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The role of permanently resident populations in the two-patches SIR model with commuters

ISMAIL MIMOUNI AND ALAIN RAPAPORT*

MISTEA, Univ. Montpellier, INRAE, Institut Agro, Montpellier, France E-mails: mimouni.ismail@yahoo.com, alain.rapaport@inrae.fr

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Abstract

We consider a two-patches SIR model where communication occurs thru commuters, distinguishing explicitly permanently resident populations from commuters populations. We give an explicit formula of the reproduction number, and show how the proportions of permanently resident populations impact it. We exhibit non-intuitive situations for which allowing commuting from a safe territory to another one where the transmission rate is higher can reduce the overall epidemic threshold and avoid an outbreak.

Key-words. SIR model, reproduction number, patches models, commuters.

Mathematics Subject Classification. 92D30, 34D20, 90C31.

1 Introduction

Since the pioneer work of Kermack and McKendrick [21], the SIR model has been very popular in epidemiology, as the basic model for infectious diseases with direct transmission (e.g. [1]). It retakes great importance nowadays due to the recent coronavirus pandemic. While early models were not spatialized, the importance of accounting for spatial heterogeneity has been often reported in the literature (see, e.g. [2, 29, 17, 19, 20, 23]). However, different mechanisms come into play to explain the spatial spreading of a disease. Although diffusion appears to be a natural process to describe the local propagation of an infectious agent among a population, which leads to models with partial differential equations [28], it appears to be not well suited for describing long distance spreading. In particular, transportation between cities comes into the picture as a major source of rapid spreading among non-homogeneous populations [5, 3, 30, 25, 27, 8, 36, 31, 24]. Meta-populations or multi-patches models are then more appropriate to describe the spatial characteristics of the propagation [34, 35, 6, 13, 4], as already well considered in ecology [15, 26]. These models require a precise description of the movements between patches, which are most of the time assumed to be linear and thus encoded into a connection matrix [6, 4]. Typically one obtains a system of ordinary differential equations on a graph, which couples the communication dynamics with the epidemiological one.

For diseases spreading among human populations living in different cities, "commuters" (individuals leaving in a city, traveling regularly for short periods in a neighboring city, and coming back to their home city) play a crucial role in the disease propagation among territories [18, 17, 19, 27, 36]. Such coupling between patches have been already considered in the literature, distinguishing among populations N_i attached to a city *i* the sub-population N_{ii} present in its permanent housing from other sub-populations N_{ij} temporary present in another city $j \neq i$ (it can be also seen as multi-groups models as in [9, 14, 16]). However, such models explicitly assume that the whole population housing in a given city can potentially commute to another one. We believe that this is not always fully realistic and that a sub-population that never (or very rarely) moves to another city should be distinguished from the sub-population that visits at a regular basis another city. The study of this extension, which has not been yet considered in the literature to our knowledge, and how it impacts the disease spreading, is the primary objective of the

^{*}corresponding author

present work. For this purpose, we establish an analytical expression of the reproduction number (as the epidemic threshold formerly introduced and analyzed in [10, 32, 11, 12]) for the two patches case (that is also valid for the particular case when the whole populations travel, for which the exact expression of the reproduction number has not been yet provided in the literature).

We also had in mind to consider heterogeneity among territories when disease transmission differs from one city to another one. Typically, non-pharmaceutical interventions (such as reducing physical distance in the population) could be applied with different strength in each city, providing distinct transmission rates. Then, we aim at analyzing how the proportions of commuters in each city can increase or decrease the overall reproduction number. Intuitively, one may believe that the best way to reduce the spreading is to encourage commuters from the city with the lowest transmission rate to do not travel to the other city, and on the opposite to encourage as much as possible commuters from the other city to spend time in the safer city. Indeed, we shall see that this is not always true... The second objective of the present work is thus to study the minimization of the epidemic threshold of the two-patches model with respect to these proportions, depending on the commuting rates. This analysis can potentially serve for decisions making to prevent epidemic outbreak (as in [22] for instance).

The paper is organized as follows. In the next section, we present the complete model in dimension 18 and give some preliminaries. Section 3 is devoted to the analysis of the asymptotic behavior of the solutions of the model. We give and demonstrate an explicit expression of the reproduction number, introducing four relevant quantities q_{ij} (i, j = 1, 2). In a corollary, we also give an alternative way of computation, which is useful in the following. In Section 4, we study the minimization of the reproduction number with respect to the proportions of commuters in each patch. Finally, Section 5 gives a numerical illustration of the results, considering two territories with intrinsic basic reproduction numbers lower and higher than one. We depict the relative sizes of the permanently resident populations that can avoid the outbreak of the epidemic depending on the commuting rates, and discuss the various cases. We end by a conclusion.

2 The model

We follow the modeling of commuters proposed in [18] between two patches (such as cities or territories), but here we consider in addition that a part of the population in each patch do not commute (the "permanently resident" sub-population). We consider populations of size N_i whose home belongs to a patch $i \in \{1, 2\}$, structured in three groups:

- i. permanently resident, being all the time in patch i, whose population size is denoted N_{ir} ,
- ii. commuters to patch j, but located in patch i at time t, of population size denoted N_{ii} ,
- iii. commuters to patch j and located in patch j at time t, of population size denoted N_{ij} .

We shall denote $N_{ic} = N_{ii} + N_{ij}$ the size of the total population of commuters having home in patch *i*. The individuals commutes to patch *j* at a rate λ_i with a return rate μ_i . For each group $g \in \{ir, ii, ij\}$ we denote by S_g , I_g , R_g the sizes of susceptible, infected and recovered sub-populations.

We consider the SIR model assuming that the recovery parameter γ is identical everywhere while the transmission rate β_i depends on the patch *i* but is identical among each group. The model writes as

follows (with $i \neq j$ in $\{1, 2\}$).

$$\begin{split} \dot{S}_{ir} &= -\beta_i S_{ir} \frac{I_{ir} + I_{ii} + I_{ji}}{N_{ir} + N_{ii} + N_{ji}}, \\ \dot{I}_{ir} &= \beta_i S_{ir} \frac{I_{ir} + I_{ii} + I_{ji}}{N_{ir} + N_{ii} + N_{ji}} - \gamma I_{ir}, \\ \dot{R}_{ir} &= \gamma I_{ir}, \\ \dot{S}_{ii} &= -\beta_i S_{ii} \frac{I_{ir} + I_{ii} + I_{ji}}{N_{ir} + N_{ii} + N_{ji}} - \lambda_i S_{ii} + \mu_i S_{ij}, \\ \dot{I}_{ii} &= \beta_i S_{ii} \frac{I_{ir} + I_{ii} + I_{ji}}{N_{ir} + N_{ii} + N_{ji}} - \gamma I_{ii} - \lambda_i I_{ii} + \mu_i I_{ij}, \\ \dot{R}_{ii} &= \gamma I_{ii} - \lambda_i R_{ii} + \mu_i R_{ij}, \\ \dot{S}_{ij} &= -\beta_j S_{ij} \frac{I_{jr} + I_{jj} + I_{ij}}{N_{jr} + N_{jj} + N_{ij}} - \gamma I_{ij} + \lambda_i S_{ii} - \mu_i S_{ij}, \\ \dot{I}_{ij} &= \beta_j S_{ij} \frac{I_{jr} + I_{jj} + I_{ij}}{N_{jr} + N_{jj} + N_{ij}} - \gamma I_{ij} + \lambda_i I_{ii} - \mu_i I_{ij}, \\ \dot{R}_{ij} &= \gamma I_{ij} + \lambda_i R_{ii} - \mu_i R_{ij} \end{split}$$

One can straightforwardly check that the population sizes N_{ir} , and N_{ic} are constant. Moreover N_{ii} , N_{ij} fulfill the system of equations

$$\begin{cases} \dot{N}_{ii} = -\lambda_i N_{ii} + \mu_i N_{ij} \\ \dot{N}_{ij} = \lambda_i N_{ii} - \mu_i N_{ij} \end{cases}$$

whose solutions verify

$$\lim_{t \to +\infty} N_{ii}(t) = \bar{N}_{ii} := \frac{\mu_i}{\lambda_i + \mu_i} N_{ic}, \quad \lim_{t \to +\infty} N_{ij}(t) = \bar{N}_{ij} := \frac{\lambda_i}{\lambda_i + \mu_i} N_{ic}$$
(1)

We shall assume that populations are already balanced at initial time i.e. that one has $N_{ii} = \bar{N}_{ii}$, $N_{ij} = \bar{N}_{ij}$ (constant). For simplicity, we shall drop the notation⁻ in the following, and denote

$$N_{ip} := N_{ir} + N_{ii} + N_{ji}$$

which represents the (constant) size of the total population present in patch i.

3 The epidemic threshold

We denote the vectors

$$I = (I_{1r}, I_{11}, I_{12}, I_{2r}, I_{22}, I_{21})^{\top}, \quad S = (S_{1r}, S_{11}, S_{12}, S_{2r}, S_{22}, S_{21})^{\top}$$

and consider the state vector

$$X = \left[\begin{array}{c} I \\ S \end{array} \right]$$

which belongs to the invariant domain

$$\mathcal{D} := \{ X \in \mathbb{R}^{12}_+; \ \mathbb{M}X \le \mathbb{N} \}$$

where $\mathbb N$ is the vector

$$\mathbb{N} = (N_{1r}, N_{11}, N_{12}, N_{2r}, N_{22}, N_{12})^{\mathsf{T}}$$

and $\mathbb M$ the square matrix

$$\mathbb{M} = [\mathbb{I}_6, \mathbb{I}_6]$$

(where \mathbb{I}_6 denotes the identity matrix of dimension 6×6). The disease free equilibrium is defined as

$$X^{\star} = \left[\begin{array}{c} 0\\ \mathbb{N} \end{array} \right]$$

Let \mathcal{R}_i be the intrinsic reproduction number in the patch *i* (i.e. when there is no connection between patches), that is

$$\mathcal{R}_i := \frac{\beta_i}{\gamma}.$$

We give now an explicit expression of the epidemic threshold when the two patches communicates via commuters.

Proposition 1. Let

$$\mathcal{R}_{1,2} := \frac{q_{11} + q_{22} + \sqrt{(q_{22} - q_{11})^2 + 4q_{12}q_{21}}}{2} \tag{2}$$

where

$$q_{11} = \mathcal{R}_{1} \left(\frac{N_{1r}}{N_{1p}} + \frac{N_{11}}{N_{1p}} \frac{\gamma + \mu_{1}}{\gamma + \lambda_{1} + \mu_{1}} + \frac{N_{21}}{N_{1p}} \frac{\gamma + \lambda_{2}}{\gamma + \lambda_{2} + \mu_{2}} \right)$$

$$q_{22} = \mathcal{R}_{2} \left(\frac{N_{2r}}{N_{2p}} + \frac{N_{22}}{N_{2p}} \frac{\gamma + \mu_{2}}{\gamma + \lambda_{2} + \mu_{2}} + \frac{N_{12}}{N_{2p}} \frac{\gamma + \lambda_{1}}{\gamma + \lambda_{1} + \mu_{1}} \right)$$

$$q_{21} = \mathcal{R}_{1} \left(\frac{N_{11}}{N_{1p}} \frac{\lambda_{1}}{\gamma + \lambda_{1} + \mu_{1}} + \frac{N_{21}}{N_{1p}} \frac{\mu_{2}}{\gamma + \lambda_{2} + \mu_{2}} \right)$$

$$q_{12} = \mathcal{R}_{2} \left(\frac{N_{12}}{N_{2p}} \frac{\mu_{1}}{\gamma + \lambda_{1} + \mu_{1}} + \frac{N_{22}}{N_{2p}} \frac{\lambda_{2}}{\gamma + \lambda_{2} + \mu_{2}} \right)$$
(3)

Then, one has the following properties.

- i. If $\mathcal{R}_{1,2} > 1$, then X^* is unstable.
- ii. If $\mathcal{R}_{1,2} < 1$, then X^* is exponentially stable with respect to the variable¹ I.
- iii. If $\mathcal{R}_1 = \mathcal{R}_2 := \mathcal{R}$, then $\mathcal{R}_{1,2} = \mathcal{R}$.

Proof. Write the dynamics of X as $\dot{X} = f(X)$. The jacobian matrix J of f at X^* is of the form

$$J = \left[\begin{array}{cc} A & 0 \\ \star & B \end{array} \right] \text{ with } A = F - V$$

where

$$F = \begin{bmatrix} \beta_1 \frac{N_{1r}}{N_{1p}} & \beta_1 \frac{N_{1r}}{N_{1p}} & 0 & 0 & 0 & \beta_1 \frac{N_{1r}}{N_{1p}} \\ \beta_1 \frac{N_{11}}{N_{1p}} & \beta_1 \frac{N_{11}}{N_{1p}} & 0 & 0 & 0 & \beta_1 \frac{N_{11}}{N_{1p}} \\ 0 & 0 & \beta_2 \frac{N_{12}}{N_{2p}} & \beta_2 \frac{N_{12}}{N_{2p}} & \beta_2 \frac{N_{12}}{N_{2p}} & 0 \\ 0 & 0 & \beta_2 \frac{N_{2r}}{N_{2p}} & \beta_2 \frac{N_{2r}}{N_{2p}} & \beta_2 \frac{N_{2r}}{N_{2p}} & 0 \\ 0 & 0 & \beta_2 \frac{N_{22}}{N_{2p}} & \beta_2 \frac{N_{22}}{N_{2p}} & \beta_2 \frac{N_{22}}{N_{2p}} & 0 \\ 0 & 0 & \beta_2 \frac{N_{22}}{N_{2p}} & \beta_2 \frac{N_{22}}{N_{2p}} & \beta_2 \frac{N_{22}}{N_{2p}} & 0 \\ \beta_1 \frac{N_{21}}{N_{1p}} & \beta_1 \frac{N_{21}}{N_{1p}} & 0 & 0 & 0 & \beta_1 \frac{N_{21}}{N_{1p}} \end{bmatrix}, V = \begin{bmatrix} \gamma & 0 & 0 & 0 & 0 & 0 \\ 0 & \gamma + \lambda_1 & -\mu_1 & 0 & 0 & 0 \\ 0 & -\lambda_1 & \gamma + \mu_1 & 0 & 0 & 0 \\ 0 & 0 & 0 & \gamma & 0 & 0 \\ 0 & 0 & 0 & \gamma + \lambda_2 & -\mu_2 \\ 0 & 0 & 0 & 0 & -\lambda_2 & \gamma + \mu_2 \end{bmatrix}$$

and

Note that F is a non-negative matrix and V is a non-singular M-matrix. We recall (see for instance from [32]) that one has the property

$$\max \operatorname{Re}(\operatorname{Spec}(A)) < 0 \iff \rho(FV^{-1}) < 1 > 1$$

 $^{^{1}}$ We refer to [33] for the definition of partial stability.

A straightforward computation gives the following expression of the matrix $M := FV^{-1}$

$$M = \begin{bmatrix} \mathcal{R}_{1} \frac{N_{1r}}{N_{1p}} & \mathcal{R}_{1} \frac{N_{1r}(\gamma+\mu_{1})}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{1} \frac{N_{1r}\mu_{1}}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{1} \frac{N_{1r}\lambda_{2}}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{1r}(\gamma+\lambda_{2})}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ \mathcal{R}_{1} \frac{N_{11}}{N_{1p}} & \mathcal{R}_{1} \frac{N_{11}(\gamma+\mu_{1})}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{1} \frac{N_{11}\mu_{1}}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{1} \frac{N_{11}\lambda_{2}}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{11}(\gamma+\lambda_{2})}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ 0 & \mathcal{R}_{2} \frac{N_{12}\lambda_{1}}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}}{N_{2p}} & \mathcal{R}_{2} \frac{N_{12}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{2} \frac{N_{12}\mu_{2}}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ 0 & \mathcal{R}_{2} \frac{N_{2r}\lambda_{1}}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{2r}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{2r}}{N_{2p}} & \mathcal{R}_{2} \frac{N_{2r}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{2} \frac{N_{2r}\mu_{2}}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} \\ 0 & \mathcal{R}_{2} \frac{N_{22}\lambda_{1}}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{22}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{22}}{N_{2p}} & \mathcal{R}_{2} \frac{N_{22}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{2} \frac{N_{22}\mu_{2}}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} \\ \mathcal{R}_{1} \frac{N_{11}}{N_{1p}} & \mathcal{R}_{1} \frac{N_{21}(\gamma+\mu_{1})}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{1} \frac{N_{21}\mu_{1}}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{1} \frac{N_{21}\lambda_{2}}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}(\gamma+\lambda_{2})}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ \end{array} \right$$

Let us consider the diagonal matrix

$$D := \begin{bmatrix} \mathcal{R}_1 \frac{N_{1r}}{N_{1p}} & & \\ & \mathcal{R}_1 \frac{N_{11}}{N_{1p}} & & \\ & & \mathcal{R}_2 \frac{N_{12}}{N_{2p}} & & \\ & & & \mathcal{R}_2 \frac{N_{2r}}{N_{2p}} & & \\ & & & & \mathcal{R}_2 \frac{N_{22}}{N_{2p}} & \\ & & & & \mathcal{R}_1 \frac{N_{21}}{N_{1p}} \end{bmatrix}$$

and the matrix $Q = D^{-1}MD$. A straightforward computation gives the expression

$$Q = \begin{bmatrix} \mathcal{R}_{1} \frac{N_{1r}}{N_{1p}} & \mathcal{R}_{1} \frac{N_{11}(\gamma+\mu_{1})}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}\mu_{1}}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{2} \frac{N_{22}\lambda_{2}}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}(\gamma+\lambda_{2})}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ \mathcal{R}_{1} \frac{N_{1r}}{N_{1p}} & \mathcal{R}_{1} \frac{N_{11}(\gamma+\mu_{1})}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}\mu_{1}}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{2} \frac{N_{22}\lambda_{2}}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}(\gamma+\lambda_{2})}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ 0 & \mathcal{R}_{1} \frac{N_{11}\lambda_{1}}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{2r}}{N_{2p}} & \mathcal{R}_{2} \frac{N_{22}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}\mu_{2}}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ 0 & \mathcal{R}_{1} \frac{N_{11}\lambda_{1}}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{2r}}{N_{2p}} & \mathcal{R}_{2} \frac{N_{22}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}\mu_{2}}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ 0 & \mathcal{R}_{1} \frac{N_{11}\lambda_{1}}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{2r}}{N_{2p}} & \mathcal{R}_{2} \frac{N_{22}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}\mu_{2}}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ \mathcal{R}_{1} \frac{N_{1r}}{N_{1p}} & \mathcal{R}_{1} \frac{N_{11}(\gamma+\mu_{1})}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{2} \frac{N_{22}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}(\gamma+\lambda_{2})}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ \mathcal{R}_{1} \frac{N_{1r}}{N_{1p}} & \mathcal{R}_{1} \frac{N_{11}(\gamma+\mu_{1})}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{2} \frac{N_{22}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}(\gamma+\lambda_{2})}{N_{1p}(\gamma+\lambda_{2}+\mu_{2})} \\ \mathcal{R}_{1} \frac{N_{1r}}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{12}(\gamma+\lambda_{1})}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{2} \frac{N_{22}(\gamma+\mu_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{21}(\gamma+\lambda_{2})}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} \\ \mathcal{R}_{1} \frac{N_{1r}}{N_{1p}(\gamma+\lambda_{1}+\mu_{1})} & \mathcal{R}_{2} \frac{N_{2}}{N_{2p}(\gamma+\lambda_{1}+\mu_{1})} & 0 & \mathcal{R}_{2} \frac{N_{2}}{N_{2p}(\gamma+\lambda_{2}+\mu_{2})} & \mathcal{R}_{1} \frac{N_{1}}{N_{1$$

The matrix Q is non-negative and irreducible. By Perron-Frobenius Theorem (see for instance [7]), this matrix admits a unique positive eigenvector (up to a scalar multiplication) that corresponds to the simple (positive) eigenvalue $\ell = \rho(Q) = \rho(M)$.

Note that the rank of Q is two. We posit

$$Y = (1, 1, 0, 0, 0, 1)^{+}, \quad Z = (0, 0, 1, 1, 1, 0)^{+}$$

and define Q_Y , Q_Z the first and third lines, respectively, of the matrix Q. Then, for any vector $X \in \mathbb{R}^6$, on has one has $QX = (Q_Y X)Y + (Q_Z X)Z$. We look for an positive eigenvector X of the form $X = \alpha Y + (1 - \alpha)Z$ with $\alpha \in (0, 1)$. One has then

$$QX = \alpha QY + (1 - \alpha)QZ = \alpha ((Q_Y Y)Y + (Q_Z Y)Z) + (1 - \alpha) ((Q_Y Z)Y + (Q_Z Z)Z) = (\alpha (Q_Y Y) + (1 - \alpha) (Q_Y Z))Y + (\alpha (Q_Z Y) + (1 - \alpha) (Q_Z Z))Z$$
(4)

On another hand, as X is an eigenvector, one has

$$QX = \ell X = \alpha \ell Y + (1 - \alpha)\ell Z \tag{5}$$

The vectors Y and Z being orthogonal, one obtains from (4)-(5) the conditions

$$\begin{cases} \alpha Q_Y Y + (1-\alpha)Q_Y Z = \alpha \ell \\ \alpha Q_Z Y + (1-\alpha)Q_Z Z = (1-\alpha)\ell \end{cases}$$
(6)

Let $r = \frac{1-\alpha}{\alpha}$. Eliminating ℓ in the two previous equations, r is the positive solution of the polynomial

$$r^2 Q_Y Z + r(Q_Y Y - Q_Z Z) - Q_Z Y = 0$$

and $\ell = Q_Y Y + r Q_Y Z$. One obtains straightforwardly the expression of the eigenvalue

$$\ell = \frac{Q_Y Y + Q_Z Z + \sqrt{(Q_Y Y - Q_Z Z)^2 + 4(Q_Y Z)(Q_Z Y)}}{2}$$

Finally, from the expression of Q, one gets

$$q_{11} = Q_Y Y = \mathcal{R}_1 \left(\frac{N_{1r}}{N_{1p}} + \frac{N_{11}}{N_{1p}} \frac{\gamma + \mu_1}{\gamma + \lambda_1 + \mu_1} + \frac{N_{21}}{N_{1p}} \frac{\gamma + \lambda_2}{\gamma + \lambda_2 + \mu_2} \right)$$

$$q_{22} = Q_Z Z = \mathcal{R}_2 \left(\frac{N_{2r}}{N_{2p}} + \frac{N_{22}}{N_{2p}} \frac{\gamma + \mu_2}{\gamma + \lambda_2 + \mu_2} + \frac{N_{12}}{N_{2p}} \frac{\gamma + \lambda_1}{\gamma + \lambda_1 + \mu_1} \right)$$

$$q_{21} = Q_Z Y = \mathcal{R}_1 \left(\frac{N_{11}}{N_{1p}} \frac{\lambda_1}{\gamma + \lambda_1 + \mu_1} + \frac{N_{21}}{N_{1p}} \frac{\mu_2}{\gamma + \lambda_2 + \mu_2} \right)$$

$$q_{12} = Q_Y Z = \mathcal{R}_2 \left(\frac{N_{12}}{N_{2p}} \frac{\mu_1}{\gamma + \lambda_1 + \mu_1} + \frac{N_{22}}{N_{2p}} \frac{\lambda_2}{\gamma + \lambda_2 + \mu_2} \right)$$

and thus $\ell = \mathcal{R}_{1,2}$, which is exactly $\rho(M)$.

i. When $\mathcal{R}_{1,2} > 1$, the matrix A has at least one eigenvalue with positive real part and the matrix J as well. The equilibrium X^* is thus unstable on \mathcal{D} .

ii. When $\mathcal{R}_{1,2} < 1$, the matrix A is Hurwitz, but X^* is not an hyperbolic equilibrium. However, on can write the dynamics of the vector I as an non-autonomous system

$$\dot{I} = g(t, I) := \begin{pmatrix} \beta_1 S_{1r}(t) \frac{I_{1r} + I_{11} + I_{21}}{N_{1p}} - \gamma I_{1r} \\ \beta_1 S_{11}(t) \frac{I_{1r} + I_{11} + I_{21}}{N_{1p}} - (\gamma + \lambda_1) I_{11} + \mu_1 I_{12} \\ \beta_2 S_{12}(t) \frac{I_{2r} + I_{22} + I_{12}}{N_{2p}} + \lambda_1 I_{11} - (\gamma + \mu_1) I_{12} \\ \beta_2 S_{2r}(t) \frac{I_{2r} + I_{22} + I_{12}}{N_{2p}} - \gamma I_{2r} \\ \beta_2 S_{22}(t) \frac{I_{2r} + I_{22} + I_{12}}{N_{2p}} - (\gamma + \lambda_2) I_{22} + \mu_2 I_{21} \\ \beta_1 S_{21}(t) \frac{I_{1r} + I_{11} + I_{21}}{N_{1p}} + \lambda_2 I_{22} - (\gamma + \mu_2) I_{21} \end{pmatrix}$$

Note that this dynamics is cooperative and as for any $t \ge 0$ one has $S_{ij}(t) \le N_{ij}$ for $ij \in \{1r, 11, 12, 2r, 22, 21\}$, one get

$$g(t,I) \le \bar{g}(I) := AI, \ I \ge 0$$

Therefore, any solution $I(\cdot)$ of $\dot{I} = g(t, I)$ with $I(0) = I_0 \ge 0$ verifies $0 \le I(t) \le \bar{I}(t)$ for any $t \ge 0$, where $\bar{I}(\cdot)$ is solution of the linear dynamics $\dot{\bar{I}} = \bar{g}(\bar{I})$ with $\bar{I}(0) = I_0$. As A is Hurwitz, we conclude that X^* is exponentially stable with respect to I, which proves point ii.

iii. For the particular case $\mathcal{R}_1 = \mathcal{R}_2 := \mathcal{R}$, the transpose of the matrix M writes

$$M^{\top} = \mathcal{R} \begin{bmatrix} \frac{N_{1r}}{N_{1p}} & \frac{N_{11}}{N_{1p}} & 0 & 0 & 0 & \frac{N_{21}}{N_{1p}} \\ \frac{N_{1r}(\gamma+\mu_1)}{N_{1p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{11}(\gamma+\mu_1)}{N_{1p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{12}\lambda_1}{N_{2p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{2r}\lambda_1}{N_{2p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{22}\lambda_1}{N_{2p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{21}(\gamma+\mu_1)}{N_{1p}(\gamma+\lambda_1+\mu_1)} \\ \frac{N_{1r}\mu_1}{N_{1p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{11}\mu_1}{N_{1p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{12}(\gamma+\lambda_1)}{N_{2p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{2r}(\gamma+\lambda_1)}{N_{2p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{22}(\gamma+\lambda_1)}{N_{2p}(\gamma+\lambda_1+\mu_1)} & \frac{N_{21}\mu_1}{N_{1p}(\gamma+\lambda_1+\mu_1)} \\ 0 & 0 & \frac{N_{12}}{N_{2p}} & \frac{N_{2r}}{N_{2p}} & \frac{N_{22}}{N_{2p}} & 0 \\ \frac{N_{1r}\lambda_2}{N_{1p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{11}\lambda_2}{N_{1p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{12}(\gamma+\mu_2)}{N_{2p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{22}(\gamma+\mu_2)}{N_{2p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{21}\lambda_2}{N_{1p}(\gamma+\lambda_2+\mu_2)} \\ \frac{N_{1r}(\gamma+\lambda_2)}{N_{1p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{11}(\gamma+\lambda_2)}{N_{1p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{12}\mu_2}{N_{1p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{22}\mu_2}{N_{1p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{21}(\gamma+\lambda_2)}{N_{1p}(\gamma+\lambda_2+\mu_2)} & \frac{N_{21}(\gamma+\lambda_2)}{N_{21}(\gamma+\lambda_2+\mu_2)} & \frac{N_{21}(\gamma+\lambda_2)}{N_{21}(\gamma+\lambda_2$$

One can check that one has $M^{\top}U = \mathcal{R}U$ where $U = (1, 1, 1, 1, 1, 1)^{\top}$. As U is a positive vector, we deduce from the Perron-Frobenius Theorem that one has $\rho(M) = \rho(M^T) = \mathcal{R}$, which ends the proof. \Box

Remark 1. The explicit expression (2) of the epidemic threshold given in Proposition 1 is also relevant in absence of permanently resident populations, which has not been yet provided explicitly in the literature (up to our knowledge).

Corollary 1. One has

$$\min\left(\mathcal{R}_1, \mathcal{R}_2\right) \le \mathcal{R}_{1,2} \le \max\left(\mathcal{R}_1, \mathcal{R}_2\right)$$

Proof. Denote by $M(\mathcal{R}_1, \mathcal{R}_2)$ the matrix FV^{-1} for the parameters $\mathcal{R}_1, \mathcal{R}_2$, and let $\mathcal{R}_- := \min(\mathcal{R}_1, \mathcal{R}_2)$, $\mathcal{R}_+ := \max(\mathcal{R}_1, \mathcal{R}_2)$. From the expression of the non-negative matrices M, one gets

$$M(\mathcal{R}_{-},\mathcal{R}_{-}) \leq M(\mathcal{R}_{1},\mathcal{R}_{2}) \leq M(\mathcal{R}_{+},\mathcal{R}_{+})$$

which implies (see for instance [7]) the inequalities

$$\rho(M(\mathcal{R}_{-},\mathcal{R}_{-})) \leq \rho(M(\mathcal{R}_{1},\mathcal{R}_{2})) \leq \rho(M(\mathcal{R}_{+},\mathcal{R}_{+}))$$

and thus

$$\mathcal{R}_{-} \leq \mathcal{R}_{1,2} \leq \mathcal{R}_{+}.$$

Alternatively, the number $\mathcal{R}_{1,2}$ can be determined as follows.

Corollary 2. Assume $\mathcal{R}_2 > \mathcal{R}_1$. Then, one has

$$\mathcal{R}_{1,2} = \alpha \mathcal{R}_1 + (1-\alpha) \mathcal{R}_2 \tag{7}$$

where $\alpha \in [0,1)$ is the smallest root of the polynomial

$$P(\alpha) = \alpha^{2}(\mathcal{R}_{2} - \mathcal{R}_{1}) - \alpha(\mathcal{R}_{2} - \mathcal{R}_{1} + q_{12} + q_{21}) + q_{12}$$

Proof. One can check, from expressions (3), that one has $q_{11} + q_{21} = \mathcal{R}_1$ and $q_{22} + q_{12} = \mathcal{R}_2$. Then, from (6), one get

$$\mathcal{R}_{1,2} = l = \alpha \mathcal{R}_1 + (1 - \alpha) \mathcal{R}_2 \tag{8}$$

where α is a root of the polynomial P obtained from (6) by eliminating l, that is

$$P(\alpha) = \alpha^{2}(\mathcal{R}_{2} - \mathcal{R}_{1}) - \alpha(\mathcal{R}_{2} - \mathcal{R}_{1} + q_{12} + q_{21}) + q_{12}$$

From Corollary 1, we know that α belongs to [0, 1]. Note that one has $P(0) = q_{12} \ge 0$ and $P(1) = -q_{21} \le 0$. Therefore, when $\mathcal{R}_2 - \mathcal{R}_1 > 0$, P admits exactly one root in [0, 1) and another one in $[1, \rightarrow)$. However, if $\alpha = 1$ one should have $q_{21} = 0$ and thus $\lambda_1 = 0$, $\mu_2 = 0$, which implies $N_{11} = N_{1c}$, $N_{12} = 0$, $N_{22} = 0$, $N_{21} = N_{2c}$. Then, one obtains $q_{11} = \mathcal{R}_1$, $q_{22} = \mathcal{R}_2$ and from the expression (2) on gets $\mathcal{R}_{1,2} = \max(\mathcal{R}_1, \mathcal{R}_2) = \mathcal{R}_2$ which contradicts $\alpha = 1$. We conclude that α belongs to [0, 1) and is thus the smallest root of P.

Remark 2. When there is no communication between patches (that is $N_{1r} = N_{1p} = N_1$, $N_{2r} = N_{2p} = N_2$), one has $q_{21} = 0$ and $q_{12} = 0$. If $\mathcal{R}_2 > \mathcal{R}_1$, resp. $\mathcal{R}_1 > \mathcal{R}_2$, one has $\alpha = 0$, resp. $\alpha = 1$, which gives

$$\mathcal{R}_{1,2} = \max(\mathcal{R}_1, \mathcal{R}_2).$$

We look now for a characterization of the minimum value of the threshold $\mathcal{R}_{1,2}$.

4 Minimization of the epidemic threshold

In this section, we assume that the mixing is fast compared to the recovery rate (as its is often considered in the literature), which amounts to have numbers λ_i , μ_i large compared to γ . Our objective is to study how the proportions of commuters in the populations impact the value of $\mathcal{R}_{1,2}$.

Given \mathcal{R}_1 , \mathcal{R}_2 , we consider the approximation $\mathcal{R}_{1,2}$ of the threshold $\mathcal{R}_{1,2}$ which consists in keeping $\gamma = 0$ in the expressions (3). For convenience, we posit the numbers

$$\eta_i := \frac{\lambda_i}{\lambda_i + \mu_i} \in (0, 1) \qquad (i = 1, 2)$$

One has a first result about the variations of $\mathcal{R}_{1,2}$ with respect to N_{1c} , N_{2c} .

Proposition 2. Fix parameters N_i , β_i , γ , λ_i , μ_i (i = 1, 2) such that $\mathcal{R}_2 > \mathcal{R}_1$.

- i. For any $N_{1c} \in (0, N_1)$, the map $N_{2c} \mapsto \tilde{\mathcal{R}}_{1,2}(N_{1c}, N_{2c})$ is decreasing.
- ii. The map $N_{1c} \mapsto \tilde{\mathcal{R}}_{1,2}(N_{1c}, N_{2c})$ is increasing at (N_{1c}, N_{2c}) when

$$\eta_2(1-\eta_2)N_{2c} > (1-\eta_1)(N_2-\eta_2N_{2c}) \tag{9}$$

iii. The map $N_{1c} \mapsto \tilde{\mathcal{R}}_{1,2}(N_{1c}, N_{2c})$ is increasing, resp. decreasing, at (N_{1c}, N_{2c}) if the numbers A and B are negative, resp. positive, where

$$\begin{split} A &:= \mathcal{R}_2 \frac{\frac{N_2}{2} - \eta_1 (\frac{1}{2} - \eta_1) N_{1c} - (\frac{3}{2} - \eta_2) \eta_2 N_{2c}}{N_2 - \eta_2 N_{2c} + \eta_1 N_{1c}} - \mathcal{R}_1 \frac{\frac{N_1}{2} - (\frac{3}{2} - \eta_1) \eta_1 N_{1c} - \eta_2 (\frac{1}{2} - \eta_2) N_{2c}}{N_1 - \eta_1 N_{1c} + \eta_2 N_{2c}}, \\ B &:= \mathcal{R}_2 \frac{(1 - \eta_1) (N_2 - \eta_2 N_{2c}) - \eta_2 (1 - \eta_2) N_{2c}}{(N_2 - \eta_2 N_{2c} + \eta_1 N_{1c})^2} - \mathcal{R}_1 \frac{(1 - \eta_1) (N_1 + \eta_2 N_{2c}) + \eta_2 (1 - \eta_2) N_{2c}}{(N_1 - \eta_1 N_{1c} + \eta_2 N_{2c})^2} \end{split}$$

Proof. Following Corollary 2, one has

$$\tilde{\mathcal{R}}_{1,2} = \tilde{\alpha}\mathcal{R}_1 + (1 - \tilde{\alpha})\mathcal{R}_2 \tag{10}$$

where $\tilde{\alpha}$ is the smallest root of the polynomial

$$\tilde{P}(\alpha) = \alpha^2 (\mathcal{R}_2 - \mathcal{R}_1) - \alpha (\mathcal{R}_2 - \mathcal{R}_1 + \tilde{q}_{12} + \tilde{q}_{21}) + \tilde{q}_{12}$$

where \tilde{q}_{12} , \tilde{q}_{21} are the approximations of q_{12} , q_{21} defined in (3). Let us note that one can write $N_{ii} = (1 - \eta_i)N_{ic}$, $N_{ij} = \eta_i N_{ic}$ (for $j \neq i$) and also $N_{ip} = N_i - \eta_i N_{ic} + \eta_j N_{jc}$, which leads to the following expressions of \tilde{q}_{12} , \tilde{q}_{21}

$$\tilde{q}_{21} = \mathcal{R}_1 \frac{(1 - \eta_1)\eta_1 N_{1c} + \eta_2 (1 - \eta_2) N_{2c}}{N_1 - \eta_1 N_{1c} + \eta_2 N_{2c}}, \quad \tilde{q}_{12} = \mathcal{R}_2 \frac{\eta_1 (1 - \eta_1) N_{1c} + (1 - \eta_2) \eta_2 N_{2c}}{N_2 - \eta_2 N_{2c} + \eta_1 N_{1c}}$$
(11)

For simplicity, we shall drop the notation $\tilde{}$ in the rest of the proof. Note than α being the smallest root of P, it verifies

$$\alpha < \frac{\mathcal{R}_2 - \mathcal{R}_1 + q_{12} + q_{21}}{2(\mathcal{R}_2 - \mathcal{R}_1)} \tag{12}$$

Let us differentiate the equality $P(\alpha) = 0$ with respect to q_{12} and q_{21} :

$$2\alpha \frac{\partial \alpha}{\partial q_{12}} (\mathcal{R}_2 - \mathcal{R}_1) - \frac{\partial \alpha}{\partial q_{12}} (\mathcal{R}_2 - \mathcal{R}_1 + q_{12} + q_{21}) - \alpha + 1 = 0$$
$$2\alpha \frac{\partial \alpha}{\partial q_{21}} (\mathcal{R}_2 - \mathcal{R}_1) - \frac{\partial \alpha}{\partial q_{21}} (\mathcal{R}_2 - \mathcal{R}_1 + q_{12} + q_{21}) - \alpha = 0$$

which gives

$$\frac{\partial \alpha}{\partial q_{12}} = \frac{1-\alpha}{\mathcal{R}_2 - \mathcal{R}_1 + q_{12} + q_{21} - 2\alpha(\mathcal{R}_2 - \mathcal{R}_1)}$$
$$\frac{\partial \alpha}{\partial q_{21}} = \frac{-\alpha}{\mathcal{R}_2 - \mathcal{R}_1 + q_{12} + q_{21} - 2\alpha(\mathcal{R}_2 - \mathcal{R}_1)}$$

Then, one can write

$$\frac{\partial \alpha}{\partial N_{ic}} = \frac{\partial \alpha}{\partial q_{12}} \frac{\partial q_{12}}{\partial N_{ic}} + \frac{\partial \alpha}{\partial q_{21}} \frac{\partial q_{21}}{\partial N_{ic}} = \frac{(1-\alpha)\frac{\partial q_{12}}{\partial N_{ic}} - \alpha \frac{\partial q_{21}}{\partial N_{ic}}}{\mathcal{R}_2 - \mathcal{R}_1 + q_{12} + q_{21} - 2\alpha(\mathcal{R}_2 - \mathcal{R}_1)} \qquad (i = 1, 2)$$

and from inequality (12), we obtain that the signs of the derivatives $\frac{\partial \alpha}{\partial N_{ic}}$ are given by the sign of the numbers

$$\sigma_i := (1 - \alpha) \frac{\partial q_{12}}{\partial N_{ic}} - \alpha \frac{\partial q_{21}}{\partial N_{ic}} \qquad (i = 1, 2)$$
(13)

We begin by the dependency with respect to N_{2c} . One has first

$$\frac{\partial q_{12}}{\partial N_{2c}} = \mathcal{R}_2 \eta_2 \frac{(1-\eta_2)(N_2+\eta_1 N_{1c}) + \eta_1 (1-\eta_1) N_{1c}}{(N_2+\eta_1 N_{1c} - \eta_2 N_{2c})^2} > 0$$

Note that one has

$$q_{21} = \frac{\mathcal{R}_1(N_2 + \eta_1 N_{1c} - \eta_2 N_{2c})}{\mathcal{R}_2(N_1 - \eta_1 N_{1c} + \eta_2 N_{2c})} q_{12} \tag{14}$$

and thus

$$\frac{\partial q_{21}}{\partial N_{2c}} = \frac{\mathcal{R}_1(N_2 + \eta_1 N_{1c} - \eta_2 N_{2c})}{\mathcal{R}_2(N_1 - \eta_1 N_{1c} + \eta_2 N_{2c})} \frac{\partial q_{12}}{\partial N_{2c}} - \frac{\mathcal{R}_1 \eta_2(N_1 + N_2)}{\mathcal{R}_2(N_1 - \eta_1 N_{1c} + \eta_2 N_{2c})^2} q_{12}$$

Then, one gets the inequality

$$\sigma_2 > \left(1 - \alpha - \alpha \frac{\mathcal{R}_1(N_2 + \eta_1 N_{1c} - \eta_2 N_{2c})}{\mathcal{R}_2(N_1 - \eta_1 N_{1c} + \eta_2 N_{2c})}\right) \frac{\partial q_{12}}{\partial N_{2c}}$$

On another hand, one gets from $P(\alpha) = 0$ the inequality

$$(1-\alpha)q_{12} - \alpha q_{21} = \alpha(1-\alpha)(\mathcal{R}_2 - \mathcal{R}_1) > 0$$

and with (14)

$$(1-\alpha)q_{12} - \alpha q_{21} = \left(1 - \alpha - \alpha \frac{\mathcal{R}_1(N_2 + \eta_1 N_{1c} - \eta_2 N_{2c})}{\mathcal{R}_2(N_1 - \eta_1 N_{1c} + \eta_2 N_{2c})}\right)q_{12} > 0$$

We then conclude that σ_2 is positive, and from (10) we deduce that the map $N_{2c} \mapsto \mathcal{R}_{1,2}$ is decreasing. This proves the point i.

We study now the dependency with respect to N_{1c} . A straightforward calculation gives

$$\frac{\partial q_{12}}{\partial N_{1c}} = \mathcal{R}_2 \eta_1 \frac{(1-\eta_1)(N_2 - \eta_2 N_{2c}) - \eta_2 (1-\eta_2) N_{2c}}{(N_2 - \eta_2 N_{2c} + \eta_1 N_{1c})^2}$$
(15)

and

$$\frac{\partial q_{21}}{\partial N_{1c}} = \mathcal{R}_1 \eta_1 \frac{(1-\eta_1)(N_1+\eta_2 N_{2c}) + \eta_2 (1-\eta_2) N_{2c}}{(N_1-\eta_1 N_{1c}+\eta_2 N_{2c})^2} > 0$$
(16)

When $\frac{\partial q_{12}}{\partial N_{1c}} < 0$, we can conclude that σ_1 is negative and $\mathcal{R}_{1,2}$ is thus increasing with respect to N_{1c} . This condition is equivalent to (9). This proves the point ii. When this last condition is not satisfied, having $\frac{\partial q_{12}}{\partial N_{1c}} < \frac{\partial q_{21}}{\partial N_{1c}}$ with $\alpha > \frac{1}{2}$ is another sufficient condition to obtain $\sigma_1 < 0$ from expression (13). However, having $\alpha > \frac{1}{2}$ amounts to have $P(\frac{1}{2}) > 0$, that is

$$\frac{\mathcal{R}_2 - \mathcal{R}_1}{4} - \frac{\mathcal{R}_2 - \mathcal{R}_1 + q_{12} + q_{21}}{2} + q_{12} > 0$$

or equivalently

$$\frac{\mathcal{R}_2}{2} - q_{12} < \frac{\mathcal{R}_1}{2} - q_{21}$$

One can straightforwardly check that this last condition is equivalent to A < 0 and that the condition $\frac{\partial q_{12}}{\partial N_{1c}} < \frac{\partial q_{21}}{\partial N_{1c}}$ is equivalent to B < 0. In the same manner, having A > 0 and B > 0 implies $\alpha < \frac{1}{2}$ and $\frac{\partial q_{12}}{\partial N_{1c}} > \frac{\partial q_{21}}{\partial N_{1c}}$, which is a sufficient condition to have $\sigma_1 > 0$, and thus $\mathcal{R}_{1,2}$ increasing with respect to N_{1c} . This proves the point iii.

This result suggests that the map $N_{1c} \mapsto \tilde{\mathcal{R}}_{1,2}(N_{1c}, N_{2c})$ is not necessarily monotonic, differently to the map $N_{2c} \mapsto \tilde{\mathcal{R}}_{1,2}(N_{1c}, N_{2c})$. We show now that the possibilities of its variations are limited.

Proposition 3. Under hypotheses of Proposition 2, for each $N_{2c} \in (0, N_2)$ the map $N_{1c} \mapsto \tilde{\mathcal{R}}_{1,2}(N_{1c}, N_{2c})$ possesses one of the three properties

- a. it is decreasing on $(0, N_1)$,
- b. it is increasing on $(0, N_1)$,
- c. there exists $N_{1c}^{\star} \in (0, N_1)$ such that it is decreasing on $(0, N_{1c}^{\star})$ and increasing on (N_{1c}^{\star}, N_1) .

Proof. Fix $N_{2c} \in (0, N_2)$. If the map $N_{1c} \mapsto \tilde{\mathcal{R}}_{1,2}(N_{1c}, N_{2c})$ is not monotonic, there exists $\hat{N}_{1c} \in (0, N_1)$ such that $\frac{\partial \tilde{\mathcal{R}}_{1,2}}{\partial N_{1c}}(\hat{N}_{1c}, N_{2c}) = 0$. For simplicity, we shall drop the notation $\tilde{}$ in the rest of the proof. Following the proof of Proposition 2, one has $\mathcal{R}_{1,2} = \alpha \mathcal{R}_1 + (1-\alpha)\mathcal{R}_2$ with

$$\frac{\partial \alpha}{\partial N_{1c}} = \frac{(1-\alpha)\frac{\partial q_{12}}{\partial N_{1c}} - \alpha \frac{\partial q_{21}}{\partial N_{1c}}}{\mathcal{R}_2 - \mathcal{R}_1 + q_{12} + q_{21} - 2\alpha(\mathcal{R}_2 - \mathcal{R}_1)} := \frac{\sigma_1}{\nu}$$

where $\nu > 0$. Therefore, one has $\frac{\partial \alpha}{\partial N_{1c}} = 0$ and $\sigma_1 = 0$ at $N_{1c} = \hat{N}_{1c}$, and thus

$$\left. \frac{\partial^2 \alpha}{\partial N_{1c}^2} \right|_{N_{1c} = \hat{N}_{1c}} = \left. \frac{\frac{\partial \sigma_1}{\partial N_{1c}}}{\nu} \right|_{N_{1c} = \hat{N}_{1c}} = \left. \frac{(1-\alpha)\frac{\partial^2 q_{12}}{\partial N_{1c}^2} - \alpha \frac{\partial^2 q_{21}}{\partial N_{1c}^2}}{\nu} \right|_{N_{1c} = \hat{N}_{1c}}$$

From expressions (15) and (16), a straightforward calculation gives

$$\frac{\partial^2 q_{12}}{\partial N_{1c}^2} = \frac{-2\eta_1 \frac{\partial q_{12}}{\partial N_{1c}}}{N_1 - \eta_1 N_{1c} + \eta_2 N_{2c}}, \quad \frac{\partial^2 q_{21}}{\partial N_{1c}^2} = \frac{2\eta_1 \frac{\partial q_{21}}{\partial N_{1c}}}{N_2 - \eta_2 N_{2c} + \eta_1 N_{1c}}$$

where $\frac{\partial q_{21}}{\partial N_{1c}} > 0$ and from $\sigma_1 = 0$ one gets $\frac{\partial q_{12}}{\partial N_{1c}} > 0$ for $N_{1c} = \hat{N}_{1c}$. Finally, one obtains

$$\frac{\partial^2 \mathcal{R}_{1,2}}{\partial N_{1c}^2}(\hat{N}_{1c}, N_{2c}) = -(\mathcal{R}_2 - \mathcal{R}_1) \frac{\partial^2 \alpha}{\partial N_{1c}^2}(\hat{N}_{1c}, N_{2c}) < 0$$

Consequently, any extremum of the map $N_{1c} \mapsto \mathcal{R}_{1,2}(N_{1c}, N_{2c})$ is a local minimizer, which implies that this map has at most one local minimizer.

Finally, we give conditions for which the minimization of the threshold $\mathcal{R}_{1,2}$ presents a trichotomy.

Proposition 4. Let parameters β_i , γ be such that $\mathcal{R}_2 > \mathcal{R}_1$ and assume that N_1 , N_2 satisfy $N_1\mathcal{R}_2 > N_2\mathcal{R}_1$. Then, provided that γ is small enough compared to λ_i and μ_i , the function $(N_{1c}, N_{2c}) \mapsto \mathcal{R}_{1,2}(N_{1c}, N_{2c})$ admits an unique minimum at (N_{1c}^*, N_{2c}^*) with $N_{2c}^* = N_2$. Moreover, one has the following properties.

- 1. $N_{1c}^{\star} = 0$ if $\eta_2 > 1 \eta_1$,
- 2. $N_{1c}^{\star} = N_1$ if η_1 and η_2 are sufficiently small,
- 3. there exists η_1, η_2 for which $N_{1c}^{\star} \in (0, N_1)$.

Proof. We first show that the announced properties are satisfied for the approximate function $\tilde{\mathcal{R}}_{1,2}$.

From Propositions 2 and 3, we know that $\tilde{\mathcal{R}}_{1,2}$ admits an unique minimum at $(\hat{N}_{1c}, \hat{N}_{2c})$ with $\hat{N}_{2c} = N_2$. For $N_{2c} = N_2$, the condition (9) simply writes $\eta_2 > 1 - \eta_1$ which implies from point ii. of Proposition 2 that one has $\hat{N}_{1c} = 0$ when this condition is fulfilled. This shows that point 1 is verified for the function $\tilde{\mathcal{R}}_{1,2}$.

One obtains the limits

$$\lim_{\eta_1,\eta_2 \to 0} A = \frac{\mathcal{R}_2}{2} - \frac{\mathcal{R}_1}{2} > 0, \quad \lim_{\eta_1,\eta_2 \to 0} B = \frac{\mathcal{R}_2}{N_2} - \frac{\mathcal{R}_1}{N_1} > 0$$

which show that numbers A and B are positive when η_1 , η_2 are small, and thus one has $\hat{N}_{1c} = N_1$ from point iii of Proposition 2. This shows that point 2 is verified for the function $\tilde{\mathcal{R}}_{1,2}$.

Take now any $N_{1c} \in (0, N_1)$. When $\eta_2 > 1 - \eta_1$, one has $\frac{\partial \tilde{\mathcal{R}}_{1,2}}{\partial N_{1c}}(N_{1c}, N_2) > 0$, and for η_1, η_2 small, $\frac{\partial \tilde{\mathcal{R}}_{1,2}}{\partial N_{1c}}(N_{1c}, N_2) < 0$ is verified. Then, by continuity of the function $\tilde{\mathcal{R}}_{1,2}$ with respect to parameters η_1 , η_2 , one deduce that the existence of values $\hat{\eta}_1, \hat{\eta}_2$ for which $\frac{\partial \tilde{\mathcal{R}}_{1,2}}{\partial N_1}(N_{1c}, N_2) = 0$. As the function $\tilde{\mathcal{R}}_{1,2}$ cannot have more than a local extremum (see Proposition 3), we deduce that N_{1c} realizes the minimum of the function $N_{1c} \mapsto \tilde{\mathcal{R}}_{1,2}(N_{1c}, N_2)$ when $\eta_1 = \hat{\eta}_1$ and $\eta_2 = \hat{\eta}_2$. This shows that point 3 is verified for the function $\tilde{\mathcal{R}}_{1,2}$.

Finally, note that the exact threshold $\mathcal{R}_{1,2}$ amounts to replace in the expression of \tilde{q}_{12} , \tilde{q}_{21} the numbers η_i by $\frac{\lambda_i + \gamma}{\lambda_i + \mu_i + \gamma}$, which is continuous with respect to γ and equal to η_i for $\gamma = 0$. By continuity of $\tilde{\mathcal{R}}_{1,2}$ with respect to \tilde{q}_{12} , \tilde{q}_{21} , we deduce that uniqueness of the minimizer of $\mathcal{R}_{1,2}$ and properties 1. to 3. are also fulfilled by the function $(N_{1c}, N_{2c}) \mapsto \mathcal{R}_{1,2}$, provided that γ is small enough.

5 Numerical illustration

We consider two territories of same population size $N = N_1 = N_2$ with different transmission rates such that one has $\mathcal{R}_1 < 1 < \mathcal{R}_2$ (values are given in Table 1). Typically, some precautionary measures (such as social distance) are taken in the first territory so that the disease cannot spread in this territory if it is closed, while the epidemic can spread in the second territory in absence of communication with territory 1. We aim at studying how the epidemic can die out when commuting occur between territories, depending on the proportions of resident in each population, denoted

$$p_i = := \frac{N_{ir}}{N} = 1 - \frac{N_{ic}}{N}, \quad (i = 1, 2)$$

(in other words, how to obtain $\mathcal{R}_{1,2} < 1$ playing with p_1, p_2). Note that when $N_1 = N_2$, the threshold $\mathcal{R}_{1,2}$ depends on the proportions p_1, p_2 independently of N.

γ	β_1	β_2	\mathcal{R}_1	\mathcal{R}_2
0.3	0.24	0.33	0.9	1.1

Table 1: Characteristics numbers of the epidemic

Conditions of Proposition 4 are satisfied provided that commuting parameters λ_i , μ_i are large enough. We have considered three sets of these parameters, given in Table 2, that correspond to the three possible situations depicted in Proposition 4.

case	λ_1	μ_1	λ_2	μ_2	η_1	η_2
Α	10	10	10	1	0.5	0.9090909
В	10	100	10	100	0.009901	0.009901
С	10	10	10	70	0.5	0.125

Table 2: Three sets of commuting parameters

The approximate expression $\mathcal{R}_{1,2}$ turns out to be a very good approximation of the exact value $\mathcal{R}_{1,2}$, even in case A for which γ is not so small compared to μ_2 (see Table 3).

case	A	В	С	
$\max_{p_1,p_2} \tilde{\mathcal{R}}_{1,2} - \mathcal{R}_{1,2} $	1.910^{-3}	1.410^{-4}	610^{-4}	

Table 3: Quality of the approximation $\tilde{\mathcal{R}}_{1,2}$

Figures 1, 2, 3 show families of curves $p_1 \mapsto \mathcal{R}_{1,2}$ for different values of $p_2 \in [0,1]$. One can observe that theses curves possess the properties given by Propositions 2 and 3:

- they are either decreasing, increasing or decreasing down to a minimum and then increasing,
- they are ordered and the lower one is obtained for $p_2 = 0$ (i.e. $N_{2c} = N_2$).

This last feature is suite intuitive: the more there are commuters from territory 2 (that spend time in territory 1 where the conditions of transmission disease is lower), the less the epidemic spreads. A way to reduce the value of $\mathcal{R}_{1,2}$ is thus to encourage commuting towards territory 1 (whatever are the commuting rates). However, the role of the resident population in territory 1 is far less intuitive because it does depends on the commuting rates.

- 1. In case A, commuters from territory 2 return more rarely to home than commuters from territory 1 do. The condition of point 1. of Proposition 4 is fulfilled. Then, the threshold $\mathcal{R}_{1,2}$ can be made small (and below 1) when the proportion of resident in territory 1 is high i.e. when the inhabitants of territory 1 are encouraged not to commute.
- 2. In case B, both commuters return rapidly to their home. This means that the numbers of commuters from one territory present in the other one at a given time is low. Then the condition of point 2 of Proposition 4 is fulfilled. Here, it is better to encourage inhabitants of territory 1 to commute

to the other territory where the disease spreads yet more easily... which is counter-intuitive at first sight. Indeed, commuters do not send much time in the other territory, and have thus heuristically less time to meet and transmit the disease...

3. In case C, commuters from territory 2 return more rapidly to home than commuters from territory 1 do, on the opposite of case A. Conditions of points 1 and 2 of Proposition 4 are not fulfilled here and we are in an intermediate situation for which point 3 of Proposition 4 occurs. It is theoretically possible to have $\mathcal{R}_{1,2} < 1$ on the condition that the proportion of commuters of territory 1 is well balanced.

Finally, this example shows that playing only with the return rates μ_1 , μ_2 provides the three possible scenarios, but other changes could also exhibit them.

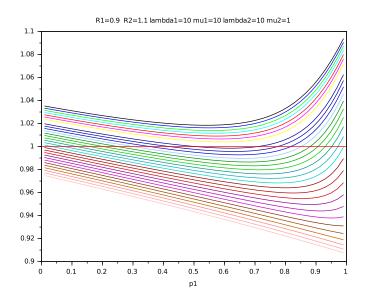


Figure 1: $\mathcal{R}_{1,2}$ as a function of p_1 in case A (each curve corresponds to a value of $p_2 \in [0,1]$)

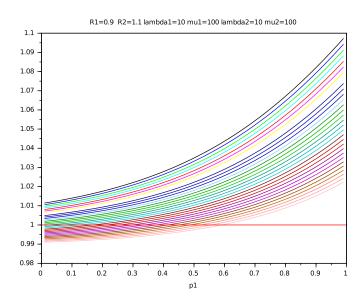


Figure 2: $\mathcal{R}_{1,2}$ as a function of p_1 in case B (each curve corresponds to a value of $p_2 \in [0,1]$)

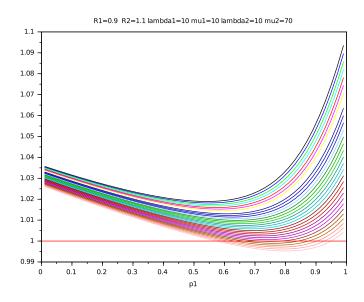


Figure 3: $\mathcal{R}_{1,2}$ as a function of p_1 in case C (each curve corresponds to a value of $p_2 \in [0,1]$)

6 Conclusion

In this work, we have been able to provide an explicit expression of the reproduction number, although the model is in dimension 18. This expression has allowed us to study its minimization with respect to the proportions of permanently resident populations in each patch. We discovered a trichotomy of cases, with some counter intuitive situations. In each case, it is always beneficial to have commuters traveling to a safer city where the transmission rate is lower. However, for the safer city, three situations occurs: either it is better to avoid commuting to the other city, or on the opposite encouraging commuting to the more risky city reduces the reproduction number, and in a third case there exists an optimal intermediate proportion of commuters of the safer city which minimizes the epidemic threshold. In some sense, the permanently resident populations, which have been ignored in former modeling, can play an hidden role in an epidemic outbreak. This is illustrated on an example for which only right proportions of commuters (or permanently resident) avoid the outbreak. This suggests that counter-intuitive situations may also occur when considering networks with more than two nodes. The extension of the present results to more general networks is an open problem, that might be the matter of a future work.

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References

- [1] ANDERSON, R.M. AND MAY, R.M. Infectious Diseases of Humans, Oxford University, 1991.
- [2] ANGULO, J.J, TAKIGUTI, C.K, PEDERNEIRAS, C.A, CARVALHO-DE-SOUZA, A.M, OLIVEIRA-DE-SOUZA, M.C. AND MEGALE, P. Identification of pattern and process in the spread of a contagious disease. Social Science & Medicine, 13D(3):183–9, 1979.
- [3] ARINO, J., JORDAN, R. AND VAN DEN DRIESSCHE, P. Quarantine in a multi-species epidemic model with spatial dynamics. Mathematical Biosciences, 206(1), 46–60, 2007.
- [4] ARINO, J. Diseases in Metapopulations. In Modeling and Dynamics of Infectious Diseases, World Scientific, 64–122, 2009.

- [5] ARINO, J AND VAN DEN DRIESSCHE, P. A multi-city epidemic model. Mathematical Population Studies, 10(3), 175–193, 2003.
- [6] ARINO, J AND VAN DEN DRIESSCHE, P. Disease spread in metapopulations. Fields Institute Communications, 48, 2006.
- [7] BERMAN, A. AND PLEMMONS, R.J. Nonnegative Matrices in the Mathematical Sciences, SIAM, 1994.
- [8] CHEN, Y., YAN, M., AND XIANG, Z., Transmission Dynamics of a Two-City SIR Epidemic Model with Transport-Related Infections. Journal of Applied Mathematics, ID 764278, 2014.
- CLANCY, D. Carrier-borne epidemic models incorporating population mobility. Mathematical Biosciences, 132(2), 185–204, 1996.
- [10] DIEKMANN, O., HEESTERBEEK, J.A.P. AND METZ, J.A.J. On the definition and computation of the basic reproduction ratio R0 in models for infectious diseases in heterogeneous populations. Journal of Mathematical Biology, 28, 365–382 1990.
- [11] DIEKMANN, O., HEESTERBEEK, J.A.P. AND ROBERT M.G., The construction of next-generation matrices for compartmental epidemic models. Journal of the Royal Society Interface, 7, 873–885, 2007.
- [12] DHIRASAKDANON, T., THIEME, H.R. AND VAN DEN DRIESSCHE, P. A sharp threshold for disease persistence in host metapopulations. Journal of Biological Dynamics, 1(4), 363–378, 2007.
- [13] GAO, D. How Does Dispersal Affect the Infection Size? SIAM Journal on Applied Mathematics, 80, 2144–2169, 2007.
- [14] GUO, H., LI, M. AND SHUAI, Z. Global stability of the endemic equilibrium of multigroup SIR epidemic models. Canadian Applied Mathematics Quarterly, 14(3), 259–284, 2006.
- [15] HANSKI, I. Metapopulation Ecology. Oxford University Press, 1999.
- [16] IGGIDR, A., SALLET, G. AND TSANOU, B. Global Stability Analysis of a Metapopulation SIS Epidemic Model. Mathematical Population Studies, 19(3), 115–129, 2012.
- [17] KEELING, M. J., BJRNSTAD, O. N. AND GRENFELL, B. T. Metapopulation dynamics of infectious diseases. Chapter 17 in I. Hanski, and O. Gaggiotti (eds.). Ecology, Genetics, and Evolution of Metapopulations, Elsevier, 415–445, 2004.
- [18] KEELING, M.J. AND ROHANI, P. Estimating Spatial Coupling in Epidemiological Systems: a Mechanistic Approach. Ecology Letters, 5, 20–29, 2002.
- [19] KEELING, M.J. AND ROHANI, P. Modeling Infectious Diseases in Humans and Animals. Princeton University Press, 2007.
- [20] KELLY, M.R., TIEN, J.H., EISENBERG, M.C AND LENHART, S. The impact of spatial arrangements on epidemic disease dynamics and intervention strategies. Journal of Biological Dynamics, 10, 222– 49, 2016.
- [21] KERMACK, W. AND MCKENDRICK, A. A contribution to the mathematical theory of epidemics.. Proceedings of the Royal Society, A 115, 700–721, 1927.
- [22] KNIPL, D. A new approach for designing disease intervention strategies in metapopulation models. Journal of Biological Dynamics, 10(1), 71–94, 2016.
- [23] LI, J., XIANG, T. AND HE, L. Modeling epidemic spread in transportation networks: A review, Journal of Traffic and Transportation Engineering, 8(2), 139–152, 2021.
- [24] LIPSHTAT, A., ALIMI, R. AND BEN-HORIN, Y. Commuting in metapopulation epidemic modeling. Scientific Reports, 11, 15198, 2021.
- [25] LIU, X. AND STECHLINSKI, P. Transmission dynamics of a switched multi-city model with transportrelated infections. Nonlinear Analysis: Real World Applications, 14(1), 264–279, 2013.

- [26] MACARTHUR, R.H. AND AND WILSON, E.O. The Theory of Island Biogeography, Princeton University Press, 2001.
- [27] MPOLYA, E.A., YASHIMA, K., OHTSUKI, H AND SASAKI, A. Epidemic dynamics of a vector-borne disease on a villages-and-city star network with commuters. Journal of Theoretical Biology, 343, 120–26, 2014.
- [28] MURRAY, J.D. Mathematical Biology II: Spatial Models and Biomedical Applications. Springer, 2003.
- [29] SATTENSPIEL, L. AND DIETZ, K. A structured epidemic model incorporating geographic mobility among regions. Mathematical Biosciences, 128 (1-2), 71–91, 1995.
- [30] TAKEUCHI, Y., LIU, X. AND CUI, J. Global dynamics of SIS models with transport-related infection. Journal of Mathematical Analysis and Applications, 329(2), 1460-71, 2007.
- [31] TOCTO-ERAZO, M.R., OLMOS-LICEAGA, D. AND MONTOYA-LAOS, J.A. Effect of daily human movement on some characteristics of dengue dynamics. Mathematical Biosciences, 332, 108531, 2021.
- [32] VAN DEN DRIESSCHE, P. AND WATMOUGH, J. Reproduction numbers and sub-threshold endemic equilibria for compartmental models of disease transmission. Mathematical Biosciences, 180 (1-2), 29–48, 2002.
- [33] VOROTNIKOV, V.I Partial Stability and Control, Birkäuser, 1998.
- [34] WANG, W. AND MULONE, G. Threshold of disease transmission in a patch environment. Journal of Mathematical Analysis and Applications, 285(1), 321–335, 2003.
- [35] WANG, W. AND ZHAO, X.Q. An epidemic model in a patchy environment. Mathematical Biosciences, 190(1), 97–112, 2004.
- [36] YIN, Q., WANG, Z., XIA, C., DEHMER, M., EMMERT-STREIB, F. AND JIN, Z. A novel epidemic model considering demographics and intercity commuting on complex dynamical networks. Applied Mathematics and Computation, 386, N. 125517, 2020.