L};$$

(3) the minimization condition:

$$H(X^{\diamond}(t), U^{\diamond}(t), \lambda^{\diamond}(t)) = \min_{0 < u_i \le u_{i\max}} H(X^{\diamond}(t), U(t), \lambda^{\diamond}(t)),$$

(4) hold for almost all $t \in [0, T]$, where the function *H*, defined by

$$\begin{split} H(X, U, \lambda) = &AI + \frac{B_1}{2}u_1^2 + \frac{B_2}{2}u_2^2 + \frac{B_3}{2}u_3^2 + \frac{B_4}{2}u_4^2 \\ &+ \lambda_1 \left\{ \mu N - \left[(1 - u_1)\beta_h I + \frac{\beta_H B_H}{k_H + B_H} + \frac{\beta_L B_L}{k_L + B_L} \right] S - (\mu + u_2) S \right\} \\ &+ \lambda_2 \left\{ u_2 S - \sigma \left[(1 - u_1)\beta_h I + \frac{\beta_H B_H}{k_H + B_H} + \frac{\beta_L B_L}{k_L + B_L} \right] V - \mu V \right\} \\ &+ \lambda_3 \left\{ \left[(1 - u_1)\beta_h I + \frac{\beta_H B_H}{k_H + B_H} + \frac{\beta_L B_L}{k_L + B_L} \right] (S + \sigma V) - (u_3 + \mu) I \right\} \\ &+ \lambda_4 (u_3 I - \mu R) + \lambda_5 \left[\xi I - (u_4 + \chi) B_H \right] \\ &+ \lambda_6 \left[\chi B_H - (u_4 + \delta_L) B_L \right], \end{split}$$

is called the Hamiltonian.

(5) Moreover, the following transversality conditions also hold:

$$\lambda_i(T) = 0, \quad i = 1, \dots, 6.$$

From [18, 19, 20, 21], it is not difficult to show the following theorem.

Theorem 5.1. The optimal control problem (5.2) with fixed final time T admits a unique optimal solution $(S^{\circ}(\cdot), V^{\circ}(\cdot), I^{\circ}(\cdot), R^{\circ}(\cdot), B_{H}^{\circ}(\cdot), B_{L}^{\circ}(\cdot))$ associated with an optimal control U(t) for $t \in [0, T]$. Moreover, there exist adjoint functions $\lambda_i^{\circ}(\cdot)$ (i = 1, ..., 6) such that

$$\begin{aligned} \frac{d\lambda_{1}^{\circ}}{dt} &= \lambda_{1}^{\circ} \left(\mu + u_{2}\right) + \lambda_{1}^{\circ} \left[(1 - u_{1})\beta_{h}I^{\circ} + \frac{\beta_{H}B_{H}^{\circ}}{k_{H} + B_{H}^{\circ}} + \frac{\beta_{L}B_{L}^{\circ}}{k_{L} + B_{L}^{\circ}} \right] \\ &\quad - \lambda_{2}^{\circ}u_{2} - \lambda_{3}^{\circ} \left[(1 - u_{1})\beta_{h}I^{\circ} + \frac{\beta_{H}B_{H}^{\circ}}{k_{H} + B_{H}^{\circ}} + \frac{\beta_{L}B_{L}^{\circ}}{k_{L} + B_{L}^{\circ}} \right] , \\ \frac{d\lambda_{2}^{\circ}}{dt} &= \lambda_{2}^{\circ}\sigma \left[(1 - u_{1})\beta_{h}I^{\circ} + \frac{\beta_{H}B_{H}^{\circ}}{k_{H} + B_{H}^{\circ}} + \frac{\beta_{L}B_{L}^{\circ}}{k_{L} + B_{L}^{\circ}} \right] + \lambda_{2}^{\circ}\mu \\ &\quad - \lambda_{3}^{\circ}\sigma \left[(1 - u_{1})\beta_{h}I^{\circ} + \frac{\beta_{H}B_{H}^{\circ}}{k_{H} + B_{H}^{\circ}} + \frac{\beta_{L}B_{L}^{\circ}}{k_{L} + B_{L}^{\circ}} \right] , \\ \frac{d\lambda_{3}^{\circ}}{dt} &= -A + \lambda_{1}^{\circ} \left(1 - u_{1} \right)\beta_{h}S^{\circ} + \lambda_{2}^{\circ} \left(1 - u_{1} \right)\beta_{h}\sigma V^{\circ} \\ &\quad - \lambda_{3}^{\circ}\beta_{h} \left(1 - u_{1} \right) \left(S^{\circ} + \sigma V^{\circ} \right) + \lambda_{3}^{\circ} \left(u_{3} + \mu \right) - \lambda_{4}^{\circ}u_{3} - \lambda_{5}^{\circ}\xi, \\ \frac{d\lambda_{4}^{\circ}}{dt} &= \lambda_{4}^{\circ}\beta_{H}S^{\circ} \frac{k_{H}}{\left(k_{H} + B_{H}^{\circ}\right)^{2}} + \lambda_{2}^{\circ}\beta_{H}\sigma V^{\circ} \frac{k_{H}}{\left(k_{H} + B_{H}^{\circ}\right)^{2}} \\ &\quad - \lambda_{3}^{\circ}\beta_{H} \left(S^{\circ} + \sigma V^{\circ} \right) \frac{k_{H}}{\left(k_{H} + B_{H}^{\circ}\right)^{2}} + \lambda_{5}^{\circ} \left(u_{4} + \chi \right) - \lambda_{6}^{\circ}\chi, \\ \frac{d\lambda_{6}^{\circ}}{dt} &= \lambda_{1}^{\circ}\beta_{L}S^{\circ} \frac{k_{L}}{\left(k_{L} + B_{L}^{\circ}\right)^{2}} + \lambda_{2}^{\circ}\beta_{L}\sigma V^{\circ} \frac{k_{L}}{\left(k_{L} + B_{L}^{\circ}\right)^{2}} + \lambda_{6}^{\circ} \left(u_{4} + \delta_{L} \right), \\ &\quad - \lambda_{3}^{\circ}\beta_{L} \left(S^{\circ} + \sigma V^{\circ} \right) \frac{k_{L}}{\left(k_{L} + B_{L}^{\circ}\right)^{2}} + \lambda_{6}^{\circ} \left(u_{4} + \delta_{L} \right), \end{aligned}$$

with transversality conditions

$$\lambda_i^{\diamond}(T) = 0, \quad i = 1, \dots, 6.$$

Furthermore,

$$u_i^{\diamond} = \max[0, \min(\tilde{u}_i, u_{i\max})], \tag{5.4}$$

where

$$\begin{split} \tilde{u}_1 &= \frac{\left(\lambda_3^{\circ} - \lambda_1^{\circ}\right)\beta_h S^{\circ} I^{\circ} + \left(\lambda_3^{\circ} - \lambda_2^{\circ}\right)\beta_h \sigma V^{\circ} I^{\circ}}{B_1}, \\ \tilde{u}_2 &= \frac{\left(\lambda_1^{\circ} - \lambda_2^{\circ}\right)S^{\circ}}{B_2}, \quad \tilde{u}_3 = \frac{\left(\lambda_3^{\circ} - \lambda_4^{\circ}\right)I^{\circ}}{B_3}, \quad \tilde{u}_4 = \frac{\lambda_5^{\circ} B_H^{\circ} + \lambda_6^{\circ} B_L^{\circ}}{B_4}. \end{split}$$

In the next section, we solve u_i^{\diamond} by a numerical method. Numerical techniques for optimal control problems can often be classified as either direct or indirect. In terms of disease control, indirect methods, such as the Forward-Backward Sweep Method, approximate solutions to optimal control problems by numerically solving the boundary value problem for the differential-algebraic system generated by the Maximum Principle [22].

6. Numerical simulations

In this section, we want to illustrate the theoretical results for system (1.2) by numerical simulations. Besides, we transform system (1.2) to simulate the cholera outbreak in Haiti. Furthermore, by Forwardbackward Sweep Method, we obtain the optimal control strategies and show the graph trajectories of infected individuals as well as hyperinfectious and hypoinfectious vibrios with optimal control and without optimal control.

6.1. Dynamical behaviors of the model

Following [8, 11, 24, 25, 26], we choose appropriate parameter values and simulate each of the equilibria, respectively.

Variables	Case 1	Case 2	Source
Ν	10,000	10,000	assumed
μ	0.023/year	0.023/year	[24]
eta_h	0.00005/day	0.00011/day	[24]
eta_H	0.0075/day	0.0075/day	[8]
eta_L	0.00012/day	0.00012/day	[8]
k_H	2.86×10^3 cells/ml	1.43×10^3 cells/ml	[11]
k_L	2×10^{6} cells/ml	10 ⁶ cells/ml	[26]
ϕ	70%	70%	[24]
σ	25%	25%	[24]
γ	0.5/day	0.2/day	[25]
ξ	10 cells/ml/day	10 cells/ml/day	[25]
χ	0.8/h	0.2/h	[26]
δ_L	0.132/day	0.033/day	[24]

Table 2. List of parameters.

Case 1: By simple computing, we obtain that $\mathcal{R}_0 = 0.4335 < 1$. From Theorem 3.1, we derive that disease-free equilibrium $E_0(318.12, 9681.88, 0, 0, 0, 0)$ is locally asymptotically stable. Figure 2 shows the results of simulations using parameters from Case 1 in Table 2. The six figures plot the time evolution of the six variables S(t), V(t), I(t), R(t), $B_H(t)$ and $B_L(t)$, respectively. In fact, we can observe in Figure 2 that infected and recovered individuals, as well as hyperinfectious and hypoinfectious vibrios, die out.

Case 2: Similarly, we derive that $\mathcal{R}_0 = 4.5718 > 1$. From Theorem 3.2, we obtain that endemic equilibrium $E^*(297.01, 5809.58, 401.57, 3491.84, 20078.04, 121685.53)$ is locally asymptotically stable. Figure 3 shows the results of simulations using parameters from Case 2 in Table 2. Each figure plots the time evolution of the corresponding variable, too. From Figure 3, we see that infected and recovered individuals, as well as hyperinfectious and hypoinfectious vibrios, persist at the endemic level.



Figure 2. The graph trajectories of S(t), V(t), I(t), R(t), $B_H(t)$ and $B_L(t)$ versus t of system (1.2) where $\mathcal{R}_0 = 0.4335 < 1$.



Figure 3. The graph trajectories of S(t), V(t), I(t), R(t), $B_H(t)$ and $B_L(t)$ versus *t* of system (1.2) where $\mathcal{R}_0 = 4.5718 > 1$.

6.2. Simulations of the cholera outbreak in Haiti

The first cholera cases in Haiti started to be reported on 14 October 2010 in the department of Artibonite from where the outbreak rapidly spread along the Artibonite river affecting several departments. In the beginning of November 2010, the overall case fatality rate (CFR) in hospitals was 3.8%. This high CFR reflects the lack of experience of the healthcare system to deal with case management as well as the fact that patients reached the health facilities too late [28]. In Figure 4, weekly reported cholera cases in Haiti from 1 November 2010 to 1 May 2011 are plotted in blue line.

To simulate this cholera outbreak, we remove the vaccinated individuals from system (1.2) and carry out the following subsystem without vaccination:

$$\frac{dS(t)}{dt} = \mu N - \left(\beta_h I(t) + \frac{\beta_H B_H(t)}{k_H + B_H(t)} + \frac{\beta_L B_L(t)}{k_L + B_L(t)}\right) S(t) - \mu S(t),
\frac{dI(t)}{dt} = \left(\beta_h I(t) + \frac{\beta_H B_H(t)}{k_H + B_H(t)} + \frac{\beta_L B_L(t)}{k_L + B_L(t)}\right) S(t) - (\gamma + \mu) I(t),
\frac{dR(t)}{dt} = \gamma I(t) - \mu R(t),
\frac{dB_H(t)}{dt} = \xi I(t) - \chi B_H(t),
\frac{dB_L(t)}{dt} = \chi B_H(t) - \delta_L B_L(t).$$
(6.1)

Meanwhile, considering the medical treatment and public health of Haiti, we decrease the recovery rate of infected individuals from 0.2/day to 0.05/day. The constant of total population N is adjusted to 50000. From Figure 4, we observe that the numerical simulation of infected individuals in subsystem (6.1) (red dashed curve) is approximately in accord with the real cholera cases in Haiti (blue line). Once taking the vaccination control strategy, as we can see, the numerical simulation of infected individuals in system (1.2) (green curve) decreases quickly to a lower level.



Figure 4. Cholera cases in Haiti (blue line), numerical simulations of infected individuals in subsystem (6.1) (red dashed curve) and in system (1.2) (green curve), where the related parameter values are the same as Case 2 in Table 2 except for the recovery rate and the total population.

6.3. Optimal control solution

The idea exploited by Forward-Backward Sweep Method is that the initial value problem of the state equation is solved forward in time, using an estimate for the control and costate variables, then the costate final value problem is solved backward in time (see, for example, [12, 17, 23]). In [27], Lenhart

and Workman employed the Forward-Backward Sweep Method combined with progressive-regressive Runge-Kutta fourth-order schemes to get the optimal solution. Based on Lenhart and Workman's method, our numerical simulations are performed by using a MATLAB code. The MATLAB function *linspace* creates n + 1 = 101 equally spaced nodes between 0 and 1, including 0 and 1. Here, $\vec{x} = (x_1, \ldots, x_{n+1})$ and $\vec{\lambda} = (\lambda_1, \ldots, \lambda_{n+1})$ are the vector approximations for the state and adjoint. The related parameter values in systems (5.1) and (5.3) are the same as Case 1 in Table 2. A rough outline of the algorithm is given below.

(1) Make an initial guess for \vec{u} over the interval;

(2) Using the initial condition x(0) and the values for \vec{u} , solve \vec{x} forward in time according to system (5.1). Specifically, given a step size *h* and an ODE x'(t) = f(t, x(t)), the approximation of x(t+h) given x(t) is

$$x(t+h) \approx x(t) + \frac{h}{6} \left(k_1 + 2k_2 + 2k_3 + k_4 \right), \tag{6.2}$$

where

$$k_1 = f(t, x(t)), \ k_2 = f\left(t + \frac{h}{2}, x(t) + \frac{h}{2}k_1\right), \ k_3 = f\left(t + \frac{h}{2}, x(t) + \frac{h}{2}k_2\right), \ k_4 = f\left(t + h, x(t) + hk_3\right).$$

(3) Using the transversality condition $\lambda_{n+1} = \lambda(T) = 0$ and the values for \vec{u} and \vec{x} , solve $\vec{\lambda}$ backward in time according to system (5.3) by a similar way as (6.2);

(4) Update \vec{u} by entering the new \vec{x} and $\vec{\lambda}$ values into the characterization of the optimal control (5.4);

(5) Check convergence. If variable values in this iteration and the last iteration are negligibly close, output the current values as solutions. If values are not close, return to Step 2.

When all steps are complete, we obtain the optimal control strategies, which can be seen in Figure 5(a). On account of medical technology and cost, each of control strategies has limitation, thus we set $u_{1 \text{ max}} = 0.8$, $u_{2 \text{ max}} = 0.7$, $u_{3 \text{ max}} = 0.6$ and $u_{4 \text{ max}} = 0.9$. We observe that $u_2(t)$, namely, vaccination strategy, could be reduced 82 days later from the beginning of the cholera outbreak, which saves much cost of vaccination. Similarly, After 27 days, $u_4(t)$, namely, sanitation strategy, is not much necessary and could be canceled gradually. In Figure 5(b)–Figure 5(d), we compare the graph trajectories of infected individuals as well as hyperinfectious and hypoinfectious vibrios with optimal control. It is clear that infected individuals, as well as hyperinfectious and hypoinfectious vibrios, have been reduced due to the control strategies. From Figure 5, we suggest that taking control strategies at the very beginning of cholera outbreaks can reduce the number of infected individuals remarkably, which are also cost-effective optimal strategies.

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Figure 5. Figure 5(a) is the graph trajectories of four optimal control strategies; Figure 5(b)– Figure 5(d) are the graph trajectories of I(t), $B_H(t)$, $B_L(t)$ with optimal control and without optimal control.

7. Discussion

In this paper, we have considered a cholera model including hyperinfectious and hypoinfectious vibrios, both human-to-human and environment-to-human transmissions. By a complete mathematical analysis, the threshold dynamics of the model was established and it can be fully determined by the basic reproduction number. If $\mathcal{R}_0 < 1$, the disease-free equilibrium E_0 is locally and globally asymptotically stable; if $\mathcal{R}_0 > 1$, the endemic equilibrium E^* is locally and globally asymptotically stable. Numerical simulations vividly illustrate our main results of stability analysis for system (1.2). Besides, we simulated the cholera outbreak in Haiti. Furthermore, we obtained the optimal solution by the Forward-Backward Sweep Method.

At the beginning of cholera epidemic, hyperinfectious vibrios freshly-shed from infected individuals play an important role in cholera transmission, due to that they are likely to come into contact with other individuals [7]. Therefore, the strategies of quarantine and sanitation can effectively control the cholera epidemic. Besides, vaccination strategy is still the most efficient control strategy to prevent, control and eradicate cholera. As for treatment strategy, it is an essential method to fight cholera and reduce the death rate due to the disease. In contrast to the control strategies in [12, 29], Modnak or Lemos-Pai \tilde{a} o et al. only considered a single control method, vaccination or quarantine. In a word, a

combination with all the above strategies could yield the best control effect of cholera.

With the development of economy and society, the rate of population movement has accelerated in recent years. In view of this, considering the heterogeneity of each individual is an important factor in constructing more realistic models, that is, spatially heterogeneous epidemic models. The theoretical analysis of these models may be more complicated and we leave it for further investigation.

Acknowledgments

This work was supported by the National Natural Science Foundation of China (Nos. 11871316, 11801340, 11371368), the Natural Science Foundation of Shanxi Province (Nos. 201801D121006, 201801D221007).

Conflict of interest

The authors declare that they have no competing interests.

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