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Dynamical analysis and control strategies in modelling Ebola virus disease

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Abstract

Ebola virus disease (EVD) is a severe infection with an extremely high fatality rate spread through direct and indirect contacts. Recently, an outbreak of EVD in West Africa brought public attention to this deadly disease. We study the spread of EVD through a two-patch model. We determine the basic reproduction number, the disease-free equilibrium, two boundary equilibria and the endemic equilibrium when the disease persists in the two sub-populations for specific conditions. Further, we introduce time-dependent controls into our proposed model. We analyse the optimal control problem where the control system is a mathematical model for EVD that incorporates educational campaigns. The control functions represent educational campaigns in their respective patches, with one patch having more effective controls than the other. We aim to study how these control measures would be implemented for a certain time period, in order to reduce or eliminate EVD in the respective communities, while minimising the intervention implementation costs. Numerical simulations results are provided to illustrate the dynamics of the disease in the presence of controls.

Keywords: Ebola; Reproduction number; Sensitivity analysis; Educational campaign; Optimal control

1 Introduction

Ebola virus disease (EVD) is a highly infectious disease caused by the Ebola virus from the family of *Filoviridae* viruses. The Ebola virus is believed to be found in mammals of the family of Pteropodidae (fruit bats) [1]. It is a virus having a filamentous, enveloped and a non-segmented negative sense with the entry of the virus to living cells facilitated by its enveloped glycoprotein [2, 3]. The current Ebola virus disease (EVD) outbreak in West Africa is now the largest documented [4]. Ebola originated in Zaire and Sudan in 1976 with several strains known at present. The outbreak in 2015 affected mainly Guinea, Liberia and Sierra Leone (West Africa) causing more than 28 000 infections and over 11 100 deaths by December 2015 [5]. The epidemic continued increasing due to socio-economic disadvantage and shortages in the health systems of the three mainly affected countries (Guinea, Liberia and Sierra Leone) [4, 6]. Currently, in 2018 the EVD has resurfaced in the Province of Kivu in the Democratic Republic of Congo (DRC). In Kivu, there is a military conflict and there are thousands of displaced refugees [7, 8]. The affected area has a lot of trade among the different communities within, and this leads to massive inter-community movements. Due to the running battles in the Province of Kivu, health

care workers and those who might be able to assist in the fight against EVD find it difficult to reach all the communities [9]. Thus, the respective communities in the affected areas of the DRC have greatly varying intervention strategies and some are not perfect. In the Province of Kivu, rebels are against the presence of the health caregivers and UN peacekeepers, hence EVD might be difficult to curtail in the Democratic Republic of Congo [9, 10].

Partially eaten fruits and pulp dropped by the bats are then eaten by the land mammals such as monkeys, apes, baboons and gorillas. This chain of events then forms a possible transmission of the Ebola virus from bats to other mammals [1]. Infection with humans can occur through direct contact with blood or body fluids (like saliva, sweat, faeces, breast milk and semen), objects like needles and syringes that have been contaminated with the virus and infected fruit bats or other mammals [11, 12]. The survival of the virus in the environment, due to poor hygienic and sanitary conditions, is probably another source of Ebola infection in many places in Africa [13, 14]. In Africa, and particularly in the regions that were affected by Ebola outbreaks, people who live close to the rain-forests, hunt bats and monkeys and harvest forest fruits for food [15, 16]. Africans are sincere to the point that even a transmittable disease would not stop them from showing compassion for their relatives at home, caressing them and shaking hands, as this is part of their beliefs and customs. In addition, in the course of the funerals, they bath and clothe up their lifeless relatives [17]. They even share without appropriate washing the clothes of their lifeless relatives. Thus, the virus can be spread directly or indirectly.

Without effective vaccines and treatment, educational campaigns may be implemented as a countermeasure. One of the primary reasons for the spread of the virus is the poorly functioning health systems in the part of Africa where the disease occurs. Other reasons are illiteracy, poverty and lack of enough information on the mode of spread of the virus [18]. People who care for infected individuals have an increased risk of transmission. Recommended measures when caring for the infected include medical isolation through proper use of gloves, masks, gowns, boots and goggles as well as sterilisation of equipment and surfaces. Certain control strategies are employed when local, regional and international associates are informed of an establishment of a possible Ebola epidemic. The controls among them are and not restricted to the following; assessment of the worldwide health risk, establishment of social mobilisation and health education curriculum to listen and address public concerns. The application of such control procedures leads to the curtailing of the current EVD epidemic of 2014, 2015. Due to the weaker economies of the affected countries, it was a limiting factor in the fight against EVD [17]. Resource allocation needs to be optimal and the control strategies need to be implemented in such a way as to derive maximum benefits. It is worth noting that optimal control is best described by mathematical modelling. Optimal control theory has proven to be a successful tool in understanding ways to curtail the spread of infectious diseases by devising the optimal disease intervention strategies. The method consists of minimising the cost of infection or the cost of implementing the control, or both. For more on optimal control theory in epidemiology, we refer the reader to Malik and Sharomi, 2015 [19].

Mathematical modelling remains a powerful tool in describing the dynamics of disease outbreaks and making predictions. It also enables us to estimate the long term effects of interventions, integrates evidence from different scientific disciplines and investigates how

health-related practices evolve in complex systems. Statistical and biological studies have been applied in the study of the complexity of the Ebola virus life ecology [20, 21], and are worthy of attention. Some dynamic models have been proposed to study and to try to understand the dynamics of EVD [22–30]. Recently, Berge *et al.* [29] proposed a deterministic mathematical model for the transmission dynamics of EVD which involved a synergy between the epizootic phase, enzootic phase and the endemic phase. It included the direct and indirect modes of contamination, between and within the three different types of populations consisting of humans, animals and fruit bats. Work done by Berge *et al.* [29] was an extension of their previous work done in 2015 [26], in which a simple mathematical model was developed, which incorporated both the direct and the indirect Ebola virus transmission in such a way that there is a provision of Ebola viruses. Models have also been developed to try and understand various intervention strategies in trying to curtail the spread of EVD [31–40]. The impact of vaccination and vaccines was investigated in [35, 36, 39, 40] and the issue of quarantining analysed in [35, 37] through mathematical models. Further, Berge *et al.* [31] developed a mathematical model to understand the impact of contact tracing as a control strategy of EVD. In recent times, optimal control problems have generated a lot of interest from researchers. Furthermore, various techniques and intervention strategies have been applied to study optimal control problems related to EVD [41–46]. In particular, Area *et al.* [44] proposed an EVD mathematical model with the vaccination of the susceptible population, with the aim of controlling the spread of the disease and analysis of two optimal control problems related to the transmission of EVD with vaccination. A deterministic SEIR type model with additional hospitalisation, quarantine and vaccination was developed and studied by Ahmad *et al.* [43], to understand the disease dynamics.

Although substantial work has been done on the study of EVD transmission dynamics, not much consideration of meta-population study has been done; see [47, 48]. The population is subdivided into several discrete patches which are supposed to be well mixed. Then, in each patch, the population is subdivided into compartments corresponding to different epidemic status. For more on meta-population patch models, see [49]. In this work, motivated by the usefulness of and the current investigation on modelling EVD, we intend to systematically investigate the modelling and analysis of a two-patch model for the transmission dynamics of EVD. We make use of the model studied by Berge *et al.* [26] which takes into account both, the direct and indirect modes of transmission. Thus, in our model, we distinguish two patches. We assume the susceptible and the recovered individuals are the only ones who migrate, hence the two sub-populations can vary. This is consistent with the fact that in African refugee camps refugees move very little between two camps and only persons that bring them help move within the camps. These limited migrations on the susceptible individuals have taken place in places such as the Democratic Republic of Congo [9, 10]. We also support our assumption that the majority of the infectious individuals do not migrate, due to their state of health. This can also be supported by the fact that in periods of epidemics, countries tighten people's movements and their respective borders through screening and detection. Both the people who are allowed to cross a border and those who move between two refugee camps/communities are assumed to be susceptible and recovered. A mathematical model describing the optimal control of an epidemic with educational campaigns have been discussed before [50],

hence in our two-patch model, we introduce two time-dependent controls in educational campaigns, which have different effectiveness.

The paper is structured as follows. The EVD transmission model is formulated in Sect. 2 and the analytical results of the model are presented in Sect. 3. In Sect. 4 optimal control theory has been applied to the model formulated in Sect. 2. Simulation results and projection profiles of EVD are presented in Sect. 5. A summary and concluding remarks complete the paper.

2 Model formulation

We consider a two-patch model for Ebola virus disease (EVD). The model consists of two sub-populations of a large one. The recruitment in each sub-population is only in the susceptible class, and the migration between the sub-populations is by the susceptible and recovered. Having the infectious class not migrating, we assume that infection does not take place during the migration process. Each human sub-population is divided into four classes: the susceptibles $S_i(t)$, infectious $I_i(t)$, recovered $R_i(t)$ and deceased human individuals $D_i(t)$. The Ebola virus in the environment is denoted by $W_i(t)$. For all the parameters and compartmental classes, we will consider the case for $i = 1, 2$, representing patch 1 and 2, respectively. The susceptible human population is replenished by constant recruitment at rate b_i . Death for a reason that is not related to EVD is proportional to the population size and with a constant rate μ_i . The additional death due to disease affects only the class $I_i(t)$, at a rate ν_i .

It has been shown that human individuals can be infected through the contaminated environment [51]. Thus, the Ebola virus is transmitted though both direct and indirect transmission. The susceptible individuals acquire the infection at rate Λ_i with

$$\Lambda_i = \beta_{I_i} I_i + \beta_{D_i} D_i + \beta_{W_i} W_i, \quad i = 1, 2, \quad (1)$$

the form of incidence is bilinear. $\beta_{I_i} I_i + \beta_{D_i} D_i$ is the direct infection from the infectious individuals and the dead bodies. β_{W_i} is the indirect infection from the Ebola virus in the environment to the susceptible individuals. β_{I_i} is the effective contact rate of the infectious individuals, β_{D_i} is the effective contact rate of the deceased individuals and β_{W_i} is the effective contact rate of the Ebola virus (in the environment).

Motivated by the fact that it has never been reported that an individual caught EVD for the second time, we assume that the infectious individuals recover at rate ϕ_i attaining permanent disease-induced immunity. We assume that the Ebola virus finds its way into the environment only through the infectious and the deceased human individuals, which they shed into the environment especially through the urine and stool at rates δ_i and ρ_i , respectively. This assumption is supported by several references [52–54]. This happens in regions of poor sanitary facilities and/or in the regions where people do not observe appropriate hygiene practices. The Ebola virus in the environment decays and loses its infectiousness at a rate r_i . The deceased human individuals lose their infectiousness through methods such as proper handling and burial at a rate α_i . The migration rates of the susceptible and the recovered individuals, between the two populations, are a_i and b_i for $i = 1, 2$

respectively. This yields the following set of differential equations:

$$\begin{cases} S'_i = b_i + a_j S_j - \Lambda_i S_i - (\mu_i + a_i) S_i, \\ I'_i = \Lambda_i S_i - (\mu_i + \nu_i + \phi_i) I_i, \\ R'_i = \phi_i I_i + b_j R_j - (\mu_i + b_i) R_i, \\ D'_i = \nu_i I_i - \alpha_i D_i, \\ W'_i = \delta_i I_i + \rho_i D_i - r_i W_i, \quad i = 1, 2; j = 1, 2; i \neq j. \end{cases} \tag{2}$$

The total human population, the total deceased human individuals and the Ebola virus pathogens in the environment are given by $N(t) = S_1(t) + I_1(t) + R_1(t) + S_2(t) + I_2(t) + R_2(t)$, $D(t) = D_1(t) + D_2(t)$ and $W(t) = W_1(t) + W_2(t)$, respectively.

Once $S_i, I_i, D_i, W_i, i = 1, 2$ are obtained from Eqs. (2)₁, (2)₂, (2)₄, (2)₅, the functions $R_i, i = 1, 2$ are readily given by Eq. (2)₃. Thus without loss of generality, we are led to the following reduced system:

$$\begin{cases} S'_i = b_i + a_j S_j - \Lambda_i S_i - (\mu_i + a_i) S_i, \\ I'_i = \Lambda_i S_i - (\mu_i + \nu_i + \phi_i) I_i, \\ D'_i = \nu_i I_i - \alpha_i D_i, \\ W'_i = \delta_i I_i + \rho_i D_i - r_i W_i, \quad i = 1, 2; j = 1, 2; i \neq j. \end{cases} \tag{3}$$

2.1 Basic properties of the model

In this section, we study the basic properties of the solutions of system (3) which are essential in the proofs of stability.

Theorem 1 *Let the initial data be $S_i(0) > 0, I_i(0) > 0, D_i(0) > 0, W_i(0) > 0, i = 1, 2$. Then the solutions $S_i(t), I_i(t), D_i(t), W_i(t)$ for system (3) are non-negative for all $t > 0$.*

A complete proof for Theorem 1 has been outlined in Appendix 1.

Theorem 2 *System (3) is a dynamical system on the biologically feasible compact set*

$$\mathcal{G} = \left\{ S_1(t), I_1(t), D_1(t), W_1(t), S_2(t), I_2(t), D_2(t), W_2(t) : \right. \\ \left. N \leq \frac{b}{\mu_0}, D \leq \frac{\bar{\nu}b}{\underline{\alpha}\mu_0}, W \leq \frac{\bar{\alpha}\bar{\delta}b + \bar{\rho}b\bar{\nu}}{\underline{\alpha}\mu_0\underline{r}} \right\}, \tag{4}$$

where $b = b_1 + b_2, \mu_0 = \min(\mu_1, \mu_2), \bar{\nu} = \max(\nu_1, \nu_2), \bar{\delta} = \max(\delta_1, \delta_2), \bar{\rho} = \max(\rho_1, \rho_2), \bar{\alpha} = \max(\alpha_1, \alpha_2), \underline{\alpha} = \min(\alpha_1, \alpha_2), \underline{r} = \min(r_1, r_2)$.

A complete proof for Theorem 2 has been outlined in Appendix 2.

3 Analysis of the model

3.1 The disease-free equilibrium and basic reproduction number

The global behaviour for this model crucially depends on the basic reproduction number, that is, the average number of secondary cases produced by a single infective individual

which is introduced into an entirely susceptible population. System (3) has an evident equilibrium $\mathcal{E}^0 = (S_1^0, 0, 0, 0, 0, S_2^0, 0, 0, 0, 0)$ when there is no disease, where

$$\begin{cases} S_1^0 = \frac{(\mu_2 + a_2)b_1 + a_2b_2}{\mu_1\mu_2 + \mu_1a_2 + \mu_2a_1}, \\ S_2^0 = \frac{(\mu_1 + a_1)b_2 + a_1b_1}{\mu_1\mu_2 + \mu_1a_2 + \mu_2a_1}. \end{cases} \tag{5}$$

We calculate the basic reproduction number, \mathcal{R}_0 , using the next generation method approach developed in van den Driessche and Watmough [55]. Following [55], the non-negative matrix F and the non-singular matrix V for the new infection terms and the remaining transfer terms are respectively given at the disease-free equilibrium by

$$F = \begin{bmatrix} \beta_{I_1}S_1^0 & \beta_{D_1}S_1^0 & \beta_{W_1}S_1^0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & \beta_{I_2}S_2^0 & \beta_{D_2}S_2^0 & \beta_{W_2}S_2^0 \\ 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 \end{bmatrix} \quad \text{and}$$

$$V = \begin{bmatrix} k_1 & 0 & 0 & 0 & 0 & 0 \\ -v_1 & \alpha_1 & 0 & 0 & 0 & 0 \\ -\delta_1 & -\rho_1 & r_1 & 0 & 0 & 0 \\ 0 & 0 & 0 & k_2 & 0 & 0 \\ 0 & 0 & 0 & -v_2 & \alpha_2 & 0 \\ 0 & 0 & 0 & -\delta_2 & -\rho_2 & r_2 \end{bmatrix}. \tag{6}$$

Then the reproduction number \mathcal{R}_0 of system (3) is the spectral radius of the next generation matrix FV^{-1} ,

$$\mathcal{R}_0 = \rho(FV^{-1}) = \max\{\mathcal{R}_1, \mathcal{R}_2\}, \tag{7}$$

where \mathcal{R}_i ($i = 1, 2$) is the basic reproduction number of each sub-population i given by

$$\begin{aligned} \mathcal{R}_i &= \mathcal{R}_{I_i} + \mathcal{R}_{D_i} + \mathcal{R}_{W_i} \\ &= \frac{S_i^0 \beta_{D_i} v_i}{k_i \alpha_i} + \frac{S_i^0 \beta_{I_i}}{k_i} + \frac{S_i^0 \beta_{W_i} (\alpha_i \delta_i + \mu_i \rho_i)}{(k_i r_i \alpha_i)}. \end{aligned} \tag{8}$$

Following the interpretation in [56], \mathcal{R}_i for patch i , for $i = 1, 2$, is the sum of three infection contributions:

- \mathcal{R}_{I_i} is the contribution of the infectious human I_i .
- \mathcal{R}_{D_i} is the contribution of the infected corpses D_i .
- \mathcal{R}_{W_i} is the contribution due to the environmental contamination by the virus W_i .

Biologically \mathcal{R}_1 and \mathcal{R}_2 measure the average number of secondary infections generated by an Ebola virus in patch 1 and patch 2 respectively, when introduced in a disease-free population. The reproductive number \mathcal{R}_0 controls the number of equilibria of system (3).

If $\mathcal{R}_0 \leq 1$, then the disease-free equilibrium \mathcal{E}^0 is the only equilibrium in \mathcal{G} . Using Theorem 2 in van den Driessche and Watmough (2002) [55], the following result is established.

Theorem 3 *The disease-free equilibrium (DFE) \mathcal{E}^0 of system (3) is locally asymptotically stable (LAS) if $\mathcal{R}_0 < 1$ and unstable otherwise.*

We now utilise the approach of Lyapunov functions [57–59] in the analysis of the global asymptotic stability.

Theorem 4 *If $\mathcal{R}_0 \leq 1$, the DFE is globally asymptotically stable (GAS) in \mathcal{G} . If $\mathcal{R}_0 > 1$, the system is uniformly persistent.*

Proof Let $\mathcal{Y}(t) = (I_1, D_1, W_1, I_2, D_2, W_2)$. Since

$$\begin{cases} I_i' \leq \Lambda_i S_i^0 - (\mu_i + \nu_i + \phi_i)I_i, \\ D_i' \leq (\mu_i + \nu_i)I_i - \rho_i D_i, \\ W_i' \leq \delta_i I_i + \rho_i D_i - r_i W_i, \quad i = 1, 2, \end{cases} \tag{9}$$

it follows that

$$\dot{\mathcal{Y}}(t) \leq (F - V)\mathcal{Y}, \tag{10}$$

where F and V are defined in (6). It is worth noting that F and V^{-1} are non-negative. By the Perron–Frobenius theorem [60], the non-negative matrix $V^{-1}F$ has a non-negative left eigenvector $u \geq 0$ with respect to $\rho(V^{-1}F) = \rho(FV^{-1}) = \mathcal{R}_0$, that is, $u^T V^{-1}F = \mathcal{R}_0 u^T$. Motivated by [59], we consider the Lyapunov function

$$\mathcal{L} = u^T V^{-1}\mathcal{Y}. \tag{11}$$

Differentiating \mathcal{L} along solutions of (3), we have

$$\begin{aligned} \dot{\mathcal{L}}(t) &= u^T V^{-1}\dot{\mathcal{Y}} \\ &\leq u^T V^{-1}(F - V)\mathcal{Y} \\ &= (\mathcal{R}_0 - 1)u^T \mathcal{Y} \leq 0, \quad \text{if } \mathcal{R}_0 \leq 1. \end{aligned} \tag{12}$$

It can be easily verified that the largest invariant subset of \mathcal{G} where $\dot{\mathcal{L}} = 0$ is the singleton $\{\mathcal{E}^0\}$. Therefore, by LaSalle’s invariance principle [61], \mathcal{E}^0 is globally asymptotically stable in \mathcal{G} when $\mathcal{R}_0 \leq 1$.

If $\mathcal{R}_0 > 1$, then, by continuity, $\dot{\mathcal{L}} > 0$ in a neighbourhood of \mathcal{E}^0 in the interior of \mathcal{G} . Solutions in the interior of \mathcal{G} sufficiently close to \mathcal{E}^0 move away from the DFE, implying that the DFE is unstable. This completes the proof. \square

The result in Theorem 4 shows that $\mathcal{R}_0 = 1$ is a sharp threshold for disease dynamics: the disease will die out when $\mathcal{R}_0 \leq 1$, whereas the disease will persist when $\mathcal{R}_0 > 1$. We now investigate uniform persistence; we claim the following result.

Theorem 5 *If $\mathcal{R}_0 > 1$, system (3) is uniformly persistent, namely, there exists a constant $\zeta > 0$ such that*

$$\liminf_{t \rightarrow \infty} S_i(t) > \zeta, \quad \liminf_{t \rightarrow \infty} I_i(t) > \zeta, \quad \liminf_{t \rightarrow \infty} D_i(t) > \zeta, \quad \liminf_{t \rightarrow \infty} W_i(t) > \zeta,$$

where ζ is independent of the initial data in \mathcal{G} .

Proof We prove that system (3) is uniformly persistent with respect to $(X_0, \partial X_0)$, where

$$\begin{aligned} X &= \{(S_i, I_i, D_i, W_i) : S_i \geq 0, I_i \geq 0, D_i \geq 0, W_i \geq 0, i = 1, 2\}, \\ X_0 &= \{(S_i, I_i, D_i, W_i) : S_i \geq 0, I_i > 0, D_i > 0, W_i > 0, i = 1, 2\}, \\ \partial X_0 &= X/X_0. \end{aligned} \tag{13}$$

By the form of (3), it is very easy to see that both X and X_0 are positively invariant and ∂X_0 is relatively closed in X . Furthermore, system (3) is point dissipative. Let

$$M_\partial = \{(S_i(0), I_i(0), D_i(0), W_i(0)) | (S_i(t), I_i(t), D_i(t), W_i(t)) \in \partial X_0, \forall t \geq 0, i = 1, 2\}. \tag{14}$$

We now show that

$$M_\partial = \{(S_i(t), 0, 0, 0) | S_i(t) \geq 0, i = 1, 2\}. \tag{15}$$

It is worth noting that

$$\{(S_i(t), 0, 0, 0) | S_i(t) \geq 0, i = 1, 2\} \subseteq M_\partial, \tag{16}$$

so we only need to prove

$$M_\partial \subseteq \{(S_i(t), 0, 0, 0) | S_i(t) \geq 0, i = 1, 2\}. \tag{17}$$

Assume $\{(S_i(0), I_i(0), D_i(0), W_i(0))\} \in M_\partial$, for $i = 1, 2$. It suffices to show that

$$I_i(t) = D_i(t) = W_i(t) = 0, \quad \forall t \geq 0.$$

Suppose not, then there exist an $i_0, i_0 = 2$ and $t_0 \geq 0$ such that

$$(I_1(t_0), D_1(t_0), W_1(t_0))^T = 0, \quad (I_2(t_0), D_2(t_0), W_2(t_0))^T > 0 \tag{18}$$

without loss of generality, assume

$$I_2(t_0) = 0, \quad D_2(t_0) = 0, \quad W_2(t_0) > 0, \tag{19}$$

then we have

$$\frac{dW_2(t_0)}{dt} = r_2 W_2(t_0) \geq 0. \tag{20}$$

It follows that there is an $\eta > 0$ such that $W_2(t) > 0$, for $t_0 < t < t_0 + \eta$. This means that

$$(S_i(t), I_i(t), D_i(t), W_i(t))$$

does not belong to $\partial X_0, i = 1, 2$ for $t_0 < t < t_0 + \eta$, which contradicts the assumption that

$$(S_i(0), I_i(0), D_i(0), W_i(0)) \in M_\partial, \quad i = 1, 2.$$

Thus (15) holds.

It is worth noting that \mathcal{E}^0 is globally stable for system (3). It is clear that there is only an equilibria \mathcal{E}^0 in M_∂ , by the aforementioned claim, it then follows that \mathcal{E}^0 is an isolated invariant set in X , $W^S(\mathcal{E}^0) \cap X_0 = \emptyset$. Clearly, every orbit in M_∂ converges to \mathcal{E}^0 , \mathcal{E}^0 is acyclic in M_∂ . Using Theorem 4.6 in Thieme [62], we conclude that system (3) is uniformly persistent with respect to $(X_0, \partial X_0)$.

By the result of [63–65], system (3) has an equilibrium $\mathcal{E}^* = (\bar{S}_i, \bar{I}_i, \bar{D}_i, \bar{W}_i) \in X_0, i = 1, 2$. We further claim that $\bar{S}_i > 0, i = 1, 2$. Suppose that $\bar{S}_i = 0, i = 1, 2$, from the second equation of (3), we can get $\bar{I}_i = \bar{D}_i = \bar{W}_i = 0, i = 1, 2$. It is a contradiction. Then $(\bar{S}_i, \bar{I}_i, \bar{D}_i, \bar{W}_i, i = 1, 2)$, is a component-wise positive equilibrium of system (3). The proof is complete. \square

3.2 Existence of equilibria

System (3) has one disease-free equilibrium, $\mathcal{E}^0 = (S_1^0, 0, 0, 0, S_2^0, 0, 0, 0)$ and three endemic equilibria of the forms $\mathcal{E}_1^* = (S_1^*, I_1^*, D_1^*, W_1^*, S_2^*, 0, 0, 0)$, $\mathcal{E}_2^* = (S_1^{**}, 0, 0, 0, S_2^{**}, I_1^{**}, D_1^{**}, W_1^{**})$ and $\mathcal{E}^* = (\bar{S}_1, \bar{I}_1, \bar{D}_1, \bar{W}_1, \bar{S}_2, \bar{I}_2, \bar{D}_2, \bar{W}_2)$, corresponding respectively to states where the disease persists in the first sub-population and dies out in the second sub-population, the disease persists in the second sub-population and disappears in the first sub-population, and when the disease persists in the two sub-populations.

3.2.1 The first boundary equilibrium

The patch-1 system is obtained by making $I_2 = D_2 = W_2 = 0$, and it is given by

$$\begin{cases} S_1' = b_1 + a_2 S_2 - \Lambda_1 S_1 - (\mu_1 + a_1) S_1, \\ I_1' = \Lambda_1 S_1 - k_1 I_1, \\ D_1' = \nu_1 I_1 - \alpha_1 D_1, \\ W_1' = \delta_1 I_1 + \rho_1 D_1 - r_1 W_1, \\ S_2' = b_2 + a_1 S_1 - (\mu_2 + a_2) S_2. \end{cases} \tag{21}$$

Let $\mathcal{E}_1^* = (S_1^*, I_1^*, D_1^*, W_1^*, S_2^*, 0, 0, 0)$ be the first boundary endemic equilibrium with $I_1, D_1, W_1 > 0$. Then the endemic equilibrium \mathcal{E}_1^* can be obtained by setting to zero the right hand side of Eqs. (21), giving

$$\begin{cases} b_1 + a_2 S_2^* = (\beta_{I_1} I_1^* + \beta_{D_1} D_1^* + \beta_{W_1} W_1^*) S_1^* + (\mu_1 + a_1) S_1^*, \\ b_2 + a_1 S_1^* = (a_2 + \mu_2) S_2^*, \\ (\beta_{I_1} I_1^* + \beta_{D_1} D_1^* + \beta_{W_1} W_1^*) S_1^* = k_1 I_1^*, \\ \nu_1 I_1^* = \alpha_1 D_1^*, \\ \delta_1 I_1^* + \rho_1 D_1^* = r_1 W_1^*. \end{cases} \tag{22}$$

It is straightforward to prove that, when $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 \leq 1$, system (3) has exactly one non-trivial boundary endemic equilibrium $\mathcal{E}_1^* = (S_1^*, I_1^*, D_1^*, W_1^*, S_2^*, 0, 0, 0)$ where $S_1^*, I_1^*, D_1^*, W_1^*, S_2^*$ are given by

$$\begin{cases} S_1^* = \frac{S_1^0}{\mathcal{R}_1}, & S_2^* = \frac{a_1 + b_2 \mathcal{R}_1}{(a_2 + \mu_2) \mathcal{R}_1}, & I_1^* = \frac{[(a_2 + b_2) b_1 + a_2 b_2] (\mathcal{R}_1 - 1)}{(a_2 + \mu_2) k_1 \mathcal{R}_1}, \\ I_1^* = \frac{[\nu_1 ((a_2 + b_2) b_1 + a_2 b_2)] (\mathcal{R}_1 - 1)}{a_1 (a_2 + \mu_2) k_1 \mathcal{R}_1}, & & I_1^* = \frac{[(\alpha_1 \delta_1 + \nu_1 \rho_1) ((a_2 + b_2) b_1 + a_2 b_2)] (\mathcal{R}_1 - 1)}{\alpha_1 r_1 k_1 (a_2 + \mu_2) \mathcal{R}_1}. \end{cases} \tag{23}$$

We now present the local stability of \mathcal{E}_1^* when the reproduction number is close to 1. We make use of the centre manifold theory [66], but we first present the following lemma.

Lemma 1 *Consider the following general system of ordinary differential equations with parameter ϕ :*

$$\frac{dx}{dt} = f(x, \phi), \quad f : \mathbb{R}^n \times \mathbb{R} \rightarrow \mathbb{R}, \quad \text{and} \quad f \in \mathbb{C}^2 \quad (\mathbb{R}^n \times \mathbb{R}^n), \tag{24}$$

where 0 is an equilibrium of the system, that is, $f(0, \phi) = 0 \forall \phi$, and assume:

A1) $A = D_x f(0, 0) = (\frac{\partial f_i}{\partial x_j}(0, 0))$ is the linearisation of system (24) around the equilibrium 0 and ϕ evaluated at 0. Zero is a simple eigenvalue of A and other eigenvalues of A have negative real parts

A2) Matrix A has the right eigenvector u and a left eigenvector v corresponding to the zero eigenvalue. Let f_k be the k th component of f and

$$a = \sum_{k,i,j=1}^n q_k p_i p_j \frac{\partial^2 f_k}{\partial x_i \partial x_j}(0, 0),$$

$$b = \sum_{k,i=1}^n q_k p_i \frac{\partial^2 f_k}{\partial x_i \partial \phi}(0, 0). \tag{25}$$

The local dynamics of (24) around zero are totally governed by a and b .

- i. $a > 0, b > 0$. When $\phi < 0$ with $|\phi| \ll 1$, and there exists a positive unstable equilibrium, when $0 < \phi \ll 1$, 0 is unstable and there exists a negative and locally asymptotically stable equilibrium.
- ii. $a < 0, b < 0$. When $\phi < 0$ with $|\phi| \ll 1$, 0 unstable; when $0 < \phi \ll 1$, 0 is locally asymptotically stable, and there exists a positive unstable equilibrium.
- iii. $a > 0, b < 0$. When $\phi < 0$ with $|\phi| \ll 1$, 0 is unstable, and there exists a locally asymptotically stable negative equilibrium; when $0 < \phi \ll 1$, 0 is stable, and a positive unstable equilibrium appears.
- iv. $a < 0, b > 0$. When ϕ changes from negative to positive, 0 changes its stability from stable to unstable. Corresponding a negative unstable equilibrium becomes positive and locally asymptotically stable.

Theorem 6 *The unique patch 1-only boundary equilibrium, \mathcal{E}_1^* , of system (3), is locally asymptotically stable when $\mathcal{R}_1 > 1$ but only if \mathcal{R}_1 is close to 1.*

The proof of Theorem 6 is outlined in Appendix 3.

3.2.2 The second boundary equilibrium

Let $\mathcal{E}_2^{**} = (S_1^{**}, 0, 0, 0, S_2^{**}, I_2^{**}, D_2^{**}, W_2^{**})$ be the second boundary endemic equilibrium with $I_2, D_2, W_2 \neq 0$. Then the endemic equilibrium \mathcal{E}_2^{**} (steady state with $I_2, D_2, W_2 > 0$) can be obtained by setting the right hand side of equations in system (3) to zero, that

is,

$$\begin{cases} b_1 + a_2 S_2^{**} = (a_1 + \mu_1) S_1^{**}, \\ b_2 + a_1 S_1^{**} = (\beta_{I_2} I_2^{**} + \beta_{D_2} D_2^{**} + \beta_{W_2} W_2^{**}) S_2^{**} + (\mu_2 + a_2) S_2^{**}, \\ (\beta_{I_2} I_2^{**} + \beta_{D_2} D_2^{**} + \beta_{W_2} W_2^{**}) S_2^{**} = k_2 I_2^{**}, \\ v_2 I_2^{**} = \alpha_2 D_2^{**}, \\ \delta_2 I_2^{**} + \rho_2 D_2^{**} = r_2 W_2^{**}. \end{cases} \tag{26}$$

It is straightforward to prove that, when $\mathcal{R}_2 > 1$ and $\mathcal{R}_1 \leq 1$, system (3) has exactly one non-trivial boundary endemic equilibrium $\mathcal{E}_2^{**} = (S_1^{**}, 0, 0, S_2^{**}, I_2^{**}, D_2^{**}, W_2^{**})$ where $S_1^{**}, S_2^{**}, I_2^{**}, D_2^{**}, W_2^{**}$ are given by

$$\begin{cases} S_1^{**} = \frac{a_2 + b_1 \mathcal{R}_2}{(a_1 + \mu_1) \mathcal{R}_2}, & S_2^{**} = \frac{S_2^0}{\mathcal{R}_2}, & I_2^{**} = \frac{[(a_1 + b_1) b_2 + a_1 b_1] (\mathcal{R}_2 - 1)}{(a_1 + \mu_1) k_2 \mathcal{R}_2}, \\ D_2^{**} = \frac{[v_2 ((a_1 + b_1) b_2 + a_1 b_1)] (\mathcal{R}_2 - 1)}{a_2 (a_1 + \mu_1) k_2 \mathcal{R}_2}, & W_2^{**} = \frac{[(\alpha_2 \delta_2 + v_2 \rho_2) ((a_1 + b_1) b_2 + a_1 b_1)] (\mathcal{R}_2 - 1)}{\alpha_2 r_2 k_2 (a_1 + \mu_1) \mathcal{R}_2}. \end{cases} \tag{27}$$

Theorem 7 *The unique patch 2-only boundary equilibrium, \mathcal{E}_2^* , of system (3), is locally asymptotically stable when $\mathcal{R}_2 > 1$ but only if \mathcal{R}_2 is close to 1.*

Proof The proof for Theorem 7 follows the same procedure as outlined in Sect. 3.2.1 on the proof of the stability of patch 1-only boundary equilibrium, and the bifurcation parameter is computed by setting $\mathcal{R}_2 = 1$. □

3.2.3 *The interior endemic equilibrium*

$\mathcal{E}^* = (\bar{S}_1, \bar{I}_1, \bar{D}_1, \bar{W}_1, \bar{S}_2, \bar{I}_2, \bar{D}_2, \bar{W}_2)$ be the interior endemic equilibrium when both infectious of the two sub-populations co-exist, like, $I_i \neq 0, D_i \neq 0, W_i \neq 0, i = 1, 2$. To find this endemic equilibrium, we equate system (3) to zero and rewrite it as follows:

$$\begin{cases} b_i + a_j \bar{S}_j = \bar{\Lambda}_i \bar{S}_i + (\mu_i + a_i) \bar{S}_i, \\ \bar{\Lambda}_i \bar{S}_i = k_i \bar{I}_i, \\ v_i \bar{I}_i = \alpha_i \bar{D}_i, \\ \delta_i \bar{I}_i + \rho_i \bar{D}_i = r_i \bar{W}_i, \quad i = 1, 2, j = 1, 2; i \neq j. \end{cases} \tag{28}$$

Solutions of Eqs. (28) are defined by

$$\begin{cases} \bar{S}_1 = \frac{S_1^0}{\mathcal{R}_1^*}, & \bar{I}_1 = \frac{r_1 \alpha_1 (\mu_1 + a_1) (\mathcal{R}_1^* - 1)}{r_1 v_1 \beta_{D_1} + r_1 \alpha_1 \beta_{I_1} + (\alpha_1 \delta_1 + \mu_1 \rho_1) \beta_{W_1}}, \\ \bar{D}_1 = \frac{r_1 v_1 (\mu_1 + a_1) (\mathcal{R}_1^* - 1)}{r_1 v_1 \beta_{D_1} + r_1 \alpha_1 \beta_{I_1} + (\alpha_1 \delta_1 + \mu_1 \rho_1) \beta_{W_1}}, & \bar{W}_1 = \frac{(\mu_1 + a_1) (\alpha_1 \delta_1 + \mu_1 \rho_1) (\mathcal{R}_1^* - 1)}{r_1 v_1 \beta_{D_1} + r_1 \alpha_1 \beta_{I_1} + (\alpha_1 \delta_1 + \mu_1 \rho_1) \beta_{W_1}}, \\ \bar{S}_2 = \frac{S_2^0}{\mathcal{R}_2^*}, & \bar{I}_2 = \frac{r_2 \alpha_2 (\mu_2 + a_2) (\mathcal{R}_2^* - 1)}{r_2 v_2 \beta_{D_2} + r_2 \alpha_2 \beta_{I_2} + (\alpha_2 \delta_2 + \mu_2 \rho_2) \beta_{W_2}}, \\ \bar{D}_2 = \frac{r_2 v_2 (\mu_2 + a_2) (\mathcal{R}_2^* - 1)}{r_2 v_2 \beta_{D_2} + r_2 \alpha_2 \beta_{I_2} + (\alpha_2 \delta_2 + \mu_2 \rho_2) \beta_{W_2}}, & \bar{W}_2 = \frac{(\mu_2 + a_2) (\alpha_2 \delta_2 + \mu_2 \rho_2) (\mathcal{R}_2^* - 1)}{r_2 v_2 \beta_{D_2} + r_2 \alpha_2 \beta_{I_2} + (\alpha_2 \delta_2 + \mu_2 \rho_2) \beta_{W_2}}, \end{cases} \tag{29}$$

where

$$\mathcal{R}_1^* = \frac{a_1 S_1^0 + b_1 \mathcal{R}_2}{(\mu_1 + a_1) S_1^0} \mathcal{R}_2 \quad \text{and} \quad \mathcal{R}_2^* = \frac{a_2 S_2^0 + b_2 \mathcal{R}_1}{(\mu_2 + a_2) S_2^0} \mathcal{R}_1. \tag{30}$$

Note that, when $\mathcal{R}_1^* > 1$ and $\mathcal{R}_2^* > 1$, one has $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 > 1$. Then system (3) has a non-trivial interior endemic equilibrium $\mathcal{E}^* = (\bar{S}_1, \bar{I}_1, \bar{D}_1, \bar{W}_1, \bar{S}_2, \bar{I}_2, \bar{D}_2, \bar{W}_2)$ when $\mathcal{R}_1^* > 1$ and $\mathcal{R}_2^* > 1$.

We now investigate the local stability of the interior endemic equilibrium (29), and we shall make use of Lemma 1. We claim the following result.

Theorem 8 *The unique interior equilibrium, \mathcal{E}^* , of system (3), is locally asymptotically stable when $\mathcal{R}_0 > 1$ but only if the corresponding reproduction number is close to 1.*

The proof of Theorem 8 is outlined in Appendix 4.

The existence of the endemic equilibrium of system (3) is summarised in the following theorem.

Theorem 9 *System (3) has:*

1. *A boundary endemic equilibrium of the form $\mathcal{E}_1^* = (S_1^*, I_1^*, D_1^*, W_1^*, S_2^*, 0, 0, 0)$ whenever $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 \leq 1$. This means that the disease is endemic in the first sub-population and dies out in the second population.*
2. *A boundary endemic equilibrium of the form $\mathcal{E}_2^{**} = (S_1^{**}, 0, 0, 0, S_2^{**}, I_2^{**}, D_2^{**}, W_2^{**})$ whenever $\mathcal{R}_1 \leq 1$ and $\mathcal{R}_2 > 1$. This means that the disease is endemic in the second sub-population and dies out in the first population.*
3. *An interior endemic equilibrium of the form $\mathcal{E}^* = (\bar{S}_1, \bar{I}_1, \bar{D}_1, \bar{W}_1, \bar{S}_2, \bar{I}_2, \bar{D}_2, \bar{W}_2)$ whenever $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 > 1$ which corresponds to the case when the disease persists in the two sub-populations.*

4 Optimal control in educational campaigns

In this section, an optimal control problem is formulated by incorporating two intervention strategies into our basic model (3). Optimal control has been applied in the study of infectious diseases such as in [67–69]. Stopping the EVD transmission chain is still the main strategy to limit the spread of the disease. This can be achieved through educational campaigns that encourage behavioural changes like safe burial practices, self-isolation and proper hygiene [70]. To include educational campaigns into patch 1 and patch 2 as controls, u_1 and u_2 respectively, we extend system (3). The controlling effort u_1 represents a time-dependent control in educational campaigns with respect to patch 1 and u_2 representing a time-dependent control in educational campaigns with respect to patch 2. *Educational campaigns, in this case, are aimed at encouraging the uninfected to have some protective behaviours.* We assume that $u_1 > u_2$ since some neighbouring countries or communities might have lesser efforts compared to their neighbours in terms of control efforts. The extended model is given by

$$\begin{cases} S'_i = b_i + a_j S_j - (1 - u_i) \Lambda_i S_i - (\mu_i + a_i) S_i, \\ I'_i = (1 - u_i) \Lambda_i S_i - (\mu_i + \nu_i + \phi_i) I_i, \\ R'_i = \phi_i I_i + b_j R_j - (\mu_i + b_i) R_i, \\ D'_i = \nu_i I_i - \alpha_i D_i, \\ W'_i = \delta_i I_i + \rho_i D_i - r_i W_i, \quad i = 1, 2; j = 1, 2; i \neq j. \end{cases} \tag{31}$$

A successful control strategy is one that reduces the number of individuals infected with Ebola while minimising the costs associated with these efforts. Thus, our goal is to find a

pair of controls (u_1^*, u_2^*) that minimise the number of infected individuals and the cost of educational campaigns. Our objective functional is therefore formulated as

$$J(u_1, u_2) = \int_0^T [A_1 I_1 + A_2 I_2 + B_1 u_1^2 + B_2 u_2^2] dt, \tag{32}$$

subject to the constraints given by (31) and where A_1, A_2, B_1, B_2 are positive balancing coefficients transferring integrals into monetary quantity over a finite period of time T weeks. The first two terms in (32) with coefficients A_1 and A_2 represent the weights of the individuals I_1 and I_2 , respectively. The coefficients B_1 and B_2 represent the cost associated with the efforts in educational campaigns for patch 1 and 2, respectively. We seek the pair $(u_1^*, u_2^*) \in \mathcal{U}$ such that

$$\mathcal{J}(u_1^*, u_2^*) = \inf_{(u_1, u_2) \in \mathcal{U}} \mathcal{J}(u_1, u_2)$$

subject to the state system given by (31), where

$$\mathcal{U} = \{ (u_1(t), u_2(t)) : \text{is Lebesgue measurable on } [0, T], 0 \leq u_i(t) \leq u_{i\max}, i = 1, 2 \} \tag{33}$$

is the control set. Next, we derive the optimality system.

4.1 The optimality system

Using the Pontryagin maximum principle [71], we now derive the necessary conditions that an optimal control and corresponding states must satisfy. This principle converts (31)–(33) into a problem of minimising pointwise a Hamiltonian \mathcal{H} , with respect to $(u_1(t), u_2(t))$:

$$\begin{aligned} \mathcal{H} = & [A_1 I_1 + A_2 I_2 + B_1 u_1^2 + B_2 u_2^2] \\ & + \lambda_1 [b_1 + a_2 S_2 - (1 - u_1) \Lambda_1 S_1 - (\mu_1 + a_1) S_1] \\ & + \lambda_2 [(1 - u_1) \Lambda_1 S_1 - (\mu_1 + \nu_1 + \phi_1) I_1] \\ & + \lambda_3 [\phi_1 I_1 + b_2 R_2 - (\mu_1 + b_1) R_1] + \lambda_4 [\nu_1 I_1 - \alpha_1 D_1] + \lambda_5 [\delta_1 I_1 + \rho_1 D_1 - r_1 W_1] \\ & + \lambda_6 [b_2 + a_1 S_1 - (1 - u_2) \Lambda_2 S_2 - (\mu_2 + a_2) S_2] \\ & + \lambda_7 [(1 - u_2) \Lambda_2 S_2 - (\mu_2 + \nu_2 + \phi_2) I_2] \\ & + \lambda_8 [\phi_2 I_2 + b_1 R_1 - (\mu_2 + b_2) R_2] + \lambda_9 [\nu_2 I_2 - \alpha_2 D_2] + \lambda_{10} [\delta_2 I_2 + \rho_2 D_2 - r_2 W_2], \end{aligned}$$

where $\lambda_i, i = 1, 2, \dots, 10$, are the adjoint variables. In the following theorem, we present the adjoint system and control characterisation.

Theorem 10 *Given an optimal control (u_1^*, u_2^*) and corresponding state solutions S_i, I_i, R_i, D_i, W_i of the corresponding state system (31) that minimises $\mathcal{J}(u_1, u_2)$ over \mathcal{U} , there exist*

adjoint variables (functions), $\lambda_i(t)$, for $i = 1, 2, \dots, 10$, satisfying

$$\begin{cases} \lambda'_1(t) = (1 - u_1)(\lambda_1 - \lambda_2)A_1 + a_1(\lambda_1 - \lambda_6) + \mu_1\lambda_1, \\ \lambda'_2(t) = -A_1 + (1 - u_1)\beta_{I_1}S_1(\lambda_1 - \lambda_2) + \phi_1(\lambda_2 - \lambda_3) + v_1(\lambda_2 - \lambda_4) - \delta_1\lambda_5 + \mu_1\lambda_2, \\ \lambda'_3(t) = b_1(\lambda_3 - \lambda_8) + \mu_1\lambda_3, \\ \lambda'_4(t) = (1 - u_1)\beta_{D_1}S_1(\lambda_1 - \lambda_2) - \rho_1\lambda_5 + \alpha_1\lambda_4, \\ \lambda'_5(t) = (1 - u_1)\beta_{W_1}S_1(\lambda_1 - \lambda_2) + r_1\lambda_5, \\ \lambda'_6(t) = (1 - u_2)(\lambda_6 - \lambda_7)A_2 + a_2(\lambda_6 - \lambda_1) + \mu_2\lambda_6, \\ \lambda'_7(t) = -A_2 + (1 - u_2)\beta_{I_2}S_2(\lambda_6 - \lambda_7) + \phi_2(\lambda_7 - \lambda_8) + v_2(\lambda_8 - \lambda_9) - \delta_2\lambda_{10} + \mu_2\lambda_7, \\ \lambda'_8(t) = b_2(\lambda_8 - \lambda_3) + \mu_2\lambda_8, \\ \lambda'_9(t) = (1 - u_2)\beta_{D_2}S_2(\lambda_6 - \lambda_7) - \rho_2\lambda_{10} + \alpha_2\lambda_9, \\ \lambda'_{10}(t) = (1 - u_2)\beta_{W_2}S_2(\lambda_6 - \lambda_7) + r_2\lambda_{10}, \end{cases} \tag{34}$$

with terminal conditions

$$\lambda_i(t) = 0, \quad i = 1, 2, \dots, 10. \tag{35}$$

Furthermore, the optimal controls u_1^* and u_2^* are represented by

$$\begin{cases} u_1^*(t) = \max\{0, \min(u_{1\max}, \frac{(\lambda_2 - \lambda_1)A_1S_1}{2C_1})\}, \\ u_2^*(t) = \max\{0, \min(u_{2\max}, \frac{(\lambda_7 - \lambda_6)A_2S_2}{2C_2})\}. \end{cases} \tag{36}$$

Proof The existence of optimal control follows from Corollary 4.1 of [72] since the integrand of \mathcal{J} is a convex function of (u_1, u_2) and the state system satisfies the *Lipshitz* property with respect to the state variables. The following can be derived from the Pontryagin maximum principle [71]:

$$\lambda'_1 = \frac{\partial \mathcal{H}}{\partial S_1}, \quad \lambda'_2 = \frac{\partial \mathcal{H}}{\partial I_1}, \quad \dots, \quad \lambda'_{10} = \frac{\partial \mathcal{H}}{\partial W_2}, \tag{37}$$

with $\lambda_i(T) = 0$ for $i = 1, 2, \dots, 10$ evaluated at the optimal controls and corresponding states, which results in the adjoint system (34). The Hamiltonian \mathcal{H} is minimised with respect to the controls at the optimal controls, so we differentiate \mathcal{H} with respect to u_1 and u_2 on the set \mathcal{U} , respectively, giving the following optimality conditions:

$$\begin{cases} \frac{\partial \mathcal{H}}{\partial u_1} = (\lambda_1 - \lambda_2)A_1S_1 + 2C_1u_1 = 0, \\ \frac{\partial \mathcal{H}}{\partial u_2} = (\lambda_6 - \lambda_7)A_2S_2 + 2C_2u_2 = 0. \end{cases} \tag{38}$$

Hence, we obtain

$$\begin{cases} u_1^*(t) = \frac{(\lambda_2 - \lambda_1)A_1S_1}{2C_1}, \\ u_2^*(t) = \frac{(\lambda_7 - \lambda_6)A_2S_2}{2C_2}. \end{cases} \tag{39}$$

Taking into account the bounds on the controls, we obtain the desired characterisations. \square

Table 1 Model parameters, $i = 1, 2$, for patch 1 and 2 respectively. The time unit is in weeks

Definition	Symbol	Baseline values	Source
Effective contact rate of human individuals	β_i	0.16	[74]
Effective contact rate of deceased human individuals	β_{D_i}	0.489	[74]
Effective contact rate of Ebola virus	β_{W_i}	0.062	[74]
Rate of migration from first sub-population to the second sub-population (susceptible)	a_i	(0, 0.3)	Assumed
Rate of migration from first sub population to the second sub population (recovered)	b_i	(0, 0.3)	Assumed
Natural death rate of human individuals	μ_i	(0, 1)	[75]
Recovery rate of human individuals	ϕ_i	0.018 (0.16–0.202)	[76]
Disease-induced death rate of human individuals	v_i	0.5	[74]
Decay rate of Ebola virus in the environment	r_i	(0, ∞)	[77, 78]
Shedding rate of infectious human individuals	δ_i	(0, ∞)	Assumed
Shedding rate of deceased human individuals	ρ_i	(0, ∞)	Assumed
Burial rate of deceased human individuals	α_i	(0, ∞)	[79, 80]

5 Numerical simulations

In this section, we now provide some numerical simulations. The existence of an optimal control is provided and the behaviour of the optimality system made of 10 ordinary differential equations is evaluated through numerical simulations done with Matlab. The optimality system is solved using an iterative method with the Runge–Kutta fourth order scheme. Starting with a guess for the adjoint variables, the state equations are solved forward in time. Then these state values are used to solve the adjoint equations backward in time, and the iterations continue until convergence.

To illustrate the results of the foregoing analysis, we have simulated system (3) using the parameters in Table 1. Parameters in Table 1 are taken from the recent works on the West African Ebola outbreaks. We then assume some of the parameters in the realistic range for illustrative purposes. We are using the same table of parameters for both patches. Among the estimated parameters, are the balancing coefficients which have been arbitrarily chosen for illustration purposes. These weight parameters determine the importance of variables in the objective functional [73]. Thus $A_1 = 100, A_2 = 100, B_1 = B_2 = 90$.

We first consider a scenario where we have net migration on a_2 , thus, $a_2 > a_1$ implying that we have $\mathcal{R}_1 > \mathcal{R}_2 > 1$. We take $a_1 = 0.03, a_2 = 0.3$ and we assume that all other parameters take the same values for the two patches (see Table 1).

Figures 1(a) and 1(b) generally show that the controls have an impact on the control of EVD. Figure 1(a) shows that the controls are highly effective in patch 1 since in the presence of the control the infected population is very low for the whole period under study. Figure 1(b) depicts that the controls are effective only after 9 weeks. Figure 1(c) and 1(d) represents the controls u_1 and u_2 respectively. Both controls are at the upper bound for the whole period under study, 32 weeks. Thus under the given scenario, $a_2 > a_1$, both controls are feasible and effective for the whole period under study, hence more effort should be devoted to both controls in implementation.

We now consider a scenario where we have 0 net migration, thus $a_2 = a_1$, implying that we have $\mathcal{R}_1 = \mathcal{R}_2 > 1$. We take $a_1 = 0.165, a_2 = 0.165$ and we assume that all other parameters take the same values for the two patches (see Table 1).

Figure 2(a) and 2(b) shows that the controls are effective in the reduction of EVD. Figure 2(a) shows that the control u_1 becomes effective after 14 weeks and Fig. 2(a) shows that control u_2 becomes effective after 2 weeks. It is worth noting that Fig. 2(a) and 2(b)

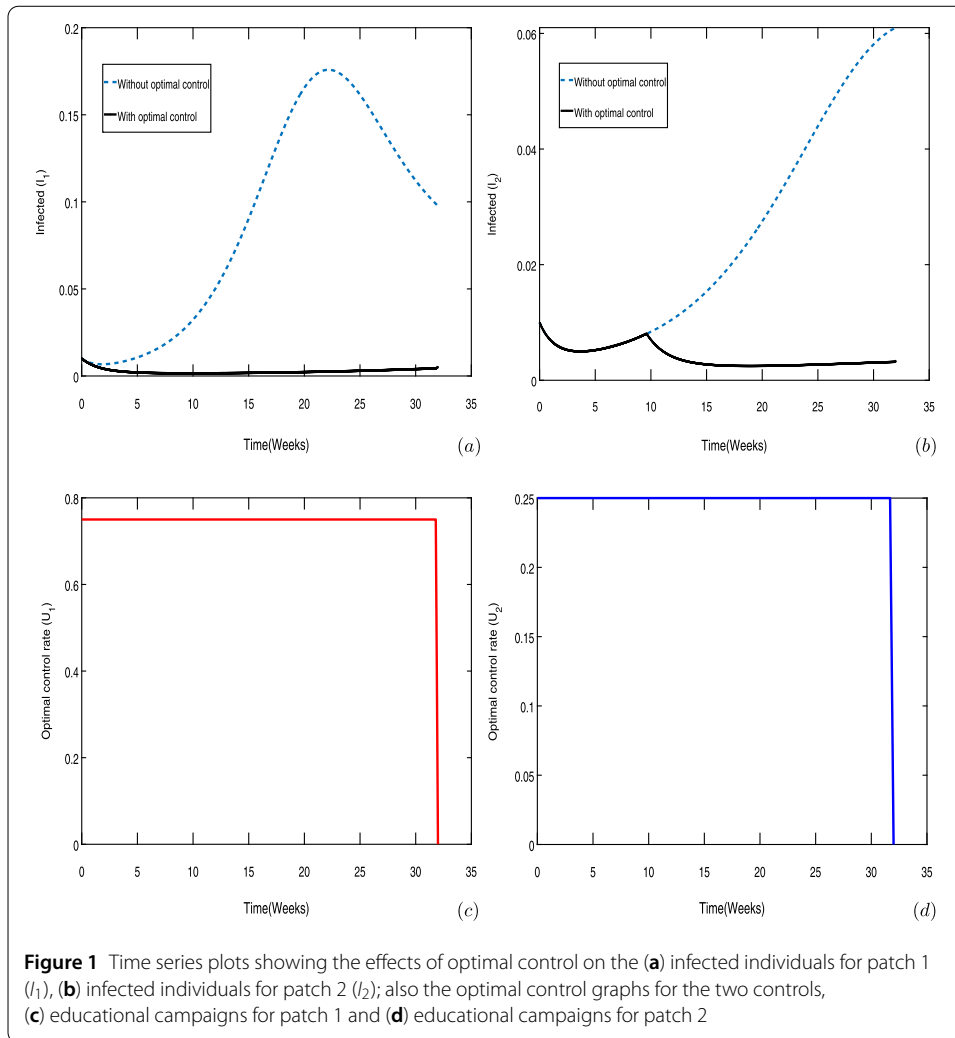
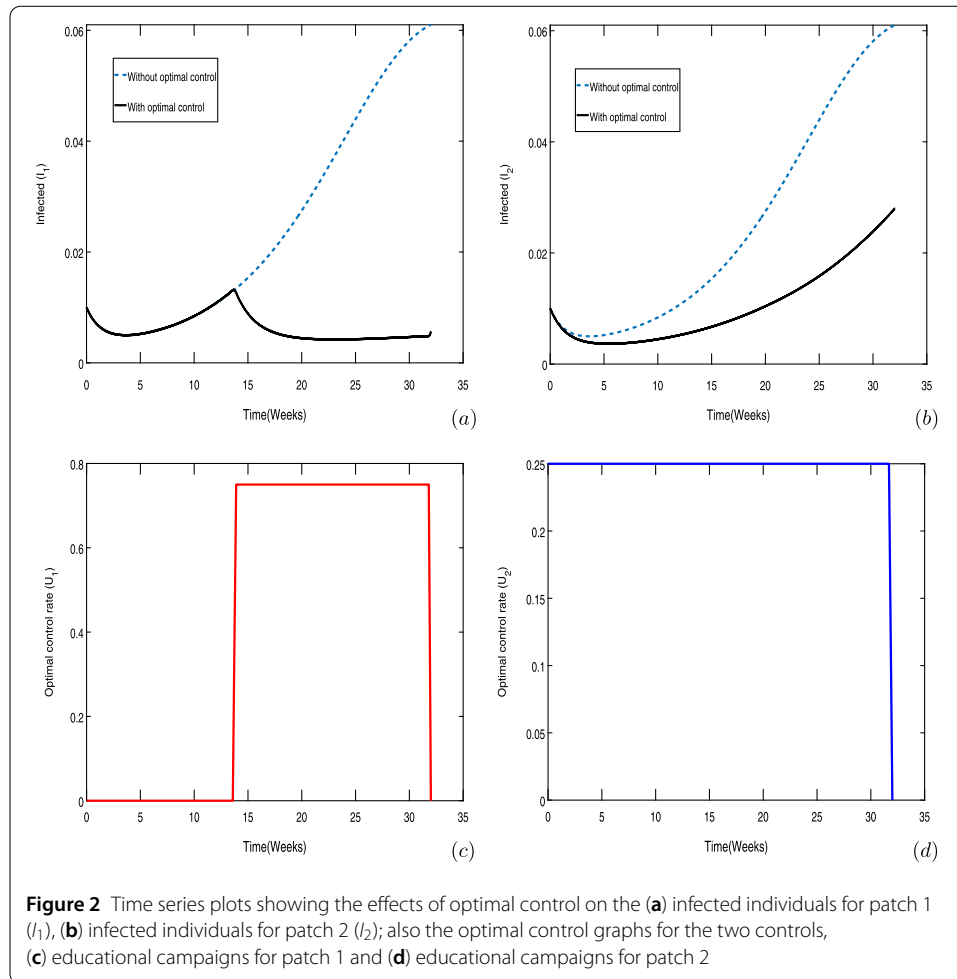


Figure 1 Time series plots showing the effects of optimal control on the (a) infected individuals for patch 1 (I_1), (b) infected individuals for patch 2 (I_2); also the optimal control graphs for the two controls, (c) educational campaigns for patch 1 and (d) educational campaigns for patch 2

shows that the controls are significant to a certain extent in the reduction of EVD. Figure 2(c) and 2(d) shows that the controls may not be sustainable for certain time intervals. In Fig. 2(c) the control u_1 is at the lower bound for 14 weeks. Thus, the control u_1 is initiated after 14 weeks under this scenario. Control u_2 starts at the upper bound and stays at the upper bound for the rest of the period under study. In this scenario, $a_2 = a_1$, more effort should be devoted to control u_2 . It is worth noting that control u_2 is the less efficient one, hence devoting more effort to control u_2 in controlling EVD might not be ideal.

Lastly, we consider a scenario where we have a net migration in a_1 , thus $a_1 > a_2$, implying that we have $\mathcal{R}_2 > \mathcal{R}_1 > 1$. We take $a_1 = 0.3$, $a_2 = 0.03$ and we assume that all other parameters take the same values for the two patches (see Table 1).

Figures 3(a) and 3(b) depict that the controls are only effective up to a certain extent. Figure 3(a), shows that the controls are effective during the last 28 weeks of the whole of 32 weeks under study. Figure 3(b) shows that the controls are only effective in the first 28 weeks only. Figure 3(c) and 3(d) also shows that the controls may not be sustainable for certain time intervals. Figure 3(c) shows that the control u_1 starts at the lower bound and is initiated after 26 weeks. Figure 3(d) shows that control u_2 is at the upper bound for the rest of the period under study. Under this scenario, $a_1 > a_2$, it is worth noting that more



effort should be devoted to control u_2 . Under the given scenario control, u_1 , which is the most effective, it is not feasible.

6 Discussion

Several countries in West Africa, in particular, Sierra Leone, Guinea and Liberia experienced morbidity and mortality during the Ebola epidemic from 2013–2015. At the time of this epidemic there was no known vaccine or drug, so effective disease control required coordinated efforts that include both standard strategies, such as hospitalisation, as well as community efforts, such as safe burial practices, proper hygiene in hospitals, etc. Not only are such efforts difficult to implement in practice, but there is also added complexity with connectivity between populations that have different policies in place. These complexities may affect some communities, that is, neighbouring communities might have lesser efforts compared to their neighbours in terms of control strategies. Currently, starting in August 2018 EVD is terrorising North Kivu in the Democratic Republic of Congo, a place with high mobility among the traders from different communities [7, 8].

In this manuscript, we formulated and analysed a differential equation-based two-patch model, with the patches connected by migration. EVD features and dynamics within a population such as the infectious environment, deceased individuals and the infectious individuals are incorporated in both patches. The susceptible and recovered individuals

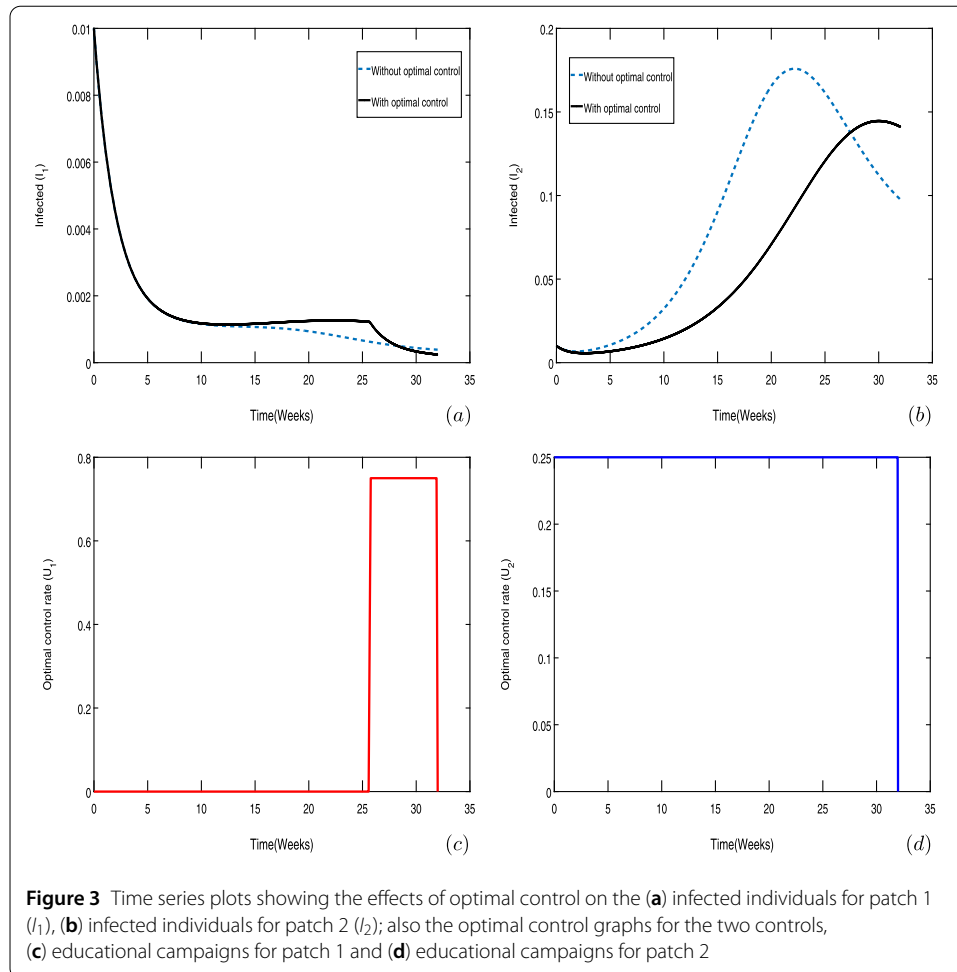


Figure 3 Time series plots showing the effects of optimal control on the (a) infected individuals for patch 1 (I_1), (b) infected individuals for patch 2 (I_2); also the optimal control graphs for the two controls, (c) educational campaigns for patch 1 and (d) educational campaigns for patch 2

are the only ones who migrate. The basic reproductive number, \mathcal{R}_0 , for the model has been computed. Our results show that \mathcal{R}_0 , can provide a sharp threshold for the disease dynamics when $\mathcal{R}_0 \leq 1$ the disease-free equilibrium is globally stable indicating that the disease would die out, but when $\mathcal{R}_0 > 1$ the disease persists. Subsequently, we have performed an optimal control study on this two-patch model to effectively design control strategies to control EVD. The control u_1 represents educational campaigns in patch 1 and the control u_2 represents educational campaigns in patch 2, with $u_1 > u_2$. We considered two communities with the same control strategy, which differ in their levels of efficacy. Our major aim is to assess the spread of the EVD epidemic within two communities connected by migration, with these communities employing the same control strategy which only differ in efficiency. The technical tool used to determine the optimal strategy is the Pontryagin maximum principle. Numerical simulations results show that the implementation of the optimal control has a huge impact on the reduction of the infected individuals in both patches, and that the outcome of the control from each patch may be different due to their different characteristics. We also realised that, given our controls $u_1 > u_2$, it is advisable to have $a_2 > a_1$, as in the net migration is into patch 1, so as to control EVD faster and with maximum effort for fewer resources. It is worth noting that for the scenario $a_2 > a_1$ we have $\mathcal{R}_1 > \mathcal{R}_2 > 1$. Thus, in the case of an EVD epidemic, we can manage to move the people who are not infected from places that have inefficient intervention strategies to

places with higher efficient intervention strategies, regardless of the relative magnitude of the respective reproduction numbers in both communities. Generally, the study finds that EVD can be controlled if optimal educational campaigns are implemented, although this might not be appropriate for certain time intervals.

However, just like any other model, we cannot say the model is complete, it can be extended to include the aspect of the refugees, rebels and health care workers, like the case in the Democratic Republic of Congo.

Appendix 1

Proof of Theorem 1 Rewriting system (3) in expanded form, we have the following system (system (3) is still the same as system (40)):

$$\begin{cases} S'_1 = b_1 + a_2S_2 - \Lambda_1S_1 - (\mu_1 + a_1)S_1, \\ I'_1 = \Lambda_1S_1 - (\mu_1 + \nu_1 + \phi_1)I_1, \\ D'_1 = \nu_1I_1 - \alpha_1D_1, \\ W'_1 = \delta_1I_1 + \rho_1D_1 - r_1W_1, \\ S'_2 = b_1 + a_1S_1 - \Lambda_2S_2 - (\mu_2 + a_2)S_2, \\ I'_2 = \Lambda_2S_2 - (\mu_2 + \nu_2 + \phi_2)I_2, \\ D'_2 = \nu_2I_2 - \alpha_2D_2, \\ W'_2 = \delta_2I_2 + \rho_2D_2 - r_2W_2. \end{cases} \tag{40}$$

It is worth noting that, since (40)₁ and (40)₅ are first order linear equations in S_1 and S_2 , respectively, it is easy to see that $S_1(t) > 0$ if and only if $S_2(t) > 0$. With this remark in mind, we shall prove below that $S_1(t) > 0$ for $t \geq 0$. To this end, put $t_1^0 = \sup\{t > 0, S_1(t) > 0\}$ and $t_2^0 = \sup\{t > 0, S_2(t) > 0\}$.

If $t_1^0 = +\infty$ and $t_2^0 = +\infty$, we use the above-mentioned remark to conclude that both $S_1(t)$ and $S_2(t)$ are positive for all $t \geq 0$. If $t_1^0 < \infty$ and $t_2^0 < \infty$, we are going to prove that this leads to a contradiction. By a continuity argument, the solution functions $S_1(t)$ and $S_2(t)$ change sign at least once in the intervals $J_1 = [t_1^0, +\infty)$ and $J_2 = [t_2^0, +\infty)$, respectively.

Denote by $t_1^m \in J_1$ and $t_2^m \in J_2$ the first real numbers such that $S_1(t_1^m) = 0$ and $S_2(t_2^m) = 0$, respectively. We then have

$$\begin{aligned} \forall t, 0 < t < t_1^{m1}, \quad S_1(t) > 0, \quad S_1(t_1^m) = 0 \quad \text{and} \\ \forall t, 0 < t < t_2^{m2}, \quad S_2(t) > 0, \quad S_2(t_2^m) = 0. \end{aligned} \tag{41}$$

Without loss of generality, suppose that $t_1^m \leq t_2^m$. Then, from system (40), we have

$$S'_1(t_1^m) = b_1 + a_2S_2(t_1^m) > 0. \tag{42}$$

Equation (42) implies that there exists a positive number $t_1^{m1} > t_1^m$ such that

$$S_1(t) > 0, \quad \forall 0 < t < t_1^{m1}. \tag{43}$$

Putting Eqs. (41) and (43) together and using the continuity of $S_1(t)$, we conclude that t_1^m is an extremum of $S_1(t)$. Moreover, since $S_1(t)$ is a differentiable function on \mathbb{R} , one has $S'(t_1^m) = 0$. This is a contradiction to (42). Therefore, $t_1^0 = +\infty$, which implies that $t_2^0 = +\infty$ as well.

To prove that $I_1(t), D_1(t), W_1(t), I_2(t), D_2(t), W_2(t) \geq 0$ for all $t \geq 0$, whenever

$$I_1(0), D_1(0), W_1(0), I_2(0), D_2(0), W_2(0) \geq 0,$$

we rewrite the corresponding equations in (40) in the form

$$x'(t) = \mathcal{M}x(t), \quad \text{where } x(t) = (I_1(t), D_1(t), W_1(t), I_2(t), D_2(t), W_2(t))^T, \tag{44}$$

$$\mathcal{M} = \begin{bmatrix} \beta_{I_1}S_1 - (\mu_1 + v_1 + \phi_1) & \beta_{D_1}S_1 & \beta_{W_1}S_1 & 0 & 0 & 0 \\ v_1 & -\alpha_1 & 0 & 0 & 0 & 0 \\ \delta_1 & \rho_1 & -r_1 & 0 & 0 & 0 \\ 0 & 0 & 0 & \beta_{I_2}S_2 - (\mu_2 + v_2 + \phi_2) & \beta_{D_2}S_2 & \beta_{W_2}S_2 \\ 0 & 0 & 0 & v_2 & -\alpha_2 & 0 \\ 0 & 0 & 0 & \delta_2 & \rho_2 & -r_2 \end{bmatrix}. \tag{45}$$

Since $S_1(t) > 0, S_2(t) > 0$ as established above, \mathcal{M} is a Metzler matrix. Thus (44) is a monotone system. Therefore \mathbb{R}_+^6 is invariant under the flow of system (44). This completes the proof of positivity of the solutions. \square

Appendix 2

Proof of Theorem 2 Adding the total human population we have

$$\begin{aligned} N' &= b_1 + b_2 - (\mu_1 + v_1 + \phi_1)I_1 - (\mu_2 + v_2 + \phi_2)I_2 - \mu_1S_1 - \mu_2S_2 \\ &\leq b - \mu_0N. \end{aligned} \tag{46}$$

Applying the Gronwall inequality to Eq. (46), we have

$$N(t) \leq \frac{b}{\mu_0} + \left(N(0) - \frac{b}{\mu_0}\right)e^{-\mu_0 t}, \quad \forall t \geq 0, \tag{47}$$

which implies that $0 \leq N(t) \leq \frac{b}{\mu_0}$ for all $t \geq 0$ if $N(0) \leq \frac{b}{\mu_0}$. Furthermore, it follows from Eq. (3)₃ that we have the relation

$$D' \leq \frac{\bar{v}b}{\mu_0} - \bar{\alpha}_1 D, \tag{48}$$

to which another application of the Gronwall inequality yields the bounds

$$0 \leq D(t) \leq \frac{\bar{v}b}{\alpha_1 \mu_0} + \left(D(0) - \frac{\bar{v}b}{\alpha_1 \mu_0}\right)e^{-\bar{v}t} \leq \frac{\bar{v}b}{\alpha_1 \mu_0}, \quad \forall t \geq 0, \tag{49}$$

where $D(0) \leq \frac{\bar{v}b}{\alpha_1 \mu_0}$.

The boundedness of $W(t)$ is proved similarly. \square

Appendix 3

Proof of Theorem 6 To apply Lemma 1, the following simplifications and change of variables are carried out. Let $S_1 = x_1, I_1 = x_2, D_1 = x_3, W_1 = x_4, S_2 = x_5, S_1^0 = x_1^0$. Furthermore, by using the vector notation $\mathcal{X} = (f_1, f_2, f_3, f_4, f_5)$, system (21) can be written in the form $\frac{d\mathcal{X}}{dt} = F(x)$, with $F = (f_1, f_2, f_3, f_4, f_5)^T$, such that

$$\begin{cases} \frac{dx_1}{dt} = f'_1 = b_1 + a_2x_5 - (\beta_{I_1}x_2 + \beta_{D_1}x_3 + \beta_{W_1}x_4)x_1 - (\mu_1 + a_1)x_1, \\ \frac{dx_2}{dt} = f'_2 = (\beta_{I_1}x_2 + \beta_{D_1}x_3 + \beta_{W_1}x_4)x_1 - k_1x_2, \\ \frac{dx_3}{dt} = f'_3 = v_1x_2 - \alpha_1x_3, \\ \frac{dx_4}{dt} = f'_4 = \delta_1x_2 + \rho_1x_3 - r_1x_4, \\ \frac{dx_5}{dt} = f'_5 = b_2 + a_1x_1 - (\mu_2 + a_2)x_2. \end{cases} \tag{50}$$

The Jacobian matrix of system (50) at \mathcal{E}^0 is given by

$$\begin{pmatrix} -(\mu_1 + a_1) & -\beta_{I_1}x_1^0 & -\beta_{D_1}x_1^0 & -\beta_{W_1}x_1^0 & a_2 \\ 0 & \beta_{I_1}x_1^0 - k_1 & \beta_{D_1}x_1^0 & \beta_{W_1}x_1^0 & 0 \\ 0 & v_1 & -\alpha_1 & 0 & 0 \\ 0 & \delta_1 & \rho_1 & -r_1 & 0 \\ a_1 & 0 & 0 & 0 & -(\mu_2 + a_2) \end{pmatrix},$$

from which it can be shown that

$$\mathcal{R}_1 = \frac{x_1^0\beta_{D_1}v_1}{k_1\alpha_1} + \frac{x_1^0\beta_{I_1}}{k_1} + \frac{x_1^0\beta_{W_1}(\alpha_1\delta_1 + \mu_1\rho_1)}{(k_1r_1\alpha_1)}, \tag{51}$$

with $S_1^0 = x_1^0$ as defined in Eq. (5).

Now, we consider $\varrho_1\beta_{I_1} = \beta_{D_1}$ and $\varrho_2\beta_{W_1} = \beta_{D_1}$ regardless of whether $\varrho_1, \varrho_2 \in (0, 1)$ or $\varrho_1, \varrho_2 \geq 1$. Taking β_{D_1} as the bifurcation parameter and considering that $\mathcal{R}_1 = 1$ and solving for β_{D_1} , we have

$$\beta^* = \beta_{D_1} = \frac{k_1\alpha_1r_1}{x_1^0(v_1r_1 + \varrho_1r_1\alpha_1 + \varrho_2(\alpha_1\delta_1 + \mu_1\rho_1))}. \tag{52}$$

The transformed system (50), with $\beta_{D_1} = \beta^*$, has a hyperbolic equilibrium point (like, the linearised system has a simple eigenvalue with zero real part, and all other eigenvalues have negative real part), so that the centre manifold theory [66] can be used to analyse the dynamics of (50) near $\beta_{D_1} = \beta^*$. It can be shown that the Jacobian of system (50) has a right eigenvector associated with the zero eigenvalue given by $p = (p_1, p_2, p_3, p_4, p_5)^T$, where

$$\begin{cases} p_1 = -\frac{\mu_2 + a_2}{\mu_2(\mu_1 + a_1) + \mu_1 a_2} \mathcal{H}p_3 < 0, & p_2 = \frac{\alpha_1}{v_1} p_3 > 0, & p_3 = p_3 > 0, \\ p_4 = \left(\frac{\delta_1\alpha_1 + \rho_1v_1}{r_1v_1}\right)p_3 > 0, & p_5 = -\frac{a_1}{\mu_2(\mu_1 + a_1) + \mu_1 a_2} \mathcal{H}p_3 < 0, \end{cases} \tag{53}$$

where

$$\mathcal{H} = \left(\frac{\alpha_1r_1\beta_{I_1}x_1^0 + r_1v_1\beta_{D_1}x_1^0 + \beta_{W_1}(\delta_1\alpha_1 + \rho_1v_1)x_1^0}{r_1v_1}\right). \tag{54}$$

The left eigenvectors of $J(\mathcal{E}^0)$ associated with the zero eigenvalue at $\beta_{D_1} = \beta^*$ is given by $q = (q_1, q_2, q_3, q_4, q_5)^T$, where

$$\begin{aligned} q_1 &= \frac{a_1}{(\mu_1 + a_1)} q_5, & q_2 &= \frac{r_1}{\beta_{W_1} S_1^0} q_4 + \frac{a_1}{(\mu_1 + a_1)} q_5, \\ q_3 &= \left(\frac{r_1 \beta_{D_1} + \rho_1 \beta_{W_1}}{\beta_{W_1}} \right) q_4, & q_4 &= q_4 > 0, & q_5 &= q_5 > 0. \end{aligned} \tag{55}$$

Computation of the bifurcation parameters a and b. For the sign of a , it can be shown that the associated non-vanishing partial derivatives of F are given by

$$\begin{aligned} \frac{\partial^2 f_1}{\partial x_1 \partial x_2} &= \frac{\partial^2 f_1}{\partial x_2 \partial x_1} = -\beta_{I_1}, & \frac{\partial^2 f_1}{\partial x_1 \partial x_3} &= \frac{\partial^2 f_1}{\partial x_3 \partial x_1} = -\beta_{D_1}, \\ \frac{\partial^2 f_1}{\partial x_1 \partial x_4} &= \frac{\partial^2 f_1}{\partial x_4 \partial x_1} = -\beta_{W_1}, & \frac{\partial^2 f_2}{\partial x_1 \partial x_2} &= \frac{\partial^2 f_2}{\partial x_2 \partial x_1} = \beta_{I_1}, \\ \frac{\partial^2 f_2}{\partial x_1 \partial x_3} &= \frac{\partial^2 f_2}{\partial x_3 \partial x_1} = \beta_{D_1}, & \frac{\partial^2 f_2}{\partial x_1 \partial x_4} &= \frac{\partial^2 f_2}{\partial x_4 \partial x_1} = \beta_{W_1}. \end{aligned} \tag{56}$$

From (56) it follows that

$$a = 2p_1(p_2\beta_{I_1} + p_3\beta_{D_1} + p_4\beta_{W_1}) \frac{r_1}{\beta_{W_1} x_1^0} q_4 < 0. \tag{57}$$

This excludes the possibility of a backward bifurcation since $a < 0$. For the sign of b , it is associated with the following non-vanishing partial derivatives of F :

$$\begin{aligned} \frac{\partial^2 f_1}{\partial x_2 \partial \beta^*} &= -\varrho_1 \beta_{D_1} x_1^0, & \frac{\partial^2 f_1}{\partial x_3 \partial \beta^*} &= -\beta_{D_1} x_1^0, & \frac{\partial^2 f_1}{\partial x_4 \partial \beta^*} &= -\varrho_2 \beta_{D_1} x_1^0, \\ \frac{\partial^2 f_2}{\partial \beta^* \partial x_2} &= \varrho_1 \beta_{D_1} x_1^0, & \frac{\partial^2 f_2}{\partial \beta^* \partial x_3} &= \beta_{D_1} x_1^0, & \frac{\partial^2 f_2}{\partial \beta^* \partial x_4} &= \varrho_2 \beta_{D_1} x_1^0. \end{aligned} \tag{58}$$

It follows from the expressions in (58) that

$$b = (p_2\beta_{I_1} + p_3\beta_{D_1} + p_4\beta_{W_1}) \frac{r_1}{\beta_{W_1} S_1^0} q_4 > 0. \tag{59}$$

Thus, $a < 0$ and $b > 0$ and applying Lemma 1 item (iv), we have established our result. The proof is complete. \square

Appendix 4

Proof of Theorem 6 To apply Lemma 1, it is convenient to make the following change of variables:

let $S_1 = x_1, I_1 = x_2, D_1 = x_3, W_1 = x_4, S_2 = x_5, I_2 = x_6, D_2 = x_7, W_2 = x_8, S_1^0 = x_1^0, S_2^0 = x_2^0$. Further, by using the vector notation $\mathcal{X} = (f_1, f_2, f_3, f_4, f_5, f_6, f_7, f_8)$, system (3) can be written

in the form $\frac{dx}{dt} = F(x)$, with $F = (f_1, f_2, f_3, f_4, f_5, f_6, f_7, f_8)^T$, such that

$$\begin{cases} \frac{dx_1}{dt} = f'_1 = b_1 + a_2x_5 - (\beta_{I_1}x_2 + \beta_{D_1}x_3 + \beta_{W_1}x_4)x_1 - (\mu_1 + a_1)x_1, \\ \frac{dx_2}{dt} = f'_2 = (\beta_{I_1}x_2 + \beta_{D_1}x_3 + \beta_{W_1}x_4)x_1 - k_1x_2, \\ \frac{dx_3}{dt} = f'_3 = v_1x_2 - \alpha_1x_3, \\ \frac{dx_4}{dt} = f'_4 = \delta_1x_2 + \rho_1x_3 - r_1x_4, \\ \frac{dx_5}{dt} = f'_5 = b_2 + a_1x_1 - (\beta_{I_2}x_6 + \beta_{D_2}x_7 + \beta_{W_2}x_8)x_5 - (\mu_2 + a_2)x_5, \\ \frac{dx_6}{dt} = f'_6 = (\beta_{I_2}x_6 + \beta_{D_2}x_7 + \beta_{W_2}x_8)x_5 - k_2x_6, \\ \frac{dx_7}{dt} = f'_7 = v_2x_6 - \alpha_2x_7, \\ \frac{dx_8}{dt} = f'_8 = \delta_2x_6 + \rho_2x_7 - r_2x_8. \end{cases} \tag{60}$$

The Jacobian matrix of system (60) at \mathcal{E}^0 is given by

$$\begin{pmatrix} -(\mu_1 + a_1) & -\beta_{I_1}x_1^0 & -\beta_{D_1}x_1^0 & -\beta_{W_1}x_1^0 & a_2 & 0 & 0 & 0 \\ 0 & \beta_{I_1}x_1^0 - k_1 & \beta_{D_1}x_1^0 & \beta_{W_1}x_1^0 & 0 & 0 & 0 & 0 \\ 0 & v_1 & -\alpha_1 & 0 & 0 & 0 & 0 & 0 \\ 0 & \delta_1 & \rho_1 & -r_1 & 0 & 0 & 0 & 0 \\ a_1 & 0 & 0 & 0 & -(\mu_2 + a_2) & -\beta_{I_2}x_2^0 & -\beta_{D_2}x_2^0 & -\beta_{W_2}x_2^0 \\ 0 & 0 & 0 & 0 & 0 & \beta_{I_2}x_2^0 - k_2 & \beta_{D_2}x_2^0 & \beta_{W_2}x_2^0 \\ 0 & 0 & 0 & 0 & 0 & v_2 & -\alpha_2 & 0 \\ 0 & 0 & 0 & 0 & 0 & \delta_2 & \rho_2 & -r_2 \end{pmatrix},$$

from which it can be shown that $\mathcal{R}_0 = \max\{\mathcal{R}_1, \mathcal{R}_2\}$, where

$$\mathcal{R}_i = \frac{S_i^0 \beta_{D_i} v_i}{k_i \alpha_i} + \frac{S_i^0 \beta_{I_i}}{k_i} + \frac{S_i^0 \beta_{W_i} (\alpha_i \delta_i + \mu_i \rho_i)}{(k_i r_i \alpha_i)}, \quad i = 1, 2. \tag{61}$$

Now, we consider $\varrho_1 \beta_{I_1} = \beta_{D_1}$, $\varrho_2 \beta_{W_1} = \beta_{D_1}$, $\varrho_3 \beta_{I_2} = \beta_{D_2}$, $\varrho_4 \beta_{W_2} = \beta_{D_2}$ regardless of whether $\varrho_1, \varrho_2, \varrho_3, \varrho_4 \in (0, 1)$ or $\varrho_1, \varrho_2, \varrho_3, \varrho_4 \geq 1$. Taking β_{D_1} as the bifurcation parameter and considering that $\mathcal{R}_0 = 1$ and solving for β_{D_1} , we have

$$\beta^* = \beta_{D_1} = \frac{k_1 \alpha_1 r_1}{S_1^0 (v_1 r_1 + \varrho_1 r_1 \alpha_1 + \varrho_2 (\alpha_1 \delta_1 + \mu_1 \rho_1))}. \tag{62}$$

Note that the linearised system of the transformed equation (60) with the bifurcation point β^* has a zero eigenvalue. Hence, the centre manifold theory [66] can be used to analyse the dynamics of system (60) near $\beta_{I_1} = \beta^*$. It can be shown that the Jacobian of system (60) has a right eigenvector associated with the zero eigenvalue given by $p = (p_1, p_2, p_3, p_4, p_5, p_6, p_7, p_8)^T$, where

$$\begin{cases} p_1 = -\frac{\mu_2 + a_2}{\mu_2(\mu_1 + a_1) + \mu_1 a_2} \mathcal{H} p_3 < 0, & p_2 = \frac{\alpha_1}{v_1} p_3 > 0, & p_3 = p_3 > 0, \\ p_4 = \left(\frac{\delta_1 \alpha_1 + \rho_1 v_1}{r_1 v_1}\right) p_3 > 0, & p_5 = -\frac{a_1}{\mu_2(\mu_1 + a_1) + \mu_1 a_2} \mathcal{H} p_3 < 0, & p_6 = \frac{\alpha_2}{v_2} p_7, \\ p_7 = p_7 > 0, & p_8 = \left(\frac{\delta_2 \alpha_2 + \rho_2 v_2}{r_2 v_2}\right) p_7 > 0, \end{cases} \tag{63}$$

with \mathcal{H} as defined in Eq. (54).

The left eigenvectors of $J(\mathcal{E}^0)$ associated with the zero eigenvalue at $\beta_{D_1} = \beta^*$ is given by $q = (q_1, q_2, q_3, q_4, q_5, q_6, q_7, q_8)^T$, where

$$\begin{cases} q_1 = \frac{a_1}{(\mu_1 + a_1)} q_5 > 0, & q_2 = \frac{r_1}{\beta_{W_1} S_1^0} q_4 + \frac{a_1}{(\mu_1 + a_1)} q_5 > 0, \\ q_3 = \left(\frac{r_1 \beta_{D_1} + \rho_1 \beta_{W_1}}{\beta_{W_1}}\right) q_4 > 0, & q_4 = q_4 > 0, \quad q_5 = q_5 > 0, \\ q_6 = q_5 + \frac{r_2}{\beta_{W_2} S_2^0} q_8 > 0, & q_7 = \left(\frac{r_2 \beta_{D_2}}{\alpha_2 \beta_{W_2}} + \frac{\rho_2}{\alpha_2}\right) q_8 > 0, \quad q_8 = q_8 > 0. \end{cases} \tag{64}$$

Computation of the bifurcation parameters a and b. For the sign of a , it can be shown that the associated non-vanishing partial derivatives of F are given by

$$\begin{aligned} \frac{\partial^2 f_1}{\partial x_1 \partial x_2} &= \frac{\partial^2 f_1}{\partial x_2 \partial x_1} = -\beta_{I_1}, & \frac{\partial^2 f_1}{\partial x_1 \partial x_3} &= \frac{\partial^2 f_1}{\partial x_3 \partial x_1} = -\beta_{D_1}, \\ \frac{\partial^2 f_1}{\partial x_1 \partial x_4} &= \frac{\partial^2 f_1}{\partial x_4 \partial x_1} = -\beta_{W_1}, & \frac{\partial^2 f_2}{\partial x_1 \partial x_2} &= \frac{\partial^2 f_2}{\partial x_2 \partial x_1} = \beta_{I_1}, \\ \frac{\partial^2 f_2}{\partial x_1 \partial x_3} &= \frac{\partial^2 f_2}{\partial x_3 \partial x_1} = \beta_{D_1}, & \frac{\partial^2 f_2}{\partial x_1 \partial x_4} &= \frac{\partial^2 f_2}{\partial x_4 \partial x_1} = \beta_{W_1}, \\ \frac{\partial^2 f_5}{\partial x_5 \partial x_6} &= \frac{\partial^2 f_5}{\partial x_6 \partial x_5} = -\beta_{I_2}, & \frac{\partial^2 f_5}{\partial x_5 \partial x_7} &= \frac{\partial^2 f_5}{\partial x_7 \partial x_5} = -\beta_{D_2}, \\ \frac{\partial^2 f_5}{\partial x_5 \partial x_8} &= \frac{\partial^2 f_5}{\partial x_8 \partial x_5} = -\beta_{W_2}, & \frac{\partial^2 f_6}{\partial x_5 \partial x_6} &= \frac{\partial^2 f_6}{\partial x_6 \partial x_5} = \beta_{I_2}, \\ \frac{\partial^2 f_6}{\partial x_5 \partial x_7} &= \frac{\partial^2 f_6}{\partial x_7 \partial x_5} = \beta_{D_2}, & \frac{\partial^2 f_6}{\partial x_6 \partial x_8} &= \frac{\partial^2 f_6}{\partial x_8 \partial x_6} = \beta_{W_2}. \end{aligned} \tag{65}$$

From (65) it follows that

$$\begin{aligned} a &= 2p_1(p_2\beta_{I_1} + p_3\beta_{D_1} + p_4\beta_{W_1}) \frac{r_1}{\beta_{W_1} S_1^0} q_4 \\ &\quad + 2p_5(p_6\beta_{I_2} + p_7\beta_{D_2} + p_8\beta_{W_2}) \frac{r_2}{\beta_{W_2} S_2^0} q_8 < 0. \end{aligned} \tag{66}$$

This excludes the possibility of a backward bifurcation since $a < 0$. For the sign of b , it is associated with the following non-vanishing partial derivatives of F :

$$\begin{aligned} \frac{\partial^2 f_1}{\partial x_2 \partial \beta^*} &= -\varrho_1 \beta_{D_1} x_1^0, & \frac{\partial^2 f_1}{\partial x_3 \partial \beta^*} &= -\beta_{D_1} x_1^0, & \frac{\partial^2 f_1}{\partial x_4 \partial \beta^*} &= -\varrho_2 \beta_{W_1} x_1^0, \\ \frac{\partial^2 f_2}{\partial \beta^* \partial x_2} &= \varrho_1 \beta_{I_1} x_1^0, & \frac{\partial^2 f_2}{\partial \beta^* \partial x_3} &= \beta_{D_1} x_1^0, & \frac{\partial^2 f_2}{\partial \beta^* \partial x_4} &= \varrho_2 \beta_{W_1} x_1^0, \\ \frac{\partial^2 f_5}{\partial x_6 \partial \beta^*} &= -\varrho_3 \beta_{I_2} x_2^0, & \frac{\partial^2 f_5}{\partial x_7 \partial \beta^*} &= -\beta_{D_2} x_2^0, & \frac{\partial^2 f_5}{\partial x_4 \partial \beta^*} &= -\varrho_2 \beta_{W_2} x_2^0, \\ \frac{\partial^2 f_6}{\partial \beta^* \partial x_6} &= \varrho_3 \beta_{I_2} x_2^0, & \frac{\partial^2 f_6}{\partial \beta^* \partial x_7} &= \beta_{D_2} x_2^0, & \frac{\partial^2 f_6}{\partial \beta^* \partial x_8} &= \varrho_4 \beta_{W_2} x_2^0. \end{aligned} \tag{67}$$

It follows from the expressions in (67), and after some algebraic manipulations, we have

$$b = (p_2\beta_{I_1} + p_3\beta_{D_1} + p_4\beta_{W_1}) \frac{r_1}{\beta_{W_1} x_1^0} q_4 + (p_6\beta_{I_2} + p_7\beta_{D_2} + p_8\beta_{W_2}) \frac{r_2}{\beta_{W_2} x_2^0} q_8 > 0. \tag{68}$$

Thus, $a < 0$ and $b > 0$ and applying Lemma 1 item (iv), we have established our result. The proof is complete. \square

Remark Suppose that β_{D_2} is chosen as a bifurcation parameter. By using the same approach as was used when β_{D_1} was taken as the bifurcation parameter in the proof of Theorem 6, we arrive at the same conclusion, that is, $a < 0$ and $b > 0$.

Acknowledgements

A. Mhlanga, the author, would like to thank the editors and the anonymous referees for their helpful comments and suggestions. He would also like to acknowledge with thanks the support of the Department of Mathematics, University of Zimbabwe.

Funding

None declared.

Availability of data and materials

The data used to support the findings of this study are included within the article and cited accordingly.

Ethics approval and consent to participate

None declared.

Competing interests

The author declares that they have no competing interests.

Consent for publication

None declared.

Authors' contributions

The author did all the work. All authors read and approved the final manuscript.

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Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Received: 23 February 2019 Accepted: 23 October 2019 Published online: 30 October 2019

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