

An optimal control problem for a metapopulation SIR model with vaccination

Gastón Beltritti, Carolina Bollo, Leopoldo Buri, Claudia Gariboldi,
Graciela Giubergia y Fernando Mazzone

November 2, 2022

Abstract

1 Introduction

The study of the dynamics of epidemics by means of mathematical models is an active area of scientific research [1, 3–5, 16]. The spread of diseases is an extremely complex social and biological process. Countless factors come into play, such as the habits of societies, the level of contact between individuals, public strategies for control and mitigation of epidemics, population structure, information campaigns, demographic changes, migrations, etc. Obviously, to this must be added, those factors based on the interaction between pathogens and the immune system and the evolutionary dynamics of the latter.

It is not easy, and perhaps not even useful, to build models that address all of these factors simultaneously. A complex model is more difficult to analyze mathematically and consequently it may not be easy to draw conclusions from it. On the other hand, the practical implementation of mathematical models requires having information about the object to be studied, for example model parameters. These parameters are usually estimated from the comparison between quantities that can be effectively observed and measured in reality and that in turn can be calculated by the model. This fit of the model to the observation may require information that is not available for a complex model.

The goals of mathematical modeling are also multiple. An issue in which this area of science can help is in providing criteria for decision making, fundamentally when these decisions involve the distribution of resources that are scarce.

In this article we are interested in the dynamics of epidemics in heterogeneous populations. Heterogeneity means that individuals do not behave in the same way with respect to the progress of the epidemic, and that the differences affect the dynamics of the process. Therefore, assuming homogeneity can lead to inaccuracies. The simplest example of this situation is when a disease is transmitted through a network of clusters of spatially separated individuals. It is known that the transmission of many diseases depends on the amount of contact between individuals. In a population separated into clusters, the rate of contact of an individual with others in the same cluster is, of course, to be

Me parece
que sería
bueno
mencionar
algo sobre
modelos
comparti-
mentados
y SIR

expected to be different from that with individuals in other clusters. But even in populations that are not spatially separated, it may be the case that individuals fulfill different roles within society, which would allow classify them into different groups with specific contact rates. The approach that we will follow in this article is known in the literature as the metapopulation models or models with patchy environments and movement between patches [4].

Vaccines are one of the most effective resources to mitigate the incidence of diseases. On some occasions, for example in a severe disease outbreaks, the supply of vaccines is limited and a decision must be made on how to distribute them. The objective of this article is to apply the mathematical theory of optimal control to the problem of determining the most efficient way to distribute a supply of vaccines within a metapopulation model. The global vaccination rate will be assumed to be a known function of time and the problem consists of how to distribute the vaccines in such a way as to minimize "cost". In this article we consider minimizing the total number of infected throughout the history of the epidemic.

We are going to briefly describe some scientific literature where problems similar to those dealt with in this article were addressed.

The so-called complex network epidemiological models [20,21] have received attention in the past. In these models each individual represents a node in a graph. The population is divided into compartments, composed of susceptible individuals, infected individuals, etc. In turn, the total population is divided according to the degree (number of neighbors) of the individual-nodes. The result is a model equivalent in certain respects to the one proposed in this article. The fact of structuring the population according to the degree of connectivity of individuals allows to analyze how this connectivity affects the spread of the disease. The analysis of control strategies (quarantine, vaccination, treatment, etc.) on complex networks has been studied in [9,13,15,19,22,26] to cite some background.

More specifically, in [2,6,7,17,23] problems of optimal control over networks-metapopulations were addressed. In [23] a optimal control problem for a metapopulation SIS model with two subpopulations was studied. The control strategies considered included treatment of the disease and restrictions on interpopulation circulation, quarantine controls that restrict the reciprocal rate of cross infection between the two regions. In [2] a problem similar to the one considered in this article was studied, i.e. a problem where vaccine doses must be distributed in a network. The objective function used in [2] accounts for the total number of individuals who were infected and the cost of the vaccination process. In [14] was studied an optimal control problem in a metapopulation SIQS quarantine controlled model. In [17] was considered optimal control in a metapopulation SIR model with treatment and state variable constraints. In [6,7] optimal control in a metapopulation SIR model with treatment and vaccination

One of the main differences between previous papers and this article is that we do not consider that access to the provision of vaccines is limited by their cost, but rather vaccine supply is limited. We assume given the quantity of vaccines per unit time for the entire network. This leads us to what in optimal control theory is called mixed variable state constraints. In [17,23] variable state constraint are introduced for treatment control optimal problems.

Es un super-borrador de introducción. Hay que trabajar mucho en mejorar la redacción y desarrollar un poca más.

2 The mathematical model

2.1 Description and consistency

By the consistency of the model we understand that the model presented has a solution, this is unique and that the model produces predictions consistent with the biological interpretation of the variables.

We will assume a population of N individuals with a birth rate equal to μ equal to death rate. In this regard, we will assume that the behavior of the population is homogeneous, that is, any subset of individuals in the population maintains the same birth and death rates. In this way the total population N is constant in time. Let's assume that this population is divided into n subpopulations according to a certain criterion (geographic, social activity, number of links in a graph of a complex network, etc.). Let us compartmentalize each subpopulation into susceptible, infected, immunized individuals, S_i , I_i and R_i respectively. We consider the following network-based SIR epidemic model:

$$\begin{cases} S'_i(t) = \mu N_i - S_i(t) \sum_{j=1}^n \beta_{ij} I_j(t) - (u_i(t) + \mu) S_i(t) + \alpha R_i(t) & (1) \\ I'_i(t) = S_i(t) \sum_{j=1}^n \beta_{ij} I_j(t) - (\mu + \gamma) I_i(t) & (2) \\ R'_i(t) = \gamma I_i(t) + u_i(t) S_i(t) - (\mu + \alpha) R_i(t) & (3) \end{cases}$$

where $1 = 1, \dots, n$ and $N_i = S_i + I_i + R_i$ is the total poblation of node i . The parameters in the model have the following meanings: β_{ii} , $i = 1, \dots, n$, β_{ij} , $i \neq j$ are the transmission rates within and between subpopulations, respectively; μ is the birth-death rate; $u_i(t)$ is the vaccination rate at the node i at moment t (control variable); γ^{-1} is the infectious period and α^{-1} is the immunity period.

We observe that summing, for each i , equations (1)-(3) we obtain $N'_i(t) = 0$, i.e. N_i is time independent. Consequently we can drop equation (3) in the system. Therefore, in this article we use the following state equations

$$\begin{cases} S'_i(t) = (\mu + \alpha) N_i - S_i(t) \left[\sum_{j=1}^n \beta_{ij} I_j(t) + u_i(t) + \mu + \alpha \right] - \alpha I_i(t) & (SIR_1) \\ I'_i(t) = S_i(t) \sum_{j=1}^n \beta_{ij} I_j(t) - (\mu + \gamma) I_i(t) & (SIR_2) \end{cases}$$

In order to address optimal control problems, the control variables u_i should not be assumed to be smooth functions. Hence, it is pertinent to explain in what sense the set of functions S_i, I_i and R_i , $i = 1, \dots, n$ are solutions of (SIR_1) -(SIR_2). Furthermore we have to discuss conditions under which problem (SIR_1) -(SIR_2) well-posed.

NUEVO:
Estamos
permi-
tiendo re-
infecciones

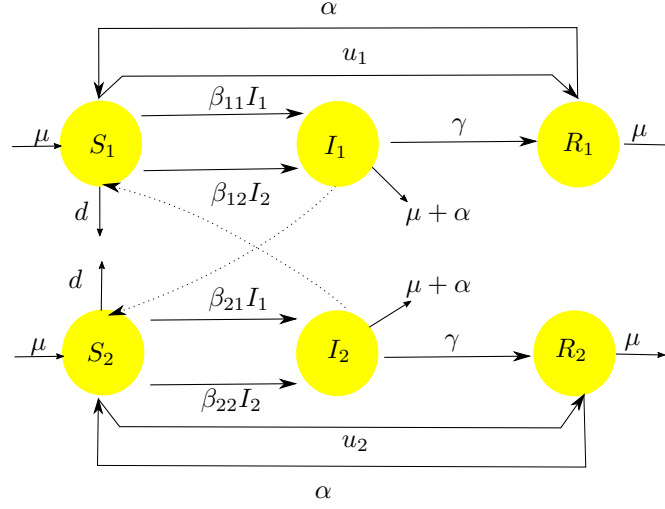


Figure 1: Flow chart model (1)-(3)

We will adopt the convention that bold symbols denote arrays objects like vectors and matrices. Thus, for example, β denotes matrix $\{\beta_{ij}\}_{i,j=1}^n$, $\mathbf{S} = (S_1, \dots, S_n)$ and so on. Given two arrays \mathbf{A} and \mathbf{B} we denote by $\mathbf{A} \times \mathbf{B}$ the element wise product of \mathbf{A} and \mathbf{B} . We write $\mathbf{x} = (\mathbf{S}, \mathbf{I})$ and

$$\begin{aligned} \mathbf{f}_S &= (\mu + \alpha)\mathbf{N} - \mathbf{S} \times (\beta\mathbf{I} + \mathbf{u}) - (\mu + \alpha)\mathbf{S} - \alpha\mathbf{I} \\ \mathbf{f}_I &= \mathbf{S} \times \beta\mathbf{I} - (\mu + \gamma)\mathbf{I}. \\ \mathbf{f} &= (\mathbf{f}_S, \mathbf{f}_I) \end{aligned}$$

Then the system $(SIR_1)-(SIR_2)$ is write in compact form $\mathbf{x}' = \mathbf{f}(\mathbf{x}, t)$. Given $\mathbf{x}_0 \in \mathbb{R}^{2n}$ we have the corresponding initial value problem

$$\begin{cases} \mathbf{x}' = \mathbf{f}(\mathbf{x}, t). \\ \mathbf{x}(0) = \mathbf{x}_0 \end{cases} \quad (E)$$

One aspect that should be noted and that is often neglected in the literature is the fact that the vector field \mathbf{f} is not continuous. This is due to the fact that, for the purposes of posing optimal control problems, the functions u_i are generally assumed to be only measurable. Fortunately, we can deal with this type of equations by means of Carathéodory theory [8, 11] of differential equations.

Following [11] we said that \mathbf{x} is a *solution* of (E) on an open interval $[0, a)$ if and only if \mathbf{x} is absolutely continuous on each subinterval $[0, \alpha] \subset [0, a)$, satisfies initial condition and $\mathbf{x}' = \mathbf{f}(\mathbf{x}, t)$ for almost everywhere $t \in (0, b)$. Alternatively, it is equivalent to say that \mathbf{x} satisfies the integral equation

$$\mathbf{x}(t) = \mathbf{x}_0 + \int_0^t \mathbf{f}(\mathbf{x}, s) ds. \quad (4)$$

We note that if $u_i \in L^1([0, +\infty))$ then \mathbf{f} is a Carathéodory function in the sense of [11, p. 3] on the domain $\Omega \times [0, +\infty)$, where Ω is an open and bounded set in \mathbb{R}^n . That means, $\mathbf{f}(\mathbf{x}, t)$ is continuous in \mathbf{x} for almost everywhere $t \in [0, +\infty)$, $\mathbf{f}(\mathbf{x}, t)$ is measurable in t for each $\mathbf{x} \in \Omega$ and there exists $m \in L^1([0, +\infty))$ such that $|\mathbf{f}(\mathbf{x}, t)| \leq m(t)$ for every $(\mathbf{x}, t) \in \Omega \times [0, +\infty)$. Consequently we can apply existence and uniqueness theory for Carathéodory equations [11, Th. 1, p. 4, Th. 2, p. 5], [8, Th. 1.1], we conclude that for any $\mathbf{x}_0 \in \Omega$ there exists a unique local solutions of the boundary value problem (E) defined in some interval $[0, \delta)$, $\delta > 0$.

We will use the following well known and elementary formula for the solution of a linear scalar equation of first order $x'(t) + p(t)x(t) = q(t)$:

$$x(t) = e^{-\int_0^t p(s)ds} \left\{ x(0) + \int_0^t e^{\int_0^s p(r)dr} q(s)ds \right\}. \quad (5)$$

For $i = 1, \dots, n$ it is convenient to set

$$\Theta_i(t) := \sum_{j=1}^n \beta_{ij} I_j, \quad \nu = \mu + \alpha.$$

Then using formula (5) we obtain that equations (SIR_1) -(SIR_2) are equivalent to the integral equations

$$\begin{aligned} S_i(t) = \exp \left\{ -\nu t - \int_0^t \Theta_i + u_i ds \right\} S_i(0) \\ + \int_0^t \exp \left\{ \nu(s-t) + \int_t^s \Theta_i + u_i dr \right\} (\nu N_i - \alpha I_i) ds \end{aligned} \quad (6)$$

$$I_i(t) = \exp(-(\mu + \gamma)t) \left\{ I_i(0) + \int_0^t \exp((\mu + \gamma)s) \Theta_i S_i ds \right\} \quad (7)$$

Equations (6)-(7) have the advantage that they are valid for every $t \geq 0$ while equations (SIR_1) -(SIR_2) holds true almost everywhere.

Although from a mathematical point of view the problem (E) makes sense for any \mathbf{x} it is only relevant for our model in the closed set

$$\Omega_0 = \{ \mathbf{x} \in \mathbb{R}^{2n} | 0 \leq x_i, x_{n+i} \leq N_i, i = 1, \dots, n \}.$$

Since $\nu N_i - \alpha I_i \geq 0$, we obtain from equations (6)-(7) that $S_i(t), I_i(t) \geq 0$ for every t where solutions are defined. On the other hand, $S_i(t) + I_i(t) \leq N_i$. Consequently, the set Ω is flow invariant for the equation (E). If Ω is any bounded an open set containing Ω_0 and $\mathbf{x}_0 \in \Omega_0$ we have that the solution $\mathbf{x}(t)$ of (E) remains inside Ω for every $t > 0$ where solution is defined. We infer using continuation theory [8, Th. 1.3] that the solition is defined for every positive time.

2.2 Propagation on the network

Desarrollaría un poco más, por ejemplo incluir definición de irreducibilidad, ver por ejemplo [18] o [27]. Considerar el modelos general, demografía y re-infecciones

We assume $\mu = d = \alpha = 0$ and $\delta = 1$. In this situation $N'_i := (S_i + I_i + R_i)' = 0$, i.e. the total population of node i remain constant. Cosequently $R_i = N - S_i - I_i$ and we can drop the equation for R_i of the system. Therefore, we can study the SIR metapopulation model with vaccination (SIRmv) system.

$$\begin{cases} S'_i = -S_i(t) \sum_{j=1}^n \beta_{ij} I_j(t) - u_i(t) S_i(t) & (\text{SIRmv1}) \\ I'_i = S_i(t) \sum_{j=1}^n \beta_{ij} I_j(t) - \gamma I_i(t) & (\text{SIRmv2}) \end{cases}$$

We note that the system of equations (SIR_1) , (SIR_1) possibly has a discontinuous right hand side. In this case, following to [11]. **Poner el significado de solución**

From (SIR_1) and (5) we obtain that

$$S_i(t) = S_i(0) \exp \left(- \int_0^t \sum_{j=1}^n \beta_{ij} I_j(s) + u(s) ds \right)$$

The following conjecture establish that if is a irreducible matrix (see [18]) the epidemic is transmitted from any node to the rest with infinite speed.

Proposition 2.1. *Suppose that β is a irreducible matrix. If there exists i such that $I_i(0) > 0$ then for every j and $t > 0$ we have $I_j(t) > 0$.*

Proof.

Hay que repensar la demostración para admitir soluciones que no son derivables. Hay que utilizar ecuaciones (6) -(7). Hagamos la demostración para el modelo full- demografía y reinfecciones. Ojo! hay un caso excepcional, cuando no hay demografía ($\mu = 0$), no hay reinfección ($\alpha = 0$) e inicialmente todos son recuperados $R_i(0) = N_i$

Let $i, j \in \{1, 2, \dots, n\}$ such that $I_i(0) > 0$ and $i \neq j$.

Suppose first that $\beta_{ij} > 0$. Let's fix $\epsilon > 0$ and see that $I_j(t)$ is positive for all $t > \epsilon$. As $I_i(0) > 0$ then $I_i(\epsilon) > 0$. For $t > \epsilon$ we have that

$$I_j(t) = e^{-\int_\epsilon^t (\gamma - \beta_{jj}) ds} \left(I_j(\epsilon) + \int_\epsilon^t e^{\int_\epsilon^s (\gamma - \beta_{jj}) dz} \left[S_j(s) \sum_{k=1, k \neq j}^n \beta_{jk} I_k(s) \right] ds \right).$$

Let's assume that $I_j(\epsilon) > 0$, then, as the functions S_k and I_k are non negative, we have that $I_j(t) > 0$ for $t > \epsilon$. If we suppose that $I_j(\epsilon) = 0$, by Taylor's theorem, we can see that

$$\begin{aligned} I_j(t) &= I_j(\epsilon) + I'_j(\epsilon)(t - \epsilon) + h_{j,\epsilon}(t) \\ &= I_j(\epsilon) + S_j(\epsilon) \sum_{k=1}^n \beta_{jk} I_k(\epsilon)(t - \epsilon) - \gamma I_j(\epsilon)(t - \epsilon) + h_{j,\epsilon}(t) \\ &\geq I_j(\epsilon) + S_j(\epsilon) \beta_{ji} I_i(\epsilon)(t - \epsilon) - \gamma I_j(\epsilon)(t - \epsilon) + h_{j,\epsilon}(t) \\ &= I_j(\epsilon) + N_j \beta_{ji} I_i(\epsilon)(t - \epsilon) + h_{j,\epsilon}(t). \end{aligned}$$

Thus

$$\frac{I_j(t) - I_j(\epsilon)}{t - \epsilon} \geq N_j \beta_{ji} I_i(\epsilon) + \frac{h_{j,\epsilon}(t)}{t - \epsilon},$$

and therefore $I_j'(\epsilon) \geq N_j \beta_{ji} I_i(\epsilon) > 0$, which implies that there exists $\epsilon_1 > \epsilon$ such that if $t \in (\epsilon, \epsilon_1]$ then $I_j(t) > 0$. Now, reasoning analogously as we did for the case where $I_j(\epsilon) > 0$, as $I_j(\epsilon_1) > 0$ then $I_j(t) > 0$ for every $t > \epsilon_1$. Thus, $I_j(t) > 0$ for all $t > \epsilon$. Since ϵ is an arbitrary positive number we have that $I_j(t) > 0$ for every $t > 0$.

Suppose now that $\beta_{ij} = 0$. Since the matrix β is irreducible, we have that there exist $j_1, j_2, \dots, j_l \in \{1, 2, \dots, n\}$ such that $\beta_{j_1 i} > 0$, $\beta_{j_r j_{r-1}} > 0$ for $r = 2, 3, \dots, l$, and $\beta_{j_l i} > 0$. Reasoning in the same way as we did in the previous paragraph we can see that, $I_{j_1}(t) > 0$ for all $t > 0$, which implies that $I_{j_2}(t) > 0$ for every $t > 0$ (note that is not necessary that $I_{j_1}(0) > 0$ to guarantee this fact), this last statement assure us that $I_{j_2}(0) > 0$, and continuing in this way we can see that $I_j(t) > 0$ for all $t > 0$. □

2.3 The number of vaccines for $t \rightarrow +\infty$

The goal of this section is to show that, in some sense, the total number of vaccinated individuals $u_i S_i$ for any i approaches zero as $t \rightarrow +\infty$. This regardless of the vaccination campaign. A very aggressive campaign, so that the vaccination rate u_i are high, will imply that the number of susceptible individuals converges to zero fast enough so that the total number of vaccinated $u_i S_i$ also tends to zero.

What we have actually shown is that certain regularizations of the functions $u_i S_i$ converge to zero at infinity. To state the result that we want to present, let us recall the concept of *regularization* or *approximation of identity* [25]. Let φ be an integrable function on \mathbb{R} satisfying $\int_{\mathbb{R}} \varphi(t) dt = 1$. In this article we will only consider functions φ with $\varphi(t) = 0$ when $t \leq 0$ and φ decreasing on $[0, +\infty)$. For $\gamma > 0$, we set $\varphi_\gamma(t) = \gamma \varphi(\gamma t)$ and for any $f \in L^1(\mathbb{R})$ with $f(t) \equiv 0$ for $t < 0$, we write

$$f_\gamma(t) = f * \varphi_\gamma(t) := \gamma \int_{-\infty}^{\infty} f(t-s) \varphi(\gamma s) ds = \gamma \int_0^t f(t-s) \varphi(\gamma s) ds.$$

Then it is well known [25] that f_γ are continuous and $f_\gamma \rightarrow f$ almost everywhere and in L^p -norm.

Suppose that $f \in L^\infty(\mathbb{R})$ and that there exists $\lim_{t \rightarrow \infty} f(t)$. Then, using the Lebesgue's dominated converge theorem

$$\lim_{t \rightarrow \infty} f_\gamma(t) = \lim_{t \rightarrow \infty} f(t). \tag{8}$$

Remark 1. Of course, the existence of the limit on the left hand side in equation (8) does not guarantee the existence of limit of $f(t)$ for $t \rightarrow \infty$. An example of this fact is shown uin appendix ?????.

The following proposition to expresses the fact that in the case when there are no demographic changes ($\mu = 0$) and reinfections ($\alpha = 0$) the epidemic is extinguished when $t \rightarrow \infty$ and, in certain sense, the total quantity of applied vaccines $u_i S_i$ goes to zero when $t \rightarrow \infty$.

Proposition 2.2. *We assume that $\mu = \alpha = 0$ and that $\varphi(t) = e^{-t}$, when $t > 0$ and $\varphi(t) = 0$ otherwise. The regularized function $(u_i S_i)_\gamma$ of the total number of appiled vaccines per unit of time goes to zero when $t \rightarrow \infty$. More concretely*

$$\lim_{t \rightarrow \infty} \gamma \int_0^t u_i(s) S_i(s) e^{\gamma(s-t)} ds = I_i(\infty) = 0.$$

Proof. Adding equations (SIR_1) and (SIR_2) we obtain

$$(S_i + I_i)' = -u_i S_i - \gamma I_i \leq 0.$$

Therefore $S_i + I_i$ is a monotone non increasing function. Hence $\lim_{t \rightarrow \infty} (S_i + I_i)$ there exists. From (SIR_1) the same considerations are true for function S_i . Consequently there exists $\lim_{t \rightarrow \infty} S_i(t) =: S_i(\infty)$. We deduce that there exists $\lim_{t \rightarrow \infty} I_i(t) =: I_i(\infty)$. If $I(\infty) > 0$, we could choose t_0 large enough for that $t \geq t_0$ implies $I_i(t) > I_i(\infty)/2 =: a > 0$. Then $(S_i(t) + I_i(t))' \leq -\gamma I_i(t) \leq -\gamma a$. This inequality implies that $S_i(t) + I_i(t) \rightarrow -\infty$, when $t \rightarrow \infty$, which is a contradiction. Consequently $I_i(\infty) = 0$.

From (7) we obtain

$$\begin{aligned} I_i(t) &= e^{-\gamma t} \left\{ I_i(0) + \int_0^t e^{\gamma s} S_i(s) \Theta_i(s) ds \right\} \\ &= e^{-\gamma t} \left\{ I_i(0) - \int_0^t e^{\gamma s} [S_i'(s) + u_i(s) S_i(s)] ds \right\} \\ &= e^{-\gamma t} \left\{ I_i(0) - \int_0^t e^{\gamma s} u_i(s) S_i(s) ds - e^{\gamma t} S_i(t) + S_i(0) - \gamma \int_0^t e^{\gamma s} S_i(s) ds \right\} \\ &= e^{-\gamma t} (S_i(0) + I_i(0)) - S_i(t) + \gamma \int_0^t e^{\gamma(s-t)} S_i(s) ds - \int_0^t e^{\gamma(s-t)} u_i(s) S_i(s) ds \end{aligned}$$

The proof is completed by taking limit for $t \rightarrow \infty$ in previous identities and using (8). \square

Conjecture 2.3.

$$\lim_{t \rightarrow \infty} u_i(s) S_i(s) = 0.$$

3 Optimal control problem

We suppose $T > 0$ a fixed time and that $M : [0, T] \rightarrow \mathbb{R}_+$ certain given non-negative function. We consider the admissible control set

$$\mathcal{U} = \{u = (u_1, \dots, u_n) : u_i \text{ measurable}, u_i \geq 0, u_1 S_1 + \dots + u_n S_n \leq M(t)\}.$$

Here, $M(t)$ represent the number of vaccines per unit of time for the total population.

The objective function given by

$$J(u) = \int_0^T \sum_{i=1}^n I_i(t) dt$$

we formulate the optimal control problem

$$\text{find } u^* \in \mathcal{U} \text{ such that } J(u^*) = \min_{u \in \mathcal{U}} J(u) \quad (9)$$

4 Existence minimizers

In this Section, we prove that the optimal control problem (9) has a solution. That is, we prove that the hypothesis of the Filippov-Cesari Theorem are satisfied (see [24]). In what follows, we will use the following notation

$$x = (S_1, \dots, S_n, I_1, \dots, I_n, R_1, \dots, R_n)$$

$$f_0(x_i, u_i, t) = f_0(S_1, \dots, S_n, I_1, \dots, I_n, R_1, \dots, R_n; u_1, \dots, u_n; t) = \sum_{i=1}^n I_i(t).$$

For $i = 1, 2, \dots, n$, we denote by

$$f_i = (f_{1i}, f_{2i}, f_{3i}), \quad f = (f_1, \dots, f_n)$$

with

$$f_{1i} = \mu N_i - S_i(t) \sum_{j=1}^n \beta_j I_j(t) - \delta u_i S_i(t) - d S_i(t)$$

$$f_{2i} = S_i(t) \sum_{j=1}^n \beta_j I_j(t) - (d + \alpha + \gamma) I_i(t)$$

$$f_{3i} = \gamma I_i(t) + \delta u_i S_i(t) - d R_i(t)$$

and we define

$$N(x, \mathcal{U}, t) = \{(f_0 + \gamma, f) : \gamma \geq 0, u \in \mathcal{U}\}.$$

Now, we will be in conditions to prove the following result.

Theorem 4.1. *The optimal control problem (9) has a solution $u^* \in \mathcal{U}$.*

Proof. We prove that $N(x, \mathcal{U}, t)$ is a convex set, for all (x, t) .

Let $(a_1, b_1), (a_2, b_2) \in N(x, \mathcal{U}, t)$ be, then there exist $\gamma_1, \gamma_2 \geq 0$ and $u_1, u_2 \in \mathcal{U}$ such that

$$(f_0(x, u_1, t) + \gamma_1; f(x, u_1, t)) = (a_1, b_1)$$

and

$$(f_0(x, u_2, t) + \gamma_2; f(x, u_2, t)) = (a_2, b_2)$$

then

$$\lambda(a_1, b_1) + (1 - \lambda)(a_2, b_2) = (\lambda a_1 + (1 - \lambda)a_2, \lambda b_1 + (1 - \lambda)b_2)$$

$$= (\lambda(f_0(x, u_1, t) + \gamma_1) + (1 - \lambda)(f_0(x, u_2, t) + \gamma_2), \lambda f(x, u_1, t) + (1 - \lambda)f(x, u_2, t)).$$

Now, we consider the second component

$$\lambda f(x, u_1, t) + (1 - \lambda)f(x, u_2, t)$$

and from the linearity of f with respect u , we have

$$\lambda f(x, u_1, t) + (1 - \lambda)f(x, u_2, t) = f(x, \lambda u_1 + (1 - \lambda)u_2, t)$$

Demostrar el teorema de existencia para n nodos y con reinfecciones y demografía

Moreover, $\bar{u} = \lambda u_1 + (1 - \lambda)u_2 \in \mathcal{U}$. In fact, $\lambda u_1 + (1 - \lambda)u_2$ is measurable, $\lambda u_1 + (1 - \lambda)u_2 \geq 0$ and if we consider $u_1 = (u_{11}, \dots, u_{1n})$ and $u_2 = (u_{21}, \dots, u_{2n})$ in \mathcal{U} , then

$$\lambda u_1 + (1 - \lambda)u_2 = (\lambda u_{11} + (1 - \lambda)u_{21}, \dots, \lambda u_{1n} + (1 - \lambda)u_{2n})$$

next

$$\begin{aligned} & (\lambda u_{11} + (1 - \lambda)u_{21}) S_1 + \dots + (\lambda u_{1n} + (1 - \lambda)u_{2n}) S_n = \\ & \lambda (u_{11} S_1 + \dots + u_{1n} S_n) + (1 - \lambda) (u_{21} S_1 + \dots + u_{2n} S_n) \leq u_{Tot} \end{aligned}$$

therefore $\lambda u_1 + (1 - \lambda)u_2 \in \mathcal{U}$.

Now, we prove that there exists $\gamma \geq 0$ such that

$$f_0(x, \bar{u}, t) + \gamma = \lambda a_1 + (1 - \lambda)a_2.$$

We note that f_0 is constant with respect to the control variable, then

$$f_0(x, \lambda u_1 + (1 - \lambda)u_2, t) = \lambda f_0(x, u_1, t) + (1 - \lambda)f_0(x, u_2, t).$$

If we define $\gamma = \lambda\gamma_1 + (1 - \lambda)\gamma_2 \geq 0$, we have that

$$\begin{aligned} f_0(x, \lambda u_1 + (1 - \lambda)u_2, t) + \gamma &= [\lambda f_0(x, u_1, t) + (1 - \lambda)f_0(x, u_2, t)] + [\lambda\gamma_1 + (1 - \lambda)\gamma_2] \\ &= \lambda f_0(x, u_1, t) + \lambda\gamma_1 + (1 - \lambda)f_0(x, u_2, t) + (1 - \lambda)\gamma_2 \\ &= \lambda (f_0(x, u_1, t) + \gamma_1) + (1 - \lambda) (f_0(x, u_2, t) + \gamma_2) \\ &= \lambda a_1 + (1 - \lambda)a_2. \end{aligned}$$

Therefore, we proved that there exists $\gamma = \lambda\gamma_1 + (1 - \lambda)\gamma_2 \geq 0$ and there exists $\bar{u} = \lambda u_1 + (1 - \lambda)u_2 \in \mathcal{U}$ such that

$$(\lambda a_1 + (1 - \lambda)a_2, \lambda b_1 + (1 - \lambda)b_2) = (f_0(x, \bar{u}, t) + \gamma, f(x, \bar{u}, t)),$$

i.e.

$$(\lambda a_1 + (1 - \lambda)a_2, \lambda b_1 + (1 - \lambda)b_2) \in N(x, \mathcal{U}, t)$$

and $N(x, \mathcal{U}, t)$ is a convex set, for all fixed (x, t) .

Moreover, \mathcal{U} is a compact set, since $0 \leq u_i \leq u_{Tot}$, $\forall i = 1, \dots, n$. Finally, taking into account that the number of susceptible, infected and removed individuals are bounded by the total quantity of individuals, we have that $\|x(t)\| \leq b$. Therefore, we have verified the hypothesis of Filippov-Cesari Existence Theorem and the thesis holds. \square

5 Necessary conditions

In this Section, we apply the Pontryagin Maximum Principle [24, Th. 4.1], [12] for to obtain necessary conditions in order to $(x^*, u^*) = (S^*, I^*, u^*)$ be a solution of the optimal problem

$$\left\{ \begin{array}{ll} \min_u \int_0^T \sum_{i=1}^n I_i(t) dt & (\text{OP}_1) \\ \text{s. t.} & \\ \mathbf{S}' = \mathbf{f}_S(\mathbf{S}, \mathbf{I}, \mathbf{u}) = \nu \mathbf{N} - \mathbf{S} \times (\beta \mathbf{I} + \mathbf{u}) - \nu \mathbf{S} - \alpha \mathbf{I} & (\text{OP}_2) \\ \mathbf{I}' = \mathbf{f}_I(\mathbf{S}, \mathbf{I}, \mathbf{u}) = \mathbf{S} \times \beta \mathbf{I} - (\mu + \gamma) \mathbf{I}. & (\text{OP}_3) \\ \mathbf{S}(0) = \mathbf{S}_0, \quad \mathbf{I}(0) = \mathbf{I}_0 & (\text{OP}_4) \\ \mathbf{S}(T), \quad \mathbf{I}(T), \quad \text{free} & \\ \mathbf{u} \cdot \mathbf{S} \leq M(t) & (\text{OP}_5) \\ \mathbf{u} \geq 0 & (\text{OP}_6) \end{array} \right.$$

As is well known the maximum principle involves the Hamiltonian formulation of the optimal problem. We consider *adjoint variables*, $\mathbf{p}_S(t) = (p_{S1}(t), \dots, p_{Sn}(t))$ and $\mathbf{p}_I(t) = (p_{I1}(t), \dots, p_{In}(t))$ and the Hamiltonian function defined by

$$\mathcal{H}(\mathbf{S}, \mathbf{I}, \mathbf{u}, \mathbf{p}_S, \mathbf{p}_I) = -p_0 \sum_{i=1}^n I_i + \mathbf{p}_S \cdot \mathbf{f}_S(\mathbf{S}, \mathbf{I}, \mathbf{u}) + \mathbf{p}_I \cdot \mathbf{f}_I(\mathbf{S}, \mathbf{I}, \mathbf{u}), \quad (11)$$

where $p_0 \in \mathbb{R}$ is independent of t . It is necessary to introduce new multipliers $q_0(t)$ and $\mathbf{q}(t) = (q_1(t), \dots, q_n(t))$ associated with each constraint and the *Lagrangian* or *generalized Hamiltonian* \mathcal{L} defined by

$$\mathcal{L}(\mathbf{S}, \mathbf{I}, \mathbf{u}, \mathbf{p}_S, \mathbf{p}_I, \mathbf{q}) = \mathcal{H}(\mathbf{S}, \mathbf{I}, \mathbf{u}, \mathbf{p}_S, \mathbf{p}_I) + \mathbf{q} \cdot \mathbf{u} - q_0 \mathbf{u} \cdot \mathbf{S}. \quad (12)$$

We suppose that $(\mathbf{S}^*, \mathbf{I}^*, \mathbf{u}^*)$ is a solution of (OP). From now on any function evaluated at $(\mathbf{S}^*, \mathbf{I}^*, \mathbf{u}^*)$ will be indicated with a $*$ as superscript.

We combine the restrictions in a vector valued function

$$\mathbf{h}(\mathbf{S}, \mathbf{u}, t) = (\mathbf{u}_1, \dots, \mathbf{u}_n, -\mathbf{u} \cdot \mathbf{S} + M(t)).$$

Then, the constraints are expressed synthetically $\mathbf{h} \geq \mathbf{0}$, where order relations between vectors mean that these relations are given component by component. Following to be able to apply the Maximum Principle it is necessary that they satisfy the According to [24, Th. 4.1], see also [12], in order to apply the maximum principle, the constraint qualification condition must be satisfied. This condition means that the gradients of the active constraints (this is $\{i : h_i = 0\}$) are linearly. In our case, the following condition must be satisfied

$$n+1 = \text{rank} \left[\begin{array}{ccc|ccc} 1 & \cdots & 0 & u_1 & \cdots & 0 & 0 \\ \vdots & \ddots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & \cdots & 1 & 0 & \cdots & u_n & 0 \\ -S_1 & \cdots & -S_n & 0 & \cdots & 0 & -\mathbf{u} \cdot \mathbf{S} + M(t) \end{array} \right] \quad (13)$$

We note that if $-\mathbf{u} \cdot \mathbf{S} + M(t) \neq 0$ then then qualification condition follows immediately. In the case that $-\mathbf{u} \cdot \mathbf{S} + M(t) = 0$ (last constraint active) we can

prove that the matrix en the rhs of (5) can be row reduced to

$$\left[\begin{array}{ccc|ccc} 1 & \cdots & 0 & u_1 & \cdots & 0 & 0 \\ \vdots & \ddots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & \cdots & 1 & 0 & \cdots & u_n & 0 \\ 0 & \cdots & 0 & u_1 S_1 & \cdots & u_n S_n & 0 \end{array} \right].$$

Since $-\mathbf{u} \cdot \mathbf{S} + M(t) = 0$, there is some i with $u_i S_i \neq 0$. This implies that matrix has completed rank.

We are in a position to apply the maximum principle. First we observe that in virtue of equations (29), (34) and (35c) in [24, Th. 4.1] we can assume that $p_0 = 1$ in (11). On the other hand, we infer that there exist functions $\mathbf{p}_S^*, \mathbf{p}_I^* : [0, T] \rightarrow \mathbb{R}^n$ and *multipliers* $\mathbf{q} = (q_0(t), \dots, q_n(t)) \in \mathbb{R}^{n+1}$ such that the following conditions are satisfied

1. *Adjoint equations*

$$\begin{cases} \dot{\mathbf{p}}_S = -\frac{\partial \mathcal{L}^*}{\partial \mathbf{S}} & = \mathbf{p}_S \times (\beta \mathbf{I}^* + \mathbf{u}^* + \nu) - \mathbf{p}_I \times \beta \mathbf{I} + q_0 \mathbf{u}^*, & (\text{ADJ}_1) \\ \dot{\mathbf{p}}_I = -\frac{\partial \mathcal{L}^*}{\partial \mathbf{I}} = p_0 \mathbf{1} + \beta^t (\mathbf{S}^* \times (\mathbf{p}_S - \mathbf{p}_I)) + \alpha \mathbf{p}_S + (\mu + \gamma) \mathbf{p}_I, & (\text{ADJ}_2) \end{cases}$$

where $\mathbf{1} = (1, \dots, 1) \in \mathbb{R}^n$.

2.

$$0 = \frac{\partial \mathcal{L}^*}{\partial \mathbf{u}} = -\mathbf{p}_S \times \mathbf{S}^* + \mathbf{q} - q_0 \mathbf{S}^* \quad (15)$$

3. *Positivity and complementary slackness conditions*

$$\mathbf{q}(t) \geq 0 \quad \text{and} \quad \mathbf{q}(t) \cdot \mathbf{h}(\mathbf{S}^*, \mathbf{u}^*, t) = 0 \quad (16)$$

4. *Terminal condition*

$$p_0 = 1 \quad \text{and} \quad \mathbf{p}_S(T) = \mathbf{p}_I(T) = 0. \quad (17)$$

5. *Maximality.* For fix $t \in [0, T]$ and for every \mathbf{u} with $\mathbf{h}(\mathbf{S}^*, \mathbf{u}, t) \geq \mathbf{0}$ we have

$$\mathcal{H}(\mathbf{S}^*, \mathbf{I}^*, \mathbf{u}^*, \mathbf{p}_S, \mathbf{p}_I) \geq \mathcal{H}(\mathbf{S}^*, \mathbf{I}^*, \mathbf{u}, \mathbf{p}_S, \mathbf{p}_I) \quad (18)$$

Corollary 5.1. *We suppose that $(\mathbf{S}^*, \mathbf{I}^*, \mathbf{u}^*)$ is a solution of (OP). Let $F(t) \subset \{1, \dots, n\}$ be the subset of indices defined by*

$$j \in F(t) \iff \mathbf{p}_{S^*j}(t) = p_{\min}(t) := \min\{\mathbf{p}_{S^*1}(t), \dots, \mathbf{p}_{S^*n}(t)\}.$$

Then

$$u_k^*(t) = 0, \quad \text{for every } k \notin F(t). \quad (19)$$

Proof. We observe that the Hamiltonian can be written

$$\mathcal{H}(\mathbf{S}, \mathbf{I}, \mathbf{u}, \mathbf{p}_S, \mathbf{p}_I) = \mathcal{H}_0(\mathbf{S}, \mathbf{I}, \mathbf{p}_S, \mathbf{p}_I) - \mathbf{p}_S \cdot (\mathbf{S} \times \mathbf{u}). \quad (20)$$

Therefore (20) implies that for each $t \in [0, T]$ and for every \mathbf{u} with $\mathbf{u} \geq \mathbf{0}$ and $\mathbf{u} \cdot \mathbf{S}^* \leq M(t)$ we have

$$\mathbf{p}_S \cdot (\mathbf{S}^* \times \mathbf{u}^*) \leq \mathbf{p}_S \cdot (\mathbf{S}^* \times \mathbf{u}) \quad (21)$$

Therefore for every \mathbf{u} with $\mathbf{u} \geq \mathbf{0}$ and $\mathbf{u} \cdot \mathbf{S}^* \leq M(t)$

$$\mathbf{p}_S \cdot (\mathbf{S}^* \times \mathbf{u}) \geq p_{\min}(t)M$$

Suppose that there exists $k \notin F(t)$ such that $u_k^* > 0$

□

6 Numerical results

.....

7 appendix

$$h = \sum_{n=0}^{\infty} \mathbb{1}_{[n, n+\frac{1}{2}]},$$

where $\mathbb{1}_A$ denotes the characteristic function of the set A . Then $h(s) + h(s - \frac{1}{2}) = 1$, for $s \in [0, +\infty]$. We define $\varphi(s) = h(e^s - 1)$. Then

$$\begin{aligned} 1 &= \lim_{t \rightarrow \infty} \gamma \int_0^t \left(\varphi(s) + \varphi\left(s - \frac{1}{2}\right) \right) e^{\gamma(s-t)} ds \\ &= \lim_{t \rightarrow \infty} \gamma \int_0^t \varphi(s) e^{\gamma(s-t)} ds + \gamma \int_0^t \varphi\left(s - \frac{1}{2}\right) e^{\gamma(s-t)} ds \quad (22) \end{aligned}$$

On the other hand

$$\begin{aligned} \gamma \int_0^t \varphi\left(s - \frac{1}{2}\right) e^{\gamma(s-t)} ds &= \gamma e^{\frac{\gamma}{2}} \int_0^{t-\frac{1}{2}} \varphi(r) e^{\gamma(r-t)} dr \\ &= \gamma e^{\frac{\gamma}{2}} \int_0^t \varphi(r) e^{\gamma(r-t)} dr - \gamma e^{\frac{\gamma}{2}} \int_{t-\frac{1}{2}}^t \varphi(r) e^{\gamma(r-t)} dr \quad (23) \end{aligned}$$

In my understanding

$$\lim_{t \rightarrow \infty} \int_{t-\frac{1}{2}}^t \varphi(r) e^{\gamma(r-t)} dr = \frac{1}{2\gamma} (1 - e^{-\gamma/2}).$$

Taking account of (22), (23) we infer that

$$\lim_{t \rightarrow \infty} \gamma \int_0^t \varphi(s) e^{\gamma(s-t)} ds = \frac{e^{\gamma/2}}{2(1 + e^{\gamma/2})}$$

References

- [1] Linda S. Allen. *Stochastic Population and Epidemic Models: Persistence and Extinction*, volume 1. Springer International Publishing, sep 2015.
- [2] Erika Asano, Louis J Gross, Suzanne Lenhart, and Leslie A Real. Optimal control of vaccine distribution in a rabies metapopulation model. *Mathematical Biosciences & Engineering*, 5(2):219, 2008.

- [3] Fred Brauer and Carlos Castillo-Chavez. *Mathematical Models in Population Biology and Epidemiology*, volume 1. Springer Science & Business Media, mar 2013.
- [4] Fred Brauer, Carlos Castillo-Chavez, and Zhilan Feng. *Mathematical Models in Epidemiology*, volume 1. Springer Nature, oct 2019.
- [5] Fred Brauer, Pauline Den van Driessche, and J. Wu. *Mathematical Epidemiology*, volume 1. Springer Science & Business Media, abr 2008.
- [6] Jitao Chen, Lijuan; Sun. Global stability and optimal control of an sirs epidemic model on heterogeneous networks. *Physica A: Statistical Mechanics and its Applications*, 410, 09 2014.
- [7] Jitao Chen, Lijuan; Sun. Optimal vaccination and treatment of an epidemic network model. *Physics Letters A*, 378, 08 2014.
- [8] Earl A. Coddington, A Coddington Earl, Coddington Mn, and Norman Levinson. *Theory of Ordinary Differential Equations*. McGraw-Hill, ago 1955.
- [9] Jose de Jesus Esquivel-Gómez and Juan Gonzalo Barajas-Ramírez. Efficiency of quarantine and self-protection processes in epidemic spreading control on scale-free networks. *Chaos: An Interdisciplinary Journal of Nonlinear Science*, 28(1):013119, Jan 2018.
- [10] Lawrence Evans. An introduction to mathematical optimal control theory version 0.2, 1985. Lecture Notes.
- [11] A. F. Filippov. *Differential Equations With Discontinuous Righthand Sides: Control Systems*. Springer Science & Business Media, sep 1988.
- [12] Richard F Hartl, Suresh P Sethi, and Raymond G Vickson. A survey of the maximum principles for optimal control problems with state constraints. *SIAM review*, 37(2):181–218, 1995.
- [13] Huiyan Kang, Kaihui Liu, and Xinchu Fu. Dynamics of an epidemic model with quarantine on scale-free networks. *Physics Letters A*, 381(47):3945–3951, 2017.
- [14] Kezan Li, Guanghu Zhu, Zhongjun Ma, and Lijuan Chen. Dynamic stability of an siqs epidemic network and its optimal control. *Communications in Nonlinear Science and Numerical Simulation*, 66, 01 2019.
- [15] N. Madar, T. Kalisky, R. Cohen, D. ben Avraham, and S. Havlin. Immunization and epidemic dynamics in complex networks. *The European Physical Journal B - Condensed Matter*, 38(2):269–276, Mar 2004.
- [16] Maia Martcheva. *An Introduction to Mathematical Epidemiology*, volume 1. Springer, oct 2015.
- [17] Martial L Ndeffo Mbah and Christopher A Gilligan. Resource allocation for epidemic control in metapopulations. *PLoS one*, (9):e24577.

- [18] Carl D. Meyer. *Matrix Analysis and Applied Linear Algebra*. SIAM, ene 2000.
- [19] Cameron Nowzari, Victor M. Preciado, and George J. Pappas. Analysis and control of epidemics: A survey of spreading processes on complex networks. *IEEE Control Systems Magazine*, 36(1):26–46, 2016.
- [20] Romualdo Pastor-Satorras and Alessandro Vespignani. Epidemic dynamics and endemic states in complex networks. *Physical Review E*, 63(6), May 2001.
- [21] Romualdo Pastor-Satorras and Alessandro Vespignani. Epidemic spreading in scale-free networks. *Physical Review Letters*, 86(14):3200–3203, Apr 2001.
- [22] Romualdo Pastor-Satorras and Alessandro Vespignani. Immunization of complex networks. *Physical Review E*, 65(3), Feb 2002.
- [23] Robert E Rowthorn, Ramanan Laxminarayan, and Christopher A Gilligan. Optimal control of epidemics in metapopulations. *Journal of the Royal Society Interface*, 6(41):1135–1144, 2009.
- [24] A. Seierstad and K. Sydsæter. *Optimal Control Theory With Economic Applications*, volume 1. Elsevier Science, feb 1987.
- [25] Elias M. Stein. *Singular Integrals and Differentiability Properties of Functions*. Princeton University Press, feb 1971.
- [26] Giovanni Strona and Claudio Castellano. Rapid decay in the relative efficiency of quarantine to halt epidemics in networks. *Physical Review E*, 97(2), Feb 2018.
- [27] Gilbert G Walter and Martha Contreras. *Compartmental Modeling With Networks*. Springer Science & Business Media, dic 2012.