



The multi-patch logistic equation with asymmetric migration

BILEL ELBETCH^a, TOUNSIA BENZEKRI^b,
DANIEL MASSART^c ✉, TEWFIK SARI^d

^aDepartment of Mathematics, University Dr. Moulay Tahar of Saida, Algeria.

^bDepartment of Mathematics, USTHB, Bab Ezzouar, Algiers, Algeria.

^cIMAG, Univ Montpellier, CNRS, Montpellier, France.

^dITAP, Univ Montpellier, INRAE, Institut Agro, Montpellier, France.

Abstract. This paper is a follow-up to a previous work where we considered a multi-patch model, each patch following a logistic law, the patches being coupled by symmetric migration terms. In this paper we drop the symmetry hypothesis. First, in the case of perfect mixing, i.e when the migration rate tends to infinity, the total population follows a logistic law with a carrying capacity which in general is different from the sum of the n carrying capacities, and depends on the migration terms. Second, we determine, in some particular cases, the conditions under which fragmentation and asymmetrical migration can lead to a total equilibrium population greater or smaller than the sum of the carrying capacities. Finally, for the three-patch model, we show numerically the existence of at least three critical values of the migration rate for which the total equilibrium population equals the sum of the carrying capacities.

Keywords: Population dynamics, asymmetrical migration, logistic equation, slow-fast systems, perfect mixing.

MSC2010: 37N25, 92D25, 34D23, 34D15.

E-mail: ^abilel.elbetch@univ-saida.dz, ^btbenzekri@usthb.dz, ^cdaniel.massart@umontpellier.fr ✉,
^dtewfik.sari@inrae.fr.

Received: 13 August 2021, Accepted: 16 November 2021.

To cite this article: B. Elbetch, T. Benzekri, D. Massart and T. Sari, The multi-patch logistic equation with asymmetric migration, *Rev. Integr. Temas Mat.*, 40 (2022), No. 1, 25–57. doi: 10.18273/revint.v40n1-2022002

La ecuación logística de múltiples sitios con migración asimétrica

Resumen. Este artículo es un seguimiento de un trabajo anterior, donde consideramos un modelo de múltiples parches, cada parche siguiendo una ley logística, los sitios estando relacionados por términos de migración simétrica. En este artículo eliminamos la hipótesis de simetría. Primero, en el caso de una mezcla perfecta, es decir, cuando la tasa de migración tiende al infinito, la población total sigue una ley logística con una capacidad de carga que en general es diferente de la suma de las capacidades de carga de los sitios, y depende de los términos de migración. En segundo lugar, determinamos, en algunos casos particulares, las condiciones bajo las cuales la fragmentación y la migración asimétrica pueden llevar a una población total de equilibrio mayor o menor que la suma de las capacidades de carga. Finalmente, para el modelo de tres sitios, mostramos numéricamente la existencia de al menos tres valores críticos de la tasa de migración para los cuales la población total de equilibrio es igual a la suma de las capacidades de carga.

Palabras clave: Dinámica de población, migración asimétrica, ecuación logística, sistemas lentos y rápidos, mezcla perfecta.

1. Introduction

The study of the dynamics of a fragmented population is fundamental in theoretical ecology, with potentially very important applied aspects: what is the effect of migration on the general population dynamics? What are the consequences of fragmentation on the persistence or extinction of the population? When is a single large refuge better or worse than several small ones (this is known as the SLOSS debate; see Hanski [19])?

The theoretical paradigm that has been used to treat these questions is that of a single population fragmented into patches coupled by migration, and the sub-population in each patch follows a local logistic law. This system is modeled by a non-linear system of differential equations of the following form:

$$\frac{dx}{dt} = f(x) + \beta\Gamma x, \quad (1)$$

where $x = (x_1, \dots, x_n)^T$, n is the number of patches in the system, x_i represents the population density in the i -th patch, $f(x) = (f_1(x_1), \dots, f_n(x_n))^T$, and

$$f_i(x_i) = r_i x_i (1 - x_i / K_i), \quad i = 1, \dots, n. \quad (2)$$

The parameters r_i and K_i are respectively the intrinsic growth rate and the carrying capacity of patch i .

The term $\beta\Gamma x$ on the right hand side of the system (1) describes the effect of the migration between the patches, where β is the migration rate and $\Gamma = (\gamma_{ij})$ is the matrix

representing the migrations between the patches. For $i \neq j$, $\gamma_{ij} > 0$ denotes the incoming flux from patch j to patch i . If $\gamma_{ij} = 0$, there is no migration. The diagonal entries of Γ satisfy the following equation

$$\gamma_{ii} = - \sum_{j=1, j \neq i}^n \gamma_{ji}, \quad i = 1, \dots, n, \tag{3}$$

which means that what comes out of a patch is distributed between the other $n - 1$ patches.

In the absence of migration, ($\beta = 0$), the system (1) admits (K_1, \dots, K_n) as a non-trivial equilibrium point. This equilibrium is globally asymptotically stable (GAS) and the total population at equilibrium is equal to the sum of the carrying capacities. The problem is whether or not the equilibrium continues to be positive and GAS, for any $\beta > 0$, and whether or not the total population at equilibrium can be greater than the sum of the carrying capacities. The case $n = 2$ and Γ symmetric

$$\Gamma = \begin{bmatrix} -1 & 1 \\ 1 & -1 \end{bmatrix},$$

where $\gamma_{12} = \gamma_{21}$ is normalized to 1 has been considered by Freedman and Waltman [14] and Holt [18]. They analyzed the model in the case of perfect mixing ($\beta \rightarrow +\infty$) and showed that the total equilibrium population can be greater than the sum of the carrying capacities $K_1 + K_2$, so that patchiness has a beneficial effect on the total equilibrium population. More recently, Arditi et al. [1] analyzed the behaviour of the system for all values of β . They showed that only three situations occur: either for any $\beta > 0$, patchiness has a beneficial effect, or this effect is always detrimental, or the effect is beneficial for lower values of the migration coefficient β and detrimental for higher values. Arditi et al. [2] extended these results to the case of two patches coupled by asymmetric migration, corresponding to the matrix

$$\Gamma = \begin{bmatrix} -\gamma_{21} & \gamma_{12} \\ \gamma_{21} & -\gamma_{12} \end{bmatrix}.$$

See also Poggiale et al. [25] who considered two patches coupled by asymmetric migration, in the particular case of perfect mixing. DeAngelis et al. [8, 11] considered the case of $n > 2$ patches in a circle, with symmetric migration between any patch and its two neighbours :

$$\frac{dx_i}{dt} = r_i x_i \left(1 - \frac{x_i}{K_i} \right) + \beta(x_{i-1} - 2x_i + x_{i+1}), \quad i = 1, \dots, n, \tag{4}$$

where we denote $x_0 = x_n$ and $x_{n+1} = x_1$, so that the same relationships hold between x_i , x_{i-1} and x_{i+1} for all values of i . This model corresponds to the matrix Γ whose non-zero off-diagonal elements are given by

$$\gamma_{1n} = \gamma_{n1} = 1 \quad \text{and} \quad \gamma_{i,i-1} = \gamma_{i-1,i} = 1, \quad \text{for} \quad 2 \leq i \leq n.$$

The system (4) is a one-dimensional discrete-patch version of the standard reaction-diffusion model. In [8, 11] the perfect mixing case is described.

In [12] we considered the general symmetric migration. We studied the system:

$$\frac{dx_i}{dt} = r_i x_i \left(1 - \frac{x_i}{K_i}\right) + \beta \sum_{j=1, j \neq i}^n \gamma_{ij} (x_j - x_i), \quad i = 1, \dots, n, \quad (5)$$

where $\beta\gamma_{ij}$ is the rate of migration between patches i and j . This system can be written in the form of System (1) with $\Gamma = (\gamma_{ij})$, the symmetric matrix whose diagonal entries are defined by (3). We studied the total population at equilibrium, as a function of the migration rate β . We gave conditions on the system parameters that ensure that migration is beneficial or detrimental, and extended several results of [1, 8, 11].

The aim of this work is to consider the case of n patches connected by asymmetric migration. Thus, we extend [2] by considering the case $n \geq 2$, and we extend [12] by considering the case where Γ is non-symmetric.

An important extension of (1) is the so called source-sink model, where the patches are of two types: the source patches, $1 \leq i \leq m$, with logistic dynamics, and the sink patches, $m + 1 \leq i \leq n$, with exponential decay

$$\begin{cases} f_i(x_i) = r_i x_i (1 - x_i/K_i), & i = 1, \dots, m, \\ f_i(x_i) = -r_i x_i, & i = m + 1, \dots, n. \end{cases} \quad (6)$$

The main problem is the number of source patches required for population persistence. For a recent study and bibliographical references the reader can consult Arino et al. [3] and Wu et al. [30].

There is another important extension of (1,2), where the dynamics on patch i is of the form

$$f_i(x_i) = r_i x_i (1 - x_i/K_i) - \gamma_i x_i, \quad i = 1, \dots, n, \quad (7)$$

with $\gamma_i > 0$. This model is the limit system (when $t \rightarrow +\infty$) of a susceptible-infected-susceptible (SIS) model in n patches connected by human migration. For details and further reading, see Section 5. Note that, when $r_i < \gamma_i$ for some patches, system (1,7) is a source-sink model. Contrary to (6), the mortality in sink patch is density-dependent. For more details and bibliographical references the reader is referred to [15].

Another example of source-sink model is the system considered by Nagahara et al. [24], called the ‘‘island chain’’ model, which is of the form:

$$\frac{dx_i}{dt} = x_i (m_i - x_i) + \beta (x_{i-1} - 2x_i + x_{i+1}), \quad i = 1, \dots, n, \quad (8)$$

where we denote $x_0 = x_1$ and $x_{n+1} = x_n$. This model is of the form (1), Γ being the matrix which verifies (3), and whose non-zero off-diagonal elements are given by

$$\gamma_{i,i-1} = \gamma_{i-1,i} = 1, \quad \text{for } 2 \leq i \leq n.$$

In the model (8) the ratios $\alpha_i = r_i/K_i$ in (2) are equal and are normalized to 1. The constant m_i represents both the intrinsic growth rate of the species in patch i and the carrying capacity of the patch. If $m_i > 0$, then patch i is favorable to the species. It is a source. The case $m_i = 0$ is permitted and corresponds to a sink. The main purpose

is to find the resource allocation (m_1, \dots, m_n) that maximizes the total population at equilibrium, under the constraint that $\sum_i m_i = m > 0$ is fixed. For more details and information on the maximization of the total population with logistic growth in a patchy environment, the reader is referred to [24] and the references therein.

For general information on the effects of patchiness and migration in both continuous and discrete cases, and the results beyond the logistic model, the reader is referred to the work of Levin [20, 21], DeAngelis et al. [8, 9, 10, 11], Freedman et al. [13], Zaker et al. [33].

It is worth noting that System (1) appears in metapopulation dynamics, involving explicit movements of the individuals between distinct locations. For the graph theoretic and dynamical system context in which metapopulation models are formulated, the reader is referred to Arino [4, Section 2].

The paper is organized as follows. In Section 2, the mathematical model of n patches, and some preliminaries results, are introduced. In Section 3, the behavior of the model is studied when the migration rate tends to infinity. In Section 4, we compare the total equilibrium population with the sum of the carrying capacities in some particular cases. In Section 5, the SIS patch model is considered, and the links with the logistic patch model are investigated. In Section 6 the three-patch model is considered, and by numerical simulations we show the existence of a new behavior for the dynamics of the total equilibrium population as a function of the migration rate. In Appendix A, we recall some results for the two-patch model with asymmetrical migration. In Appendix B, we prove some useful auxiliary results.

2. The mathematical model and preliminaries results

We consider the model of multi-patch logistic growth, coupled by asymmetric migration terms

$$\frac{dx_i}{dt} = r_i x_i \left(1 - \frac{x_i}{K_i}\right) + \beta \sum_{j=1, j \neq i}^n (\gamma_{ij} x_j - \gamma_{ji} x_i), \quad i = 1, \dots, n, \quad (9)$$

where $\gamma_{ij} \geq 0$ denotes the incoming flux from patch j to patch i , for $i \neq j$. The system (9) can be written in the form (1), where f is given by:

$$f(x) = (r_1 x_1 (1 - x_1/K_1), \dots, r_n x_n (1 - x_n/K_n))^T, \quad (10)$$

and $\Gamma := (\gamma_{ij})_{n \times n}$ is the matrix whose diagonal entries are given by (3). The matrix

$$\Gamma_0 := \Gamma - \text{diag}(\gamma_{11}, \dots, \gamma_{nn}),$$

which is the same as Γ , except that the diagonal elements are 0, is called the connectivity matrix. It is the adjacency matrix of the weighted directed graph \mathcal{G} , which has exactly n vertices (the patches), and has an arrow from patch j to patch i , with weight γ_{ij} , precisely when $\gamma_{ij} > 0$.

As to the non-negativity of the solution, we have the following proposition:

Proposition 2.1. *The domain $\mathbb{R}_+^n = \{(x_1, \dots, x_n) \in \mathbb{R}^n / x_i \geq 0, i = 1, \dots, n\}$ is positively invariant for the system (9).*

Proof. The proof is the same as in the symmetrical case [12, Proposition 2.1]. \square

When the connectivity matrix Γ_0 is irreducible, System (9) admits a unique positive equilibrium $(x_1^*(\beta), \dots, x_n^*(\beta))$, which is GAS, see [4, Theorem 2.2], [3, Theorem 1] or [12, Theorem 6.1]. In all of this work, we denote by $E^*(\beta)$ the positive equilibrium and by $X_T^*(\beta)$ the total population at equilibrium:

$$E^*(\beta) = (x_1^*(\beta), \dots, x_n^*(\beta)), \quad X_T^*(\beta) = \sum_{i=1}^n x_i^*(\beta). \quad (11)$$

Remark 2.2. The matrix Γ_0 being irreducible means that the weighted directed graph \mathcal{G} is strongly connected, which means that every patch is reachable from every other patch, either directly or through other patches. The matrix Γ is assumed to be irreducible throughout the rest of the paper.

3. Perfect mixing

In this section our aim is to study the behavior of $E^*(\beta)$ and $X_T^*(\beta)$, defined by (11), for large migration rate, i.e when $\beta \rightarrow \infty$.

3.1. The fast dispersal limit

The following lemma was proved in [3, Lemma 2]; we include a proof for the ease of the reader.

Lemma 3.1. *Let Γ be the migration matrix. Then, 0 is a simple eigenvalue of Γ and all non-zero eigenvalues of Γ have negative real part. Moreover, the kernel of the matrix Γ is generated by a positive vector. If the matrix Γ is symmetric, then $\ker \Gamma$ is generated by $u = (1, \dots, 1)^T$.*

Proof. Let $s = \max_{i=1, \dots, n} (-\gamma_{ii})$ and let B be the matrix defined by

$$B = \Gamma + sI.$$

First, we note that since the matrix Γ verifies the property (3), then Γ is a singular matrix and the vector $u = (1, \dots, 1)^T$ is an eigenvector of Γ^T associated to the eigenvalue 0. Thus u is an eigenvector of B^T , with eigenvalue s .

The matrix B^T is non-negative and irreducible, so by the Perron-Frobenius Theorem the spectral radius

$$\rho(B^T) = \max \{ |\lambda| : \lambda \text{ is an eigenvalue of } B^T \},$$

is a simple eigenvalue of the matrix B^T and it is the only eigenvalue of B^T which admits a positive eigenvector, so $s = \rho(B^T) = \rho(B)$. Therefore, $\Gamma = B - \rho(B)I$ and $\dim(\ker \Gamma) = \dim(\ker \Gamma^T) = 1$.

All other eigenvalues of B have modulus $< \rho(B)$, so their real parts are $< \rho(B)$. Since each eigenvalue of Γ is $\lambda - \rho(B)$, for some eigenvalue λ of B , all eigenvalues of Γ have negative real part.

Furthermore, according to the Perron-Frobenius theorem, there exists a positive vector δ such that $B\delta = \rho(B)\delta$, that is, $\Gamma\delta = (B - \rho(B)I)\delta = 0$. In particular, if the matrix Γ is symmetric then we may take $\delta = u$, that is, $\delta_i = 1$, for all i . \square

In all of this paper, we denote by $\delta = (\delta_1, \dots, \delta_n)^T$ a positive vector which generates the vector space $\ker \Gamma$.

Remark 3.2. The existence, uniqueness (mod. multiplicative factor), and positivity of δ were also proved in Lemma 1 of Cosner et al. [5]. On the other hand, it is shown in Guo et al. [17, Lemma 2.1] and Gao and Dong [16, Lemma 3.1] that the vector $(\Gamma_{11}^*, \dots, \Gamma_{nn}^*)^T$ is a right eigenvector of Γ associated with the zero eigenvalue. Here, Γ_{ii}^* is the cofactor of the i -th diagonal entry of Γ . Therefore, we have explicit formulae for the components of the vector δ , as functions of the coefficients of Γ , at our disposal. For two patches we have $\delta = (\gamma_{12}, \gamma_{21})^T$, and for three patches we have $\delta = (\delta_1, \delta_2, \delta_3)^T$, where

$$\begin{cases} \delta_1 = \gamma_{12}\gamma_{13} + \gamma_{12}\gamma_{23} + \gamma_{32}\gamma_{13}, \\ \delta_2 = \gamma_{21}\gamma_{13} + \gamma_{21}\gamma_{23} + \gamma_{31}\gamma_{23}, \\ \delta_3 = \gamma_{21}\gamma_{32} + \gamma_{31}\gamma_{12} + \gamma_{31}\gamma_{32}. \end{cases} \tag{12}$$

The following result asserts that when $\beta \rightarrow \infty$, the equilibrium $E^*(\beta)$ converges to an element of $\ker \Gamma$.

Theorem 3.3. For the system (9), we have

$$\lim_{\beta \rightarrow +\infty} E^*(\beta) = \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i^2 \alpha_i} (\delta_1, \dots, \delta_n),$$

where $\alpha_i = r_i/K_i$.

Proof. Denote

$$E^*(\infty) = \left(\delta_1 \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i^2 \alpha_i}, \dots, \delta_n \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i^2 \alpha_i} \right).$$

Dividing Equation 1 at the equilibrium $E^*(\beta)$ by β , for $\beta > 0$, yields

$$\text{for all } \beta > 0, \quad \frac{1}{\beta} f(E^*(\beta)) + \Gamma E^*(\beta) = 0.$$

Thus any limit point, when $\beta \rightarrow \infty$, of the set $\{E^*(\beta) : \beta > 0\}$ lies in the kernel of Γ . Now, taking the sum of all equations in

$$r_i x_i \left(1 - \frac{x_i}{K_i} \right) + \beta \sum_{j=1, j \neq i}^n (\gamma_{ij} x_j - \gamma_{ji} x_i) = 0, \quad i = 1, \dots, n,$$

we see that $E^*(\beta)$ lies in the ellipsoid

$$\mathbb{E}^{n-1} = \left\{ x \in \mathbb{R}^n : \Theta(x) := \sum_{i=1}^n r_i x_i \left(1 - \frac{x_i}{K_i} \right) = 0 \right\}.$$

The ellipsoid \mathbb{E}^{n-1} is compact, so the equilibrium $E^*(\beta)$ has at least one limit point in \mathbb{E}^{n-1} , when β goes to infinity. Since the kernel of Γ has dimension 1, and \mathbb{E}^{n-1} is the boundary of a convex set, $\mathbb{E}^{n-1} \cap \ker \Gamma$ consists of at most two points. Since the origin and $E^*(\infty)$ both lie in $\mathbb{E}^{n-1} \cap \ker \Gamma$, we get that

$$\mathbb{E}^{n-1} \cap \ker \Gamma = \{0, E^*(\infty)\}.$$

Therefore, to prove the convergence of $E^*(\beta)$ to $E^*(\infty)$, it suffices to prove that the origin cannot be a limit point of $E^*(\beta)$. We claim that for any β , there exists i such that $x_i^*(\beta) \geq K_i$, which entails that $E^*(\beta)$ is bounded away from the origin. The coordinates of the vector $\Gamma E^*(\beta)$ sum to zero, hence at least one of them, say, the i -th, is non-negative. Then

$$r_i x_i^*(\beta) \left(1 - \frac{x_i^*(\beta)}{K_i}\right) \leq 0,$$

and since $x_i^*(\beta)$ cannot be negative or 0, we have $x_i^*(\beta) \geq K_i$. \square

As a corollary of the previous theorem, we obtain the following result, which describes the total equilibrium population for perfect mixing:

Proposition 3.4. *We have*

$$X_T^*(+\infty) = \lim_{\beta \rightarrow +\infty} \sum_{i=1}^n x_i^*(\beta) = \left(\sum_{i=1}^n \delta_i \right) \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i^2 \alpha_i}. \quad (13)$$

Denote $K = (K_1, \dots, K_n)^T$. If $K = \lambda \delta$ with $\lambda > 0$, that is to say $K \in \ker \Gamma$, then $X_T^*(+\infty) = \lambda \sum_{i=1}^n \delta_i = \sum_{i=1}^n K_i$.

Proof. For the proof of (13), it suffices to sum the n components of the point $E^*(\infty)$. For the case $K \in \ker \Gamma$, it suffices to replace K_i by $\lambda \delta_i$ in (13). \square

Actually, when $K \in \ker \Gamma$, we have $X_T^*(\beta) = \sum_i K_i$ for all $\beta > 0$, see Proposition 4.5.

In the case $n = 2$, one has $\delta_1 = \gamma_{12}$ and $\delta_2 = \gamma_{21}$, as shown in Remark 3.2. Therefore, (13) becomes

$$X_T^*(+\infty) = (\gamma_{12} + \gamma_{21}) \frac{\gamma_{12} r_1 + \gamma_{21} r_2}{\gamma_{12}^2 \alpha_1 + \gamma_{21}^2 \alpha_2},$$

which is the formula given by Arditi et al. [2, Equation 7] and by Poggiale et al. [25, page 362].

If the matrix Γ is symmetric, one has $\delta_i = 1$, for all i , as shown in Lemma 3.1. Therefore, (13) specializes to the formula given in [12, Equation (24)]:

$$X_T^*(+\infty) = n \frac{\sum_{i=1}^n r_i}{\sum_{i=1}^n r_i / K_i}.$$

3.2. Two time scale dynamics

In [12] we also obtained the formula (13), in the symmetrical n-patch case (i.e the matrix Γ is symmetric), by using singular perturbation theory, see [12, Theorem 4.6].

We showed that, if $(x_1(t, \beta), \dots, x_n(t, \beta))$ is the solution of (5), with initial condition (x_1^0, \dots, x_n^0) , then, when $\beta \rightarrow \infty$, the total population $\sum x_i(t, \beta)$ is approximated by $X(t)$, the solution of the logistic equation

$$\frac{dX}{dt} = rX \left(1 - \frac{X}{nK} \right), \text{ where } r = \frac{\sum_{i=1}^n r_i}{n}, \quad K = \frac{\sum_{i=1}^n r_i}{\sum_{i=1}^n \alpha_i} \text{ and } \alpha_i = \frac{r_i}{K_i}, \quad (14)$$

with initial condition $X_0 = \sum x_i^0$. Therefore, the total population behaves like the solution of the logistic equation given by (14). In addition, one obtains the following property: with the exception of a small initial interval, the population densities $x_i(t, \beta)$ are approximated by $X(t)/n$, see [12, Formula (37)]. Therefore, this approximation shows that, when t and β tend to ∞ , the population density $x_i(t, \beta)$ tends toward $\frac{\sum r_i}{\sum \alpha_i}$, and in addition, $x_i(t, \beta)$ quickly jumps from its initial condition x_i^0 to the average X_0/n and then is very close to $X(t)/n$. Our aim is to generalize this result for the asymmetrical n-patch model (9) (i.e the matrix Γ is non-symmetric). To avoid any confusion with $X(t)$, which is the total population, we denote $Y(t)$ the solution of the logistic equation (15), and we prove that $X(t)$ is asymptotically equivalent, when β goes to infinity, to $Y(t)$. We have the following result

Theorem 3.5. *Let $(x_1(t, \beta), \dots, x_n(t, \beta))$ be the solution of the system (9) with initial condition (x_1^0, \dots, x_n^0) satisfying $x_i^0 \geq 0$ for $i = 1 \dots n$. Let $Y(t)$ be the solution of the logistic equation*

$$\frac{dX}{dt} = rX \left(1 - \frac{X}{[\sum_{i=1}^n \delta_i] K} \right), \quad (15)$$

where

$$r = \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i}, \quad K = \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i^2 \alpha_i} \text{ and } \alpha_i = \frac{r_i}{K_i}, \quad (16)$$

with initial condition $X_0 = \sum_{i=1}^n x_i^0$. Then, when $\beta \rightarrow \infty$, we have

$$\sum_{i=1}^n x_i(t, \beta) = Y(t) + o(1), \quad \text{uniformly for } t \in [0, +\infty) \quad (17)$$

and, for any $t_0 > 0$, we have

$$x_i(t, \beta) = \frac{\delta_i}{\sum_{i=1}^n \delta_i} Y(t) + o(1), \quad i = 1, \dots, n, \text{ uniformly for } t \in [t_0, +\infty). \quad (18)$$

Proof. Let $X(t, \beta) = \sum_{i=1}^n x_i(t, \beta)$. We rewrite the system (9) using the variables (X, x_1, \dots, x_{n-1}) , and get:

$$\begin{cases} \frac{dX}{dt} = \sum_{i=1}^n r_i x_i \left(1 - \frac{x_i}{K_i} \right), \\ \frac{dx_i}{dt} = r_i x_i \left(1 - \frac{x_i}{K_i} \right) + \beta \sum_{j=1, j \neq i}^n (\gamma_{ij} x_j - \gamma_{ji} x_i), \quad i = 1, \dots, n-1. \end{cases} \quad (19)$$

This system is actually a system in the variables (X, x_1, \dots, x_{n-1}) , since, whenever x_n appears in the right hand side of (19), it should be replaced by

$$x_n = X - \sum_{i=1}^{n-1} x_i. \quad (20)$$

When $\beta \rightarrow \infty$, (19) is a *slow-fast* system, with one *slow variable*, X , and $n - 1$ *fast variables*, x_i for $i = 1 \cdots n - 1$. As suggested by Tikhonov's Theorem [22, 28, 31], we consider the dynamics of the fast variables in the time scale $\tau = \beta t$. We get

$$\frac{dx_i}{d\tau} = \frac{1}{\beta} r_i x_i \left(1 - \frac{x_i}{K_i}\right) + \sum_{j=1, j \neq i}^n (\gamma_{ij} x_j - \gamma_{ji} x_i), \quad i = 1, \dots, n - 1.$$

where x_n is given by (20). In the limit $\beta \rightarrow \infty$, we find the *fast dynamics*

$$\frac{dx_i}{d\tau} = \sum_{j=1, j \neq i}^n (\gamma_{ij} x_j - \gamma_{ji} x_i), \quad i = 1, \dots, n - 1.$$

This is an $(n - 1)$ -dimensional linear differential system in the variable $Z := (x_1, \dots, x_{n-1})$, which can be rewritten in matricial form:

$$\dot{Z} = \mathcal{L}Z + XV, \quad \text{with} \quad \mathcal{L} := L - U, \quad (21)$$

where $L := (\gamma_{ij})_{n-1 \times n-1}$ is the sub matrix of the matrix Γ , obtained by dropping the last row and the last column of Γ , V is the vector defined by $V := (\gamma_{in})_{n-1 \times 1}$ and $U = (V; \dots; V)$.

By Lemma B.1, the matrix \mathcal{L} is stable, that is, all of its eigenvalues have negative real part. Therefore, it is invertible and the equilibrium of the system (21) is GAS. This equilibrium is given by

$$\left(\frac{\delta_1}{\sum_{i=1}^n \delta_i} X, \dots, \frac{\delta_{n-1}}{\sum_{i=1}^n \delta_i} X \right)^T.$$

Indeed, we denote by $L^{(i)}, U^{(i)}$ and $V^{(i)}$ the i -th row of the matrix L, U and the vector V respectively. We have:

$$\begin{aligned} \frac{\delta_n}{\sum_{i=1}^n \delta_i} \left(L^{(i)} - U^{(i)} \right) \left(\frac{\delta_1}{\delta_n} X \quad \dots \quad \frac{\delta_{n-1}}{\delta_n} X \right)^T &= -\frac{\delta_n}{\sum_{i=1}^n \delta_i} X \gamma_{in} - \frac{\sum_{i=1}^{n-1} \delta_i}{\sum_{i=1}^n \delta_i} X \gamma_{in} \\ &= -X \gamma_{in} = -X V^{(i)}. \end{aligned}$$

Thus, the slow manifold of System (19) is given by

$$x_i = \frac{\delta_i}{\sum_{i=1}^n \delta_i} X, \quad i = 1, \dots, n - 1. \quad (22)$$

As this manifold is GAS, Tikhonov's Theorem ensures that after a fast transition toward the slow manifold, the solutions of (19) are approximated by the solutions of the *reduced*

model, which is obtained by replacing (22) into the dynamics of the slow variable, that is:

$$\frac{dX}{dt} = \sum_{i=1}^n r_i \frac{X}{\sum_{i=1}^n \delta_i} \delta_i \left(1 - \frac{X}{(\sum_{i=1}^n \delta_i) K_i} \delta_i \right) = rX \left(1 - \frac{X}{(\sum_{i=1}^n \delta_i) K} \right),$$

where r and K are defined in (16). Therefore, the reduced model is (15). Since (15) admits

$$X^* = \left(\sum_{i=1}^n \delta_i \right) K = \left(\sum_{i=1}^n \delta_i \right) \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i^2 \alpha_i}$$

as a positive equilibrium point, which is GAS in the positive axis, the approximation given by Tikhonov's Theorem holds for all $t \geq 0$ for the slow variable and for all $t \geq t_0 > 0$ for the fast variables, where t_0 is as small as we want. Therefore, letting $Y(t)$ be the solution of the reduced model (15) with initial condition $Y(0) = X(0, \beta) = \sum_{i=1}^n x_i^0$, then, when $\beta \rightarrow \infty$, we have the approximations (17) and (18). \square

In the case of perfect mixing, the approximation (17) shows that the total population behaves like the solution of the single logistic equation (16) and then, when t and β tend to ∞ , the total population $\sum x_i(t, \beta)$ tends toward $(\sum_{i=1}^n \delta_i) K = (\sum_{i=1}^n \delta_i) \frac{\sum \delta_i r_i}{\sum \delta_i^2 \alpha_i}$ as stated in Proposition 3.4. The approximation (18) shows that, with the exception of a thin initial boundary layer, where the population density $x_i(t, \beta)$ quickly jumps from its initial condition x_i^0 to $\delta_i X_0 / \sum_{i=1}^n \delta_i$, each patch of the n-patch model behaves like the logistic equation

$$\frac{du}{dt} = ru \left(1 - \frac{u}{\delta_i K} \right) \text{ where } r = \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i}, \quad K = \frac{\sum_{i=1}^n \delta_i r_i}{\sum_{i=1}^n \delta_i^2 \alpha_i}, \quad \alpha_i = \frac{r_i}{K_i}. \quad (23)$$

Hence, when t and β tend to ∞ , the population density $x_i(t, \beta)$ tends toward $\delta_i \frac{\sum \delta_i r_i}{\sum \delta_i^2 \alpha_i}$, as stated in Theorem 3.3.

Remark 3.6. The single logistic equation (23) gives an approximation of the population density in each patch in the case of perfect mixing. The intrinsic growth rate r in (23) is the arithmetic mean of the r_1, \dots, r_n , weighted by $\delta_1, \dots, \delta_n$, and the carrying capacity K is the harmonic mean of K_i/δ_i , weighted by $\delta_i r_i, i = 1, \dots, n$. We point out the similarity between our expression for the carrying capacity in the limit $\beta \rightarrow \infty$, and the expression obtained in spatial homogenization, see e.g [32, Formula 81] and also [33, Formula 28].

3.3. Comparison of $X_T^*(+\infty)$ with $\sum_i K_i$.

According to Formula (13), it is clear that the total equilibrium population at $\beta = 0$ and at $\beta = +\infty$ are different in general.

In the remainder of this section, we give some conditions, in the space of parameters r_i, K_i, α_i and δ_i , for limit of the total equilibrium population when $\beta \rightarrow \infty$ to be greater or smaller than the sum of the carrying capacities. We show that all three cases are possible, i.e $X_T^*(+\infty)$ can be greater than, smaller than, or equal to $X_T^*(0)$. First, we start by giving some particular values of the parameters for which equality holds.

Proposition 3.7. Consider the system (9). If the vector $\left(\frac{1}{\alpha_1}, \dots, \frac{1}{\alpha_n}\right)^T$ lies in $\ker \Gamma$, then $X_T^*(+\infty) = \sum_i K_i$.

Proof. It is a direct consequence of the Equation (13). \square

Note that, if the matrix Γ is symmetric, then by Lemma 3.1, Proposition 3.7 says that if all α_i are equal, then $X_T^*(\infty) = \sum_i K_i$, which is [12, Proposition 4.4].

In the next proposition, we give two cases which ensure that $X_T^*(0)$ can be greater or smaller than $X_T^*(+\infty)$. This result can be stated as the following proposition:

Proposition 3.8. Consider the system (9).

1. If $\frac{K_1}{\delta_1} \leq \dots \leq \frac{K_n}{\delta_n}$ and $\delta_1 \alpha_1 \leq \dots \leq \delta_n \alpha_n$, or if $\frac{K_1}{\delta_1} \geq \dots \geq \frac{K_n}{\delta_n}$ and $\delta_1 \alpha_1 \geq \dots \geq \delta_n \alpha_n$, then $X_T^*(+\infty) \geq X_T^*(0)$.
2. If $\frac{K_1}{\delta_1} \geq \dots \geq \frac{K_n}{\delta_n}$ and $\delta_1 \alpha_1 \leq \dots \leq \delta_n \alpha_n$, or if $\frac{K_1}{\delta_1} \leq \dots \leq \frac{K_n}{\delta_n}$ and $\delta_1 \alpha_1 \geq \dots \geq \delta_n \alpha_n$, then $X_T^*(+\infty) \leq X_T^*(0)$.

In both items, if at least one of the inequalities in $\frac{K_1}{\delta_1} \leq \dots \leq \frac{K_n}{\delta_n}$ or $\frac{K_1}{\delta_1} \geq \dots \geq \frac{K_n}{\delta_n}$ is strict, then the inequality is strict in the conclusion.

Proof. Apply Lemma B.2 with the following choice: $w_i = \delta_i$, $u_i = \frac{K_i}{\delta_i}$, and $v_i = \delta_i \alpha_i$, for all $i = 1, \dots, n$. \square

If the matrix Γ is symmetric, one has $\delta_i = 1$, for all i , as shown in Lemma 3.1. Therefore Proposition 3.8 becomes

Corollary 3.9. Consider the system (9). Assume that Γ is symmetric.

1. If $K_1 \leq \dots \leq K_n$ and $\alpha_1 \leq \dots \leq \alpha_n$, or if $K_1 \geq \dots \geq K_n$ and $\alpha_1 \geq \dots \geq \alpha_n$, then $X_T^*(+\infty) \geq X_T^*(0)$.
2. If $K_1 \geq \dots \geq K_n$ and $\alpha_1 \leq \dots \leq \alpha_n$, or if $K_1 \leq \dots \leq K_n$ and $\alpha_1 \geq \dots \geq \alpha_n$, then $X_T^*(+\infty) \leq X_T^*(0)$.

This result implies Items 1 and 2 of [10, Theorem B.1], which were obtained for the model (4) in the particular case $r_i = K_i$.

4. Influence of asymmetric dispersal on total population size

In this section, we will compare, in some particular cases of the System (9), the total equilibrium population $X_T^*(\beta) = x_1^*(\beta) + \dots + x_n^*(\beta)$, with the sum of carrying capacities denoted by $X_T^*(0) = K_1 + \dots + K_n$, when the rate of migration β varies from zero to infinity. We show that the total equilibrium population, $X_T^*(\beta)$, is generally different from the sum of the carrying capacities $X_T^*(0)$. Depending on the local parameters of the patches and the kernel of the matrix Γ , $X_T^*(\beta)$ can either be greater than, smaller than, or equal to the sum of the carrying capacities.

4.1. Asymmetric dispersal may be unfavorable to the total equilibrium population

When Γ is symmetric, we have already proved that if all the growth rates are equal then dispersal is always unfavorable to the total equilibrium population, see [12, Proposition 3.1]. We also noticed that the result still holds in the general case when Γ is not necessarily symmetric, see [12, Proposition 6.2]. Hence we have the following

Proposition 4.1. *If $r_1 = \dots = r_n$ then*

$$X_T^*(\beta) = \sum_{i=1}^n x_i^*(\beta) \leq \sum_{i=1}^n K_i, \quad \text{for all } \beta \geq 0.$$

For a two-patch logistic model, this result has been proved by Arditi et al. [1, Proposition 2, item 3] for symmetric dispersal and for asymmetric dispersal [2, Proposition 1, item 3].

4.2. Asymmetric dispersal may be favorable to the total equilibrium population

In this section, we give a situation where the dispersal is favorable to the total equilibrium population. Mathematically speaking:

Proposition 4.2. *Assume that for all $j < i$, $\alpha_i \gamma_{ij} = \alpha_j \gamma_{ji}$. Then*

$$X_T^*(\beta) \geq \sum_{i=1}^n K_i \quad \text{for all } \beta \geq 0.$$

Moreover, if there exist i_0 and $j_0 \neq i_0$ such that $r_{i_0} \neq r_{j_0}$, then $X_T^*(\beta) > \sum_{i=1}^n K_i$, for all $\beta > 0$.

Proof. The equilibrium point $E^*(\beta)$ satisfies the system

$$0 = \alpha_i x_i^*(\beta) (K_i - x_i^*(\beta)) + \beta \sum_{j=1, j \neq i}^n (\gamma_{ij} x_j^*(\beta) - \gamma_{ji} x_i^*(\beta)), \quad i = 1 \dots n. \quad (24)$$

Dividing (24) by $\alpha_i x_i^*$, one obtains

$$x_i^*(\beta) = K_i + \beta \sum_{j=1, j \neq i}^n \frac{\gamma_{ij} x_j^*(\beta) - \gamma_{ji} x_i^*(\beta)}{\alpha_i x_i^*(\beta)}.$$

Taking the sum of these expressions shows that the total equilibrium population X_T^* satisfies the following relation:

$$\begin{aligned} X_T^*(\beta) &= \sum_{i=1}^n K_i + \beta \sum_{i=1}^n \sum_{j=1, j \neq i}^n \frac{\gamma_{ij} x_j^*(\beta) - \gamma_{ji} x_i^*(\beta)}{\alpha_i x_i^*(\beta)} \\ &= \sum_{i=1}^n K_i + \beta \sum_{j < i} \left(\frac{\gamma_{ij} x_j^*(\beta) - \gamma_{ji} x_i^*(\beta)}{\alpha_i x_i^*(\beta)} + \frac{\gamma_{ji} x_i^*(\beta) - \gamma_{ij} x_j^*(\beta)}{\alpha_j x_j^*(\beta)} \right) \\ &= \sum_{i=1}^n K_i + \beta \sum_{j < i} \frac{(\gamma_{ij} x_j^*(\beta) - \gamma_{ji} x_i^*(\beta)) (\alpha_j x_j^*(\beta) - \alpha_i x_i^*(\beta))}{\alpha_j \alpha_i x_j^*(\beta) x_i^*(\beta)}. \end{aligned} \quad (25)$$

The conditions $\alpha_i \gamma_{ij} = \alpha_j \gamma_{ji}$ can be written $\kappa_{ij} := \alpha_i / \gamma_{ji} = \alpha_j / \gamma_{ij}$ for all $j < i$, such that $\gamma_{ij} \neq 0$ and $\gamma_{ji} \neq 0$. Therefore, there exists $\kappa_{ij} > 0$ such that

$$\alpha_j = \kappa_{ij} \gamma_{ij} \text{ and } \alpha_i = \kappa_{ij} \gamma_{ji} \text{ for all } i, j \text{ with } \gamma_{ij} \neq 0 \text{ and } \gamma_{ji} \neq 0.$$

Replacing α_i and α_j in (25), one obtains

$$X_T^*(\beta) = \sum_{i=1}^n K_i + \beta \sum_{j < i} \frac{\kappa_{ij} (\gamma_{ij} x_j^*(\beta) - \gamma_{ji} x_i^*(\beta))^2}{\alpha_j \alpha_i x_j^*(\beta) x_i^*(\beta)} \geq \sum_{i=1}^n K_i. \quad (26)$$

Equality holds if and only if $\beta = 0$ or $\gamma_{ij} x_j^*(\beta) - \gamma_{ji} x_i^*(\beta) = 0$, for all i and j . Let us prove that if at least two patches have different growth rates, then equality cannot hold for $\beta > 0$. Suppose that there exists $\beta^* > 0$ such that the positive equilibrium satisfies

$$\forall i, j, \quad \gamma_{ij} x_j^*(\beta^*) = \gamma_{ji} x_i^*(\beta^*). \quad (27)$$

Replacing the Equation (27) in the system (24), we get that $x_i^*(\beta^*) = K_i$, for all i . Therefore, from (27), it is seen that, for all i and j , $K_j \gamma_{ij} = K_i \gamma_{ji}$. From these equations and the conditions $\alpha_i \gamma_{ij} = \alpha_j \gamma_{ji}$, we get $r_i = r_j$, for all i and j . This is a contradiction with the hypothesis that there exist two patches with different growth rates. Hence the equality in (26) holds if and only if $\beta = 0$. \square

When the matrix Γ is irreducible and symmetric, the hypothesis of Proposition 4.2 implies that $\alpha_i = \alpha_j$ for all i and j . Indeed if two patches i and j are connected (i.e. $\gamma_{ij} = \gamma_{ji} \neq 0$), then we have $\alpha_i = \alpha_j$. As the matrix Γ is irreducible, for two arbitrary patches, there exists a finite sequence (i, \dots, j) which begins in i and ends in j , such that $\gamma_{ab} \neq 0$ for all successive patches a and b in (i, \dots, j) . Hence $\alpha_a = \alpha_b$ for all a and b in (i, \dots, j) . Hence, $\alpha_i = \alpha_j$. So, when the matrix Γ is symmetric, Proposition 4.2 says that if all α_i are equal, dispersal enhances population growth, which is [12, Proposition 3.3].

Note that, when $n = 2$, Proposition 4.2 asserts that if $\alpha_2 / \alpha_1 = \gamma_{12} / \gamma_{21}$, then $X_T^*(\beta) > K_1 + K_2$, which is a result of Arditi et al. [2, Proposition 2, item b)]. See also Proposition A.1, and note that the condition $\alpha_2 / \alpha_1 = \gamma_{12} / \gamma_{21}$ implies that $(\gamma_{12}, \gamma_{21}) \in \mathcal{J}_0$.

For three patches or more, if the matrix Γ does not verify the condition ($\forall i, j, \gamma_{ij} = 0 \iff \gamma_{ji} = 0$), then the hypothesis of Proposition 4.2, that for all $j < i, \alpha_i \gamma_{ij} = \alpha_j \gamma_{ji}$ cannot be satisfied. Note that the hypothesis $\alpha_i \gamma_{ij} = \alpha_j \gamma_{ji}$ implies that, for all $i = 1, \dots, n$, one has

$$\sum_{j=1}^n \frac{\gamma_{ij}}{\alpha_j} = \sum_{j=1, j \neq i}^n \frac{\gamma_{ij}}{\alpha_j} - \sum_{j=1, j \neq i}^n \frac{\gamma_{ji}}{\alpha_i} = \sum_{j=1, j \neq i}^n \frac{\alpha_i \gamma_{ij} - \alpha_j \gamma_{ji}}{\alpha_i \alpha_j} = 0.$$

Therefore we can make the following remark:

Remark 4.3. The hypothesis of Proposition 4.2 implies that $(\frac{1}{\alpha_1}, \dots, \frac{1}{\alpha_n})^T \in \ker \Gamma$.

We make the following conjecture:

Conjecture. If $(\frac{1}{\alpha_1}, \dots, \frac{1}{\alpha_n})^T \in \ker \Gamma$ then

$$X_T^*(\beta) \geq \sum_{i=1}^n K_i, \quad \text{for all } \beta \geq 0.$$

This conjecture is true for the particular case of Proposition 4.2. It is also true for two-patch models and for n -patch models with symmetric dispersal. It agrees with Proposition 3.7.

Proposition 4.4. *The derivative of the total equilibrium population $X_T^*(\beta)$ at $\beta = 0$ is given by:*

$$\frac{dX_T^*}{d\beta}(0) = \sum_{i=1}^n \left(\frac{1}{r_i} \sum_{j=1}^n \gamma_{ij} K_j \right). \tag{28}$$

In particular, if $K \in \ker \Gamma$, where $K = (K_1, \dots, K_n)^T$, then $\frac{dX_T^*}{d\beta}(0) = 0$.

Proof. By differentiating the Equation (25) at $\beta = 0$, we get:

$$\frac{dX_T^*}{d\beta}(0) = \sum_{i=1}^n \sum_{j=1, j \neq i}^n \frac{\gamma_{ij} x_j^*(0) - \gamma_{ji} x_i^*(0)}{\alpha_i x_i^*(0)},$$

which gives (28), since $x_i^*(0) = K_i$ for all $i = 1, \dots, n$.

If $K \in \ker \Gamma$, then $\sum_{j=1}^n \gamma_{ij} K_j = 0$ for all i , so that $\frac{dX_T^*}{d\beta}(0) = 0$. ☑

Actually, when $K \in \ker \Gamma$, we prove that $X_T^*(\beta)$ is constant, so that $\frac{dX_T^*}{d\beta}(\beta) = 0$ for all $\beta \geq 0$, not only for $\beta = 0$, see Proposition 4.5.

4.3. Independence of the total equilibrium population with respect to asymmetric dispersal

In the next proposition we give sufficient and necessary conditions for the total equilibrium population not to depend on the migration rate.

Proposition 4.5. *The equilibrium $E^*(\beta)$ does not depend on β if and only if $(K_1, \dots, K_n)^T \in \ker \Gamma$. In this case, we have $E^*(\beta) = (K_1, \dots, K_n)$ for all $\beta > 0$.*

Proof. The equilibrium $E^*(\beta)$ is the unique positive solution of the equation

$$f(x) + \beta \Gamma x = 0, \tag{29}$$

where f is given by (10). Suppose that the equilibrium $E^*(\beta)$ does not depend on β , then we replace in Equation (29):

$$f(E^*(\beta)) + \beta \Gamma E^*(\beta) = 0. \tag{30}$$

The derivative of (30) with respect to β gives

$$\Gamma E^*(\beta) = 0. \tag{31}$$

Replacing the Equation (31) in the Equation (30), we get $f(E^*(\beta)) = 0$, so $E^*(\beta) = (K_1, \dots, K_n)$. From the Equation (31), we conclude that $(K_1, \dots, K_n)^T \in \ker \Gamma$.

Now, suppose that $(K_1, \dots, K_n)^T \in \ker \Gamma$, then (K_1, \dots, K_n) satisfies the Equation (29), for all $\beta \geq 0$. So, $E^*(\beta) = (K_1, \dots, K_n)$, for all $\beta \geq 0$, which proves that the total equilibrium population is independent of the migration rate β . ☑

If the matrix Γ is symmetric, the previous proposition asserts that the K_i , for $i = 1, \dots, n$, are equal if and only if $E^* = (K, \dots, K)$, where K is the common value of the K_i . This is [12, Proposition 3.2]. For $n = 2$, Proposition 4.5 asserts that if $K_1/K_2 = \gamma_{12}/\gamma_{21}$ then $X_T^*(\beta) = K_1 + K_2$ for all β , which is [2, Proposition 2, item c]. See also the last item of Proposition A.1.

4.4. Two blocks of identical patches

We consider the model (9) and we assume that there are two blocks, denoted I and J , of identical patches, such that $I \cup J = \{1, \dots, n\}$. Let p be the number of patches in I and $q = n - p$ be the number of patches in J . Without loss of generality we can take $I = \{1, \dots, p\}$ and $J = \{p + 1, \dots, n\}$. The patches being identical means that they have the same specific growth rate r_i and carrying capacity K_i . Therefore we have

$$\begin{aligned} r_1 = \dots = r_p, & & K_1 = \dots = K_p, \\ r_{p+1} = \dots = r_n, & & K_{p+1} = \dots = K_n. \end{aligned} \quad (32)$$

For each patch $i \in I$ we denote by γ_{iJ} the flux from block J to patch i , and for each patch $j \in J$ we denote by γ_{jI} the flux from block I to patch j , as defined in Table 1. For each patch i we denote by T_i the sum of all migration rates γ_{ji} from patch i to another patch $j \neq i$ (i.e. the outgoing flux of patch i) minus the sum of the migration rates γ_{ik} from patch k to patch i , where k belongs to the same block as i . Hence, we have:

$$\begin{cases} \text{If } i \in I, & \text{then } T_i = \sum_{j \in J} \gamma_{ji} + \sum_{k \in I \setminus \{i\}} (\gamma_{ki} - \gamma_{ik}). \\ \text{If } j \in J, & \text{then } T_j = \sum_{i \in I} \gamma_{ij} + \sum_{k \in J \setminus \{j\}} (\gamma_{kj} - \gamma_{jk}). \end{cases} \quad (33)$$

We make the following assumption on the migration rates:

$$\begin{aligned} \gamma_{1J} = \dots = \gamma_{pJ}, & & \gamma_{(p+1)I} = \dots = \gamma_{nI} \\ T_1 = \dots = T_p, & & T_{p+1} = \dots = T_n \end{aligned} \quad (34)$$

where γ_{iJ} , for $i \in I$ and γ_{jI} , for $j \in J$ are defined in Table 1 and T_i are given by (33).

We have the following result:

Lemma 4.6. *Assume that the conditions (34) are satisfied, then for all $i \in I$ and $j \in I$ one has*

$$\gamma_{iJ} = \gamma_{IJ}/p, \quad \gamma_{jI} = \gamma_{JI}/q, \quad T_i = \gamma_{JI}/p, \quad T_j = \gamma_{IJ}/q. \quad (35)$$

where γ_{IJ} and γ_{JI} are defined in Table 1.

Proof. The result follows from $\sum_{i \in I} \gamma_{iJ} = \gamma_{IJ}$, $\sum_{i \in J} \gamma_{jI} = \gamma_{JI}$, $\sum_{i \in I} T_i = \gamma_{JI}$ and $\sum_{i \in J} T_j = \gamma_{IJ}$. \square

In the next theorem, we will show that, at the equilibrium, and under certain conditions relating to the migration rates, we can consider the n -patch model as a 2-patch model coupled by migration terms, which are not symmetric in general. Mathematically, we can state our main result as follows:

Table 1. Definitions and notations of fluxes.

Flux	Definition
$\gamma_{iJ} = \sum_{j \in J} \gamma_{ij}$	For $i \in I$, γ_{iJ} is the flux from block J to patch i , i.e. the sum of the migration rates γ_{ij} from patch $j \in J$ to patch i .
$\gamma_{jI} = \sum_{i \in I} \gamma_{ji}$	For $j \in J$, γ_{jI} is the flux from block I to patch j , i.e. the sum of the migration rates γ_{ji} from patch $i \in I$ to patch j .
$\gamma_{IJ} = \sum_{i \in I, j \in J} \gamma_{ij}$	γ_{IJ} is the flux from block J to block I , i.e. the sum of the migration rates γ_{ij} from patch $j \in J$, to patch $i \in I$.
$\gamma_{JI} = \sum_{i \in I, j \in J} \gamma_{ji}$	γ_{JI} is the flux from block I to block J , i.e. the sum of the migration rates γ_{ji} from patch $i \in I$, to patch $j \in J$.

Theorem 4.7. Assume that the conditions (32) and (34) are satisfied. Then the equilibrium of (9) is of the form

$$x_1 = x_1^*, \dots, x_p = x_1^*, \quad x_{p+1} = x_n^*, \dots, x_n = x_n^*$$

where (x_1^*, x_n^*) is the solution of the equations

$$\begin{cases} pr_1 x_1 \left(1 - \frac{x_1}{K_1}\right) + \beta (\gamma_{IJ} x_n - \gamma_{JI} x_1) = 0, \\ qr_n x_n \left(1 - \frac{x_n}{K_n}\right) + \beta (\gamma_{JI} x_1 - \gamma_{IJ} x_n) = 0, \end{cases} \tag{36}$$

that is to say, (x_1^*, x_n^*) is the equilibrium of a 2-patch model, with specific growth rates pr_1 and qr_n , carrying capacities K_1 and K_n and migration rates γ_{JI} from patch 1 to patch 2 and γ_{IJ} from patch 2 to patch 1.

Proof. Assume that the conditions (32) are satisfied. Then the equilibrium of (9) is the unique positive solution of the set of algebraic equations

$$\begin{cases} r_1 x_i \left(1 - \frac{x_i}{K_1}\right) + \beta \sum_{k=1, k \neq i}^n (\gamma_{ik} x_k - \gamma_{ki} x_i) = 0, & i = 1, \dots, p, \\ r_n x_j \left(1 - \frac{x_j}{K_n}\right) + \beta \sum_{k=1, k \neq j}^n (\gamma_{jk} x_k - \gamma_{kj} x_j) = 0, & j = p+1, \dots, n. \end{cases} \tag{37}$$

We consider the following set of algebraic equations obtained from (37) by replacing $x_i = x_1$ for $i = 1 \dots p$ and $x_i = x_n$ for $i = p+1 \dots n$:

$$\begin{cases} r_1 x_1 \left(1 - \frac{x_1}{K_1}\right) + \beta (\gamma_{iJ} x_n - T_i x_1) = 0, & i = 1, \dots, p, \\ r_n x_n \left(1 - \frac{x_n}{K_n}\right) + \beta (\gamma_{jI} x_1 - T_j x_n) = 0, & j = p+1, \dots, n. \end{cases} \tag{38}$$

Now, using the assumptions (34), together with the relations (35), we see that the system (38) is equivalent to the set of two algebraic equations:

$$\begin{cases} r_1 x_1 \left(1 - \frac{x_1}{K_1}\right) + \beta \left(\frac{\gamma_{IJ}}{p} x_n - \frac{\gamma_{JI}}{p} x_1\right) = 0, \\ r_n x_n \left(1 - \frac{x_n}{K_n}\right) + \beta \left(\frac{\gamma_{JI}}{q} x_1 - \frac{\gamma_{IJ}}{q} x_n\right) = 0. \end{cases} \quad (39)$$

We first notice that if $x_1 = x_1^*$, $x_n = x_n^*$ is a positive solution of (39) then $x_i = x_1^*$ for $i = 1, \dots, p$ and $x_j = x_n^*$ for $j = 1, \dots, n$ is a positive solution of (37). Let us prove that (39) has a unique solution (x_1^*, x_n^*) . Indeed, by multiplying the first equation by p and the second one by q , we deduce that (39) can be written in the form (36). \square

As a corollary of the previous theorem we obtain the following result which describes the total equilibrium population in the two blocks:

Corollary 4.8. *Assume that the conditions (32) and (34) are satisfied. Then the total equilibrium population $X_T^*(\beta) = px_1^*(\beta) + qx_n^*(\beta)$ of (9) behaves like the total equilibrium population of the 2-patch model*

$$\begin{cases} \frac{dy_1}{dt} = r_1 y_1 \left(1 - \frac{y_1}{pK_1}\right) + \beta (\gamma_{12} y_n - \gamma_{21} y_1), \\ \frac{dy_n}{dt} = r_n y_n \left(1 - \frac{y_n}{qK_n}\right) + \beta (\gamma_{21} y_1 - \gamma_{12} y_n), \end{cases} \quad (40)$$

with specific growth rates r_1 and r_n , carrying capacities pK_1 and qK_n , and migration rates $\gamma_{21} = \frac{\gamma_{JI}}{p}$, $\gamma_{12} = \frac{\gamma_{IJ}}{q}$.

Proof. From Theorem 4.7, we see that (x_1^*, x_n^*) is the positive solution of (36). Hence, $(y_1^* = px_1^*, y_n^* = qx_n^*)$ is the solution of the set of equations

$$\begin{cases} r_1 y_1 \left(1 - \frac{y_1}{pK_1}\right) + \beta \left(\frac{\gamma_{IJ}}{q} y_n - \frac{\gamma_{JI}}{p} y_1\right) = 0, \\ r_n y_n \left(1 - \frac{y_n}{qK_n}\right) + \beta \left(\frac{\gamma_{JI}}{p} y_1 - \frac{\gamma_{IJ}}{q} y_n\right) = 0, \end{cases} \quad (41)$$

obtained from (36) by changing variables to $y_1 = px_1$, $y_n = qx_n$. The system (41) has a unique positive solution which is the equilibrium point of the 2-patch model (40). \square

We can describe when, under the conditions (32) and (34), the migration pattern is beneficial or detrimental in Model (9).

We consider the regions in the set of parameters γ_{IJ} and γ_{JI} , denoted \mathcal{J}_0 , \mathcal{J}_1 and \mathcal{J}_2 ,

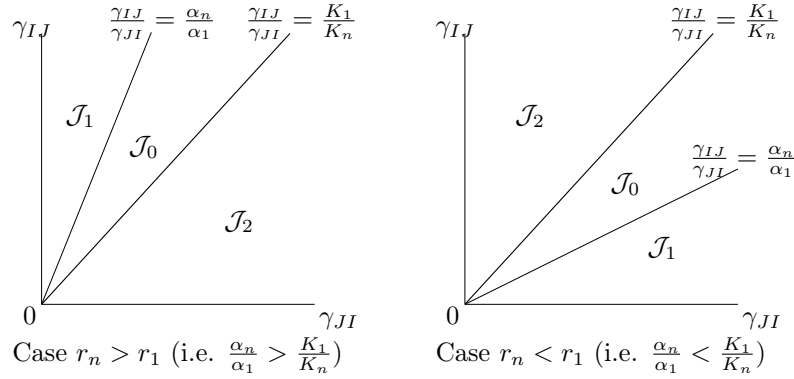


Figure 1. Qualitative properties of Model (9) under the conditions (32) and (34). In \mathcal{J}_0 , fragmentation benefits the total equilibrium population. This effect is detrimental in \mathcal{J}_2 . In \mathcal{J}_1 , the effect is beneficial for $\beta < \beta_0$ and detrimental for $\beta > \beta_0$.

depicted in Figure 1 and defined by:

$$\begin{aligned}
 &\text{If } r_n > r_1 \text{ then } \begin{cases} \mathcal{J}_1 = \left\{ (\gamma_{JI}, \gamma_{IJ}) : \frac{\gamma_{IJ}}{\gamma_{JI}} > \frac{\alpha_n}{\alpha_1} \right\} \\ \mathcal{J}_0 = \left\{ (\gamma_{JI}, \gamma_{IJ}) : \frac{\alpha_n}{\alpha_1} \geq \frac{\gamma_{IJ}}{\gamma_{JI}} > \frac{K_1}{K_n} \right\} \\ \mathcal{J}_2 = \left\{ (\gamma_{JI}, \gamma_{IJ}) : \frac{K_1}{K_n} > \frac{\gamma_{IJ}}{\gamma_{JI}} \right\} \end{cases} \\
 &\text{If } r_n < r_1 \text{ then } \begin{cases} \mathcal{J}_1 = \left\{ (\gamma_{JI}, \gamma_{IJ}) : \frac{\gamma_{IJ}}{\gamma_{JI}} < \frac{\alpha_n}{\alpha_1} \right\} \\ \mathcal{J}_0 = \left\{ (\gamma_{JI}, \gamma_{IJ}) : \frac{\alpha_n}{\alpha_1} \leq \frac{\gamma_{IJ}}{\gamma_{JI}} < \frac{K_1}{K_n} \right\} \\ \mathcal{J}_2 = \left\{ (\gamma_{JI}, \gamma_{IJ}) : \frac{K_1}{K_n} < \frac{\gamma_{IJ}}{\gamma_{JI}} \right\} \end{cases} \end{aligned} \tag{42}$$

where $\alpha_1 = r_1/K_1$ and $\alpha_n = r_n/K_n$.

Proposition 4.9. Assume that the conditions (32) and (34) are satisfied. Then the total equilibrium population $X_T^*(\beta) = px_1^*(\beta) + qx_n^*(\beta)$ of (9) satisfies the following properties

1. If $r_1 = r_n$ then $X_T^*(\beta) < pK_1 + qK_n$ for all $\beta > 0$.
2. If $r_n \neq r_1$, let $\mathcal{J}_0, \mathcal{J}_1$ and \mathcal{J}_2 , be defined by (42). Then we have:
 - if $(\gamma_{JI}, \gamma_{IJ}) \in \mathcal{J}_0$ then $X_T^*(\beta) > pK_1 + qK_n$ for any $\beta > 0$,
 - if $(\gamma_{JI}, \gamma_{IJ}) \in \mathcal{J}_1$ then $X_T^*(\beta) > pK_1 + qK_n$ for $0 < \beta < \beta_0$ and $X_T^*(\beta) < pK_1 + qK_n$ for $\beta > \beta_0$, where

$$\beta_0 = \frac{r_n - r_1}{\frac{\gamma_{IJ}}{\alpha_n} - \frac{\gamma_{JI}}{\alpha_1}} \frac{1}{\frac{\alpha_1}{p} + \frac{\alpha_n}{q}}.$$

- If $(\gamma_{JI}, \gamma_{IJ}) \in \mathcal{J}_2$ then $X_T^*(\beta) < pK_1 + qK_n$ for any $\beta > 0$.
- If $\frac{\gamma_{IJ}}{\gamma_{JI}} = \frac{K_1}{K_n}$, then $X_T^*(\beta) = pK_1 + qK_n$ for all $\beta \geq 0$.

Proof. This is a consequence of Proposition A.1 and Corollary 4.8. \square

Let us explain the result of Proposition 4.9 in the particular case where $p = n - 1$. In this case, the condition (34) becomes

$$\gamma_{1n} = \dots = \gamma_{n-1,n} \quad \text{and} \quad T_1 = \dots = T_{n-1}, \quad (43)$$

where $T_i = \gamma_{ni} + \sum_{k \neq i} (\gamma_{ki} - \gamma_{ik})$. Therefore, if the matrix Γ is symmetric, the conditions (43) are equivalent to the conditions $\gamma_{n1} = \dots = \gamma_{n,n-1}$, which mean that the fluxes of migration between the n -th patch and all $n - 1$ identical patches are equal. Hence, Proposition 4.9, showing that the n -patch model behaves like a 2-patch model, is the same as [12, Proposition 3.4], where the model (9) was considered with Γ symmetric, $n - 1$ patches are identical and the fluxes of migration between the n -th patch and all these $n - 1$ identical patches are equal. Thus Proposition 4.9 generalizes Proposition 3.4 of [12], to asymmetric dispersal and for any two identical blocks, provided that the conditions (34) are satisfied.

5. Links between SIS and logistic patch models

5.1. The SIS patch model

In [15], Gao studied the following SIS patch model in an environment of n patches connected by human migration:

$$\begin{cases} \frac{dS_i}{dt} = -\beta_i \frac{S_i I_i}{N_i} + \gamma_i I_i + \varepsilon \sum_{j=1}^n \gamma_{ij} S_j, & i = 1, \dots, n, \\ \frac{dI_i}{dt} = \beta_i \frac{S_i I_i}{N_i} - \gamma_i I_i + \varepsilon \sum_{j=1}^n \gamma_{ij} I_j, & i = 1, \dots, n, \end{cases} \quad (44)$$

where S_i and I_i are the number of susceptible and infected, in patch i , respectively; $N_i = S_i + I_i$ denotes the total population in patch i . The parameters β_i and γ_i are positive transmission and recovery rates, respectively. The matrix $\Gamma = (\gamma_{ij})$ satisfies (3) and describes the movement between patches. The coefficient ε quantifies the diffusion, as our β in (9).

Using the variables $N_i, I_i, i = 1, \dots, n$, the system (44) has a cascade structure

$$\frac{dN_i}{dt} = \varepsilon \sum_{j=1}^n \gamma_{ij} N_j, \quad i = 1, \dots, n, \quad (45)$$

$$\frac{dI_i}{dt} = \beta_i \frac{(N_i - I_i) I_i}{N_i} - \gamma_i I_i + \varepsilon \sum_{j=1}^n \gamma_{ij} I_j, \quad i = 1, \dots, n, \quad (46)$$

Therefore the infected populations I_i are the solutions of the non-autonomous system of differential equations

$$\frac{dI_i}{dt} = \beta_i I_i \left(1 - \frac{I_i}{N_i(t)} \right) - \gamma_i I_i + \varepsilon \sum_{j=1}^n \gamma_{ij} I_j, \quad i = 1, \dots, n, \quad (47)$$

where the total populations $N_i(t)$ are the solutions of the system (45). Hence, the autonomous $2n$ -dimensional system (44), is equivalent to the family of n -dimensional non-autonomous systems (47), indexed by the solutions $N_i(t)$ of (45). Note that since the γ_{ij} verify the property (3), the total population is constant: $\sum_{i=1}^n N_i(t) = N$, where $N := \sum_{i=1}^n (S_i(0) + I_i(0))$. If the matrix $\Gamma = (\gamma_{ij})$ is irreducible, then $N_i(t)$, the total population in patch i , converges towards the limit

$$\lim_{t \rightarrow +\infty} N_i(t) = N_i^* \quad \text{where } N_i^* := \frac{N}{\sum_i \delta_i} \delta_i, \quad i = 1, \dots, n, \quad (48)$$

where $\delta = (\delta_1, \dots, \delta_n)^T$ is a positive vector which generates the vector space $\ker \Gamma$. Therefore, (47) is an asymptotically autonomous system, whose limit system is obtained by replacing $N_i(t)$ in (47), by their limits N_i^* , given by (48):

$$\frac{dI_i}{dt} = \beta_i I_i \left(1 - \frac{I_i}{N_i^*} \right) - \gamma_i I_i + \varepsilon \sum_{j=1}^n \gamma_{ij} I_j, \quad i = 1, \dots, n. \quad (49)$$

The main problem for (44) is to determine the condition under which the disease free equilibrium, corresponding to the equilibrium $I = 0$ of (49), is GAS, or the endemic equilibrium, corresponding to the positive equilibrium of (49), is GAS. It is known, see [15, Theorem 2.1], that the disease free equilibrium is GAS if $\mathcal{R}_0 \leq 1$, and there exists a unique endemic equilibrium, which is GAS, if $\mathcal{R}_0 > 1$. Here \mathcal{R}_0 is the basic reproduction number of the model (44), defined as:

$$\mathcal{R}_0 = \rho(FV^{-1}) \quad \text{where } F = \text{diag}(\beta_1, \dots, \beta_n) \text{ and } V = \text{diag}(\gamma_1, \dots, \gamma_n) - \varepsilon \Gamma.$$

A reference work on the basic reproduction number for metapopulations is Arino [3], whereas Castillo-Garsow and Castillo-Chavez [7] and van den Driessche and Watmough [29] give a more general account of the subject.

5.2. Comparisons between the results on (9) and the results on (49)

Gao [15] gave many interesting results on the effect of population dispersal on total infection size. Our aim is to discuss some of the links between his results and the results of the present paper. We focus on two results on the total infection size $T_n(\varepsilon) = \sum_{i=1}^n I_i^*(\varepsilon)$, where $(I_1^*(\varepsilon), \dots, I_n^*(\varepsilon))$ is the positive equilibrium of (49). We consider the results of Gao [15] on $T_n(+\infty)$ and $T_n'(0)$.

Proposition 5.1 ([15, Theorem 3.3], [15, Theorem 3.5]). *If $\mathcal{R}_0(+\infty) > 1$, then*

$$T_n(+\infty) = \left(1 - \frac{1}{\mathcal{R}_0(+\infty)} \right) N, \quad \text{with } \mathcal{R}_0(+\infty) = \frac{\sum_i \beta_i \delta_i}{\sum_i \gamma_i \delta_i}. \quad (50)$$

If $\beta_i \neq \gamma_i$ for all i , then

$$T_n'(0) = \sum_i \left(\frac{1}{|\beta_i - \gamma_i|} \sum_j \gamma_{ij} I_j^*(0) \right). \quad (51)$$

It is worth noting that the formulas (50) and (51) involve the system (49). An important property of this system is given in the following remark.

Remark 5.2. Let $N^* = (N_1^*, \dots, N_n^*)^T$ be the vector of the carrying capacities in the system (49). One has $N^* \in \ker \Gamma$, as shown by (48).

Our aim is to compare the results given by the formulas (50) and (51) when $\gamma_i \rightarrow 0$, to our results, for the system

$$\frac{dx_i}{dt} = \beta_i x_i \left(1 - \frac{x_i}{N_i^*}\right) + \varepsilon \sum_{j=1}^n \gamma_{ij} x_j, \quad i = 1, \dots, n. \quad (52)$$

Note that the system (49) reduces to (52) when $\gamma_i = 0$ for all i . More precisely we show that, as $\gamma_i \rightarrow 0$, the formulas (50) and (51) are the same as the results predicted by Proposition 3.4 and Proposition 4.4.

Proposition 5.3. *Let $T_n(\varepsilon)$ be the total infection size of (49). Let $X_T^*(\varepsilon)$ be the total population size of (52). One has*

$$\lim_{\max_i \{\gamma_i\} \rightarrow 0} T_n(+\infty) = X_T^*(+\infty) = N, \quad \lim_{\max_i \{\gamma_i\} \rightarrow 0} T_n'(0) = \frac{dX_T^*}{d\varepsilon}(0) = 0. \quad (53)$$

Proof. When $\gamma_i \rightarrow 0$ for all i , one has $\mathcal{R}_0(+\infty) \rightarrow +\infty$ and $I_i^*(0) \rightarrow N_i^*$. Therefore, from (50) and (51) it is deduced that

$$T_n(+\infty) \rightarrow N, \quad T_n'(0) \rightarrow \sum_i \frac{1}{\beta_i} \sum_j \gamma_{ij} N_j^* = 0. \quad (54)$$

Using the property $N^* \in \ker \Gamma$, from Proposition 3.4 and Proposition 4.4, it is deduced that:

$$X_T^*(+\infty) = N, \quad \frac{dX_T^*}{d\varepsilon}(0) = 0. \quad (55)$$

From (54) and (55) we deduce (53). \square

Actually as shown in Proposition 4.5, we have the stronger result $X_T^*(\beta) = N$ for all $\beta \geq 0$. But our aim here was only the comparison between (54) and (55).

As shown in Proposition 5.2, the results of Gao [15] on the logistic patch model (49) yield results on the logistic patch model (52) by taking the limit $\gamma_i \rightarrow 0$. However, the scope of this approach is weakened by the fact that it only applies to the logistic model (52), for which the vector of carrying capacities satisfies $N^* \in \ker \Gamma$, see Remark 5.2. But this property is not true in general for our system (9), where the condition $K \in \ker \Gamma$ does not hold in general.

Our aim in this section is to show that any logistic patch model (9), without the condition $K \in \ker \Gamma$, can be written in the form (49), with the condition $N^* \in \ker \Gamma$. Indeed we have the following result:

Lemma 5.4. Consider $r_i > 0$, $K_i > 0$ and Γ as in the system (9). Let $\delta_i > 0$ be such that $(\delta_1, \dots, \delta_n)^T \in \ker \Gamma$. Let N be such that $N > \frac{\sum_i \delta_i}{\delta_i} K_i$ for $i = 1, \dots, n$. Let N_i^* defined by (48). Let $\beta_i = \frac{r_i}{K_i} N_i^*$ and $\gamma_i = \beta_i - r_i$. Then one has

$$r_i x_i (1 - x_i / K_i) = \beta_i x_i (1 - x_i / N_i^*) - \gamma_i x_i, \quad \text{for } i = 1, \dots, n \tag{56}$$

Proof. The conditions (56) are satisfied if and only if $r_i = \beta_i - \gamma_i$ and $r_i / K_i = \beta_i / N_i^*$. Therefore

$$\begin{cases} \beta_i = N_i^* \frac{r_i}{K_i} = N_i^* \alpha_i, \\ \gamma_i = \beta_i - r_i = (N_i^* - K_i) \alpha_i. \end{cases} \tag{57}$$

To ensure that $\gamma_i > 0$ for all i , just choose N in (48) such that $N_i^* > K_i$ for $i = 1, \dots, n$, that is to say, $N > \frac{\sum_i \delta_i}{\delta_i} K_i$. \square

Remark 5.5. According to the change of parameters (57), the logistic patch model (9) can be written in the form of Gao (49), i.e. with the property that $N^* \in \ker \Gamma$. For the perfect mixing case, the formula (50) and our formula (13) are the same. Indeed replacing β_i and γ_i by (57) in (50), and using (48), we get:

$$\left(1 - \frac{1}{\mathcal{R}_0(+\infty)}\right) N = \left(1 - \frac{\sum_i (N_i^* - K_i) \alpha_i}{\sum_i N_i^* \alpha_i}\right) N = \sum_i \delta_i \frac{\sum_i r_i \delta_i}{\sum_i \alpha_i \delta_i^2}.$$

For the derivative, the formula (51) and our formula (28) are the same. Indeed, if we replace β_i and γ_i by (57), in (51), we get:

$$I_j^*(0) = \frac{\beta_j - \gamma_j}{\beta_j} N_j^* = \frac{r_j}{N_j^* \alpha_j} N_j^* = K_j.$$

Therefore

$$\sum_i \left(\frac{1}{|\beta_i - \gamma_i|} \sum_j \gamma_{ij} I_j^*(0) \right) = \sum_i \left(\frac{1}{r_i} \sum_j \gamma_{ij} K_j \right).$$

The theory of asymptotically autonomous systems answers the question “under which conditions do the solutions of the original $2n$ -dimensional system (44) have the same asymptotic behavior as those of the n -dimensional limit system (49) ?”. For details and further reading on the theory of asymptotically autonomous systems, the reader is referred to Markus [23] and Thieme [26, 27]. For applications of this theory to epidemic models, see Castillo-Chavez and Thieme [6].

Hence, it is important to know whether or not some of the results of Gao [15] on the SIS model (44) can be deduced from our results on the logistic model (9). It is worth noting that the discussion in this section shows that our results on the logistic patch model imply results on the model (49) and hence, results on the original model $2n$ -dimensional system (44). However, it is needed that $\beta_i > \gamma_i$ for $i = 1, \dots, n$. Indeed, according to (57), one has $r_i = \beta_i - \gamma_i > 0$. On the other hand, the condition $\beta_i > \gamma_i$ is not required in all patches of the system (44). Another challenging problem is the study of the model (49), in the general case where $N^* = (N_1^*, \dots, N_n^*)^T$ is not necessarily in the kernel of Γ .

6. Three-patch model

In this section, we consider the model of three patches coupled by asymmetrical terms of migrations. Under the irreducibility hypothesis on the matrix Γ , there are five possible cases, modulo permutation of the three patches, see Figures 2 and 3.

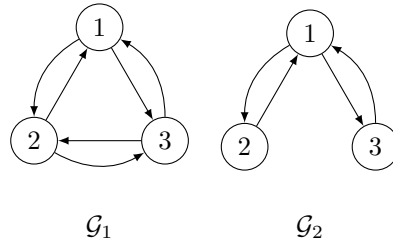


Figure 2. The two graphs \mathcal{G}_1 and \mathcal{G}_2 for which the migration matrix may be symmetric, if $\gamma_{ij} = \gamma_{ji}$.

The connectivity matrices associated to the graphs \mathcal{G}_1 and \mathcal{G}_2 are given by

$$\Gamma_0^{(1)} = \begin{bmatrix} 0 & \gamma_{12} & \gamma_{13} \\ \gamma_{21} & 0 & \gamma_{23} \\ \gamma_{31} & \gamma_{32} & 0 \end{bmatrix}, \quad \text{and} \quad \Gamma_0^{(2)} = \begin{bmatrix} 0 & \gamma_{12} & \gamma_{13} \\ \gamma_{21} & 0 & 0 \\ \gamma_{31} & 0 & 0 \end{bmatrix}.$$

For the remaining cases, the graphs $\mathcal{G}_3, \mathcal{G}_4$ and \mathcal{G}_5 , cannot be symmetrical:

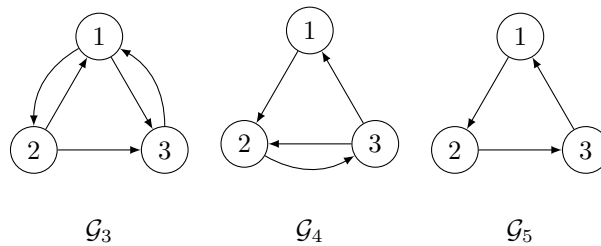


Figure 3. The three graphs $\mathcal{G}_3, \mathcal{G}_4$ and \mathcal{G}_5 for which the migration matrix cannot be symmetric.

The associated connectivity matrices are given by

$$\Gamma_0^{(3)} = \begin{bmatrix} 0 & \gamma_{12} & \gamma_{13} \\ \gamma_{21} & 0 & 0 \\ \gamma_{31} & \gamma_{32} & 0 \end{bmatrix}, \quad \Gamma_0^{(4)} = \begin{bmatrix} 0 & 0 & \gamma_{13} \\ \gamma_{21} & 0 & \gamma_{23} \\ 0 & \gamma_{32} & 0 \end{bmatrix}, \quad \Gamma_0^{(5)} = \begin{bmatrix} 0 & 0 & \gamma_{13} \\ \gamma_{21} & 0 & 0 \\ 0 & \gamma_{32} & 0 \end{bmatrix}.$$

In Table 2, we give the formula of perfect mixing $X_T^*(+\infty)$ for each of the five cases.

In the numerical simulations, we show that we can have new behaviors of $X_T^*(\beta)$. In the case $n = 2$, it was shown in [1, 2] that there exists at most one positive value of β such that $X_T^*(\beta) = K_1 + K_2$. In [12], in the case $n = 3$ and Γ is symmetric, we gave numerical values for the parameters such that there exists two positive values of β such that $X_T^*(\beta) = K_1 + K_2 + K_3$, and we were not able to find more than two values. The novelty when Γ is not symmetric is that we can find examples with three positive values.

Table 2. The generator δ of $\ker \Gamma$, for the five cases. The perfect mixing abundance $X_T^*(+\infty)$ is computed with Eq. (13).

Graphs	The formula of perfect mixing $X_T^*(+\infty)$
\mathcal{G}_1	The coefficients δ_i are given by the Equation (12)
\mathcal{G}_2	$\delta_1 = \gamma_{12}\gamma_{13}, \delta_2 = \gamma_{21}\gamma_{13}, \delta_3 = \gamma_{12}\gamma_{31},$
\mathcal{G}_3	$\delta_1 = \gamma_{12}\gamma_{13} + \gamma_{32}\gamma_{13}, \delta_2 = \gamma_{21}\gamma_{13}, \delta_3 = \gamma_{21}\gamma_{32} + \gamma_{31}\gamma_{12} + \gamma_{31}\gamma_{32},$
\mathcal{G}_4	$\delta_1 = \gamma_{32}\gamma_{13}, \delta_2 = \gamma_{21}\gamma_{13} + \gamma_{21}\gamma_{23} + \gamma_{31}\gamma_{23}, \delta_3 = \gamma_{21}\gamma_{32}.$
\mathcal{G}_5	$\delta_1 = \gamma_{32}\gamma_{13}, \delta_2 = \gamma_{21}\gamma_{13}, \delta_3 = \gamma_{21}\gamma_{32}.$

Indeed, we may have the following situation : $\frac{dX_T^*}{d\beta}(0) > 0$ and $X_T^*(+\infty) < K_1 + K_2 + K_3$, and there exist three values $0 < \beta_1 < \beta_2 < \beta_3$ for which we have

$$X_T^*(\beta) \begin{cases} > K_1 + K_2 + K_3 & \text{for } \beta \in]0, \beta_1[\cup]\beta_2, \beta_3[, \\ < K_1 + K_2 + K_3 & \text{for } \beta \in]\beta_1, \beta_2[\cup]\beta_3, +\infty[. \end{cases} \quad (58)$$

The same situation holds for each of the five graphs $\mathcal{G}_1, \mathcal{G}_2, \mathcal{G}_3, \mathcal{G}_4$ and \mathcal{G}_5 , i.e, there exist three values $0 < \beta_1 < \beta_2 < \beta_3$ for which (58) hold. See Figures 4, (for the graph \mathcal{G}_1), 5, (for the graph \mathcal{G}_2), 6-a, (for the graph \mathcal{G}_3), 6-b, (for the graph \mathcal{G}_4), and 6-c, (for the graph \mathcal{G}_5).

Table 3. The numerical values of the parameters for the logistic growth function and migration coefficients of the model (9), with $n = 3$, used in Figures 4,5,6-a,6-b and Figure 6-c. For all figures we have $(r_1, r_2, r_3, K_1, K_2, K_3) = (4, 0.7, 0.6, 5, 1, 4)$. The perfect mixing abundance $X_T^*(+\infty)$ is computed with Eq. (13) and the derivative of the total equilibrium population at $\beta = 0$ is computed with Eq. (28).

Figure	γ_{21}	γ_{12}	γ_{31}	γ_{13}	γ_{32}	γ_{23}	$\frac{dX_T^*}{d\beta}(0)$	$X_T^*(+\infty)$
4	0.15	3	0.2	0.04	11	0.1	1.06	9.21
5	14.9	10	0.2	0.04	0	0	77.20	9.86
6-a	1.44	0.01	0.2	0.04	1	0	3.11	8.93
6-b	1.52	0	0	1	1	0.002	3.52	8.72
6-c	1.51	0	0	1	1	0	3.46	8.75

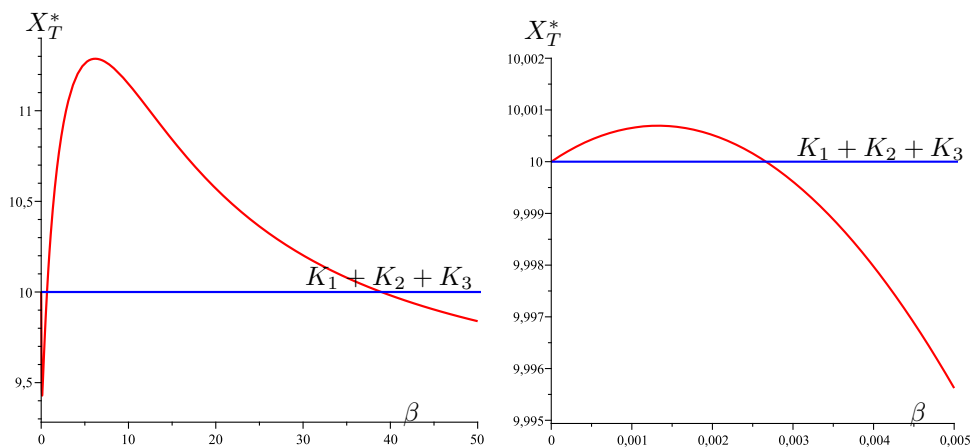


Figure 4. Total equilibrium population X_T^* of the system (9) ($n = 3$) as a function of the migration rate β . The figure on the right is a zoom, near the origin, of the figure on the left. The parameter values are given in Table 3.

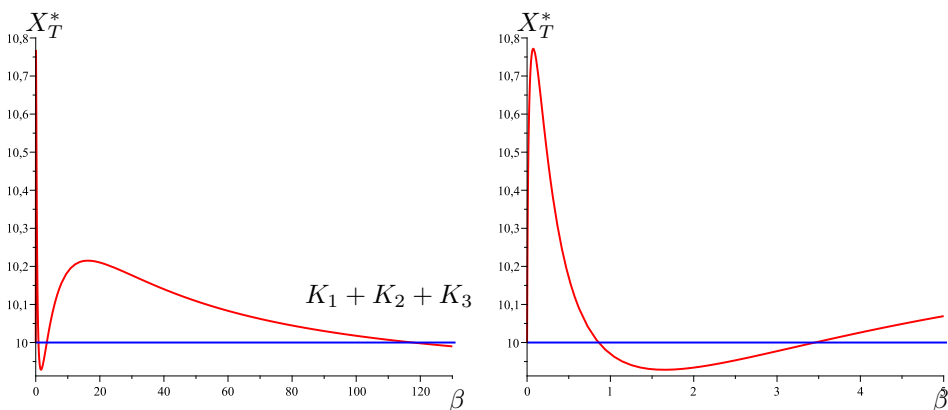


Figure 5. Total equilibrium population X_T^* of the system (9) ($n = 3$) as a function of the migration rate β . The figure on the right is a zoom, near the origin, of the figure on the left. The parameter values are given in Table 3.

7. Conclusion

The aim of this paper is to generalize, to a multi-patch model with asymmetric dispersal, the results obtained in [12] for a multi-patch model with symmetric dispersal.

In Section 3 we considered the particular case of perfect mixing, when the migration rate goes to infinity, that is, individuals may travel freely between patches. As in [12], we compute the total equilibrium population in that case, and, by perturbation arguments, we proved that the dynamics in this ideal case provides a good approximation to the case when the migration rate is large. Our results generalize those of [2] (asymmetric migration matrix, only two patches), [10] (arbitrarily many patches, but the migration matrix is symmetric and zero outside the corners and the three main diagonals), and [12] (arbitrarily many patches; arbitrary, but symmetric, migration).

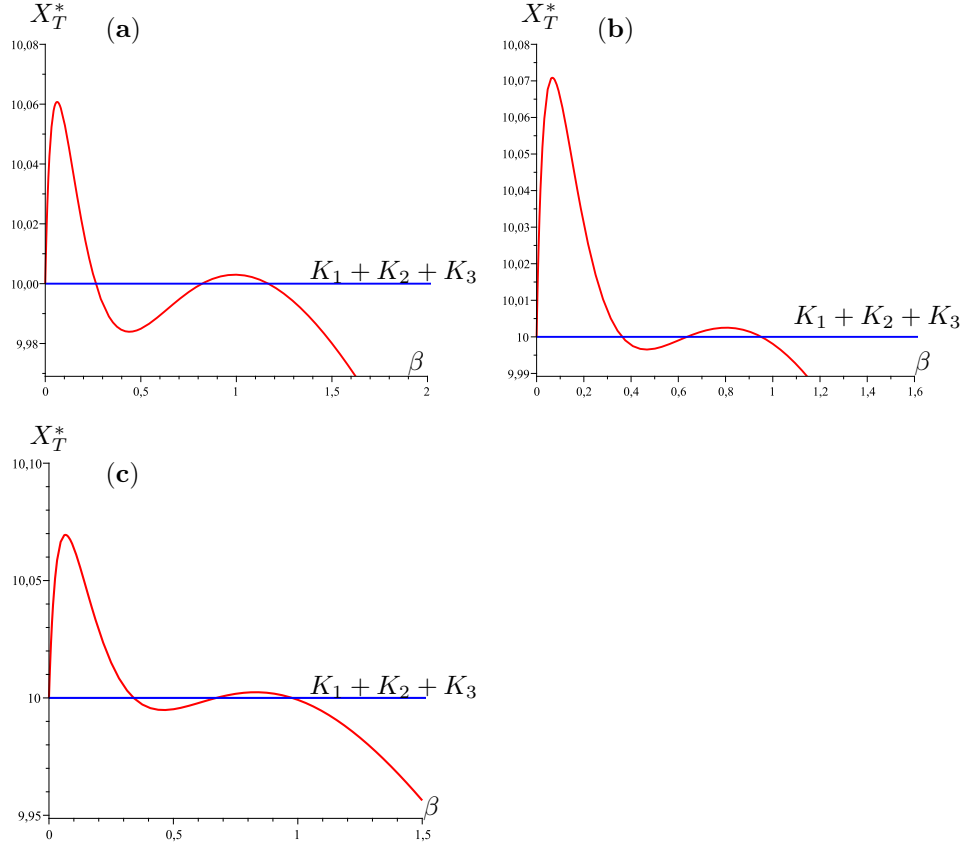


Figure 6. Total equilibrium population X_T^* of the system (9) ($n = 3$) as a function of the migration rate β . The parameter values are given in Table 3.

In Section 4 we considered the equation

$$\text{total equilibrium population} = \text{sum of the carrying capacities of the patches.} \quad (59)$$

We gave a complete solution in the case when the n patches are partitioned into two blocks of identical patches. Our results mirror those of [2], which deals with the two-patch case. Specifically, Equation (59) admits at most one non-trivial solution.

In Section 5, we consider a SIS patch model and we give the links with the logistic model.

In Section 6 we give numerical values for the dispersion parameters such that Equation (59) has at least three non-trivial solutions. In [12] we proved that for three patches and symmetric dispersal, there may be at least two solutions. A mathematical proof that, when $n=3$, Equation (59) has at most three solutions, would certainly be desirable, and could spur further work. Upper bounds for arbitrarily many patches would also be interesting.

Appendix

A. The 2-patch asymmetric model

We consider the 2-patch logistic equation with asymmetric migrations. We denote by γ_{12} the migration rate from patch 2 to patch 1, and γ_{21} from patch 1 to patch 2. The model is written:

$$\begin{cases} \frac{dx_1}{dt} = r_1 x_1 \left(1 - \frac{x_1}{L_1}\right) + \beta (\gamma_{12} x_2 - \gamma_{21} x_1), \\ \frac{dx_2}{dt} = r_2 x_2 \left(1 - \frac{x_2}{L_2}\right) + \beta (\gamma_{21} x_1 - \gamma_{12} x_2). \end{cases} \quad (60)$$

Note that the system (60) is studied in [1, 8, 13, 14, 18] in the case where the migration rates satisfy $\gamma_{21} = \gamma_{12}$, and in [2] for general migration rates. This system admits a unique equilibrium which is GAS. We denote by $E^*(\beta) = (x_1^*(\beta), x_2^*(\beta))$ this equilibrium and by $X_T^*(\beta)$ the sum of $x_i^*(\beta)$.

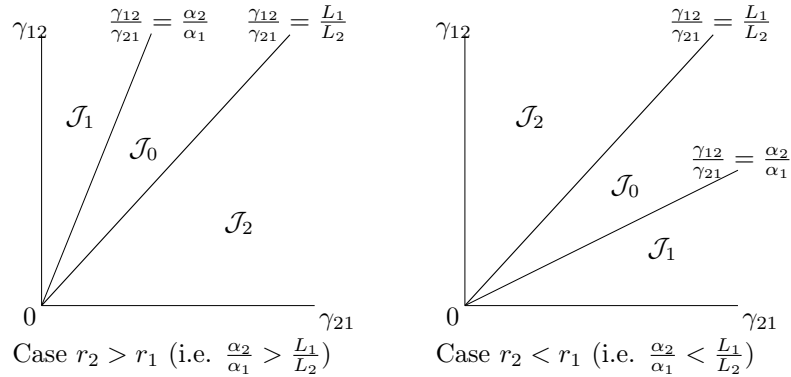


Figure 7. Qualitative properties of model (60). In \mathcal{J}_0 , patchiness has a beneficial effect on total equilibrium population. This effect is detrimental in \mathcal{J}_2 . In \mathcal{J}_1 , the effect is beneficial for $\beta < \beta_0$ and detrimental for $\beta > \beta_0$. In the figure $\alpha_1 = r_1/L_1$ and $\alpha_2 = r_2/L_2$.

We consider the regions in the set of the parameters γ_{21} and γ_{12} , denoted \mathcal{J}_0 , \mathcal{J}_1 and \mathcal{J}_2 , depicted in Fig. 7 and defined by:

$$\left\{ \begin{array}{l} \text{If } r_2 > r_1 \text{ then} \\ \text{If } r_2 < r_1 \text{ then} \end{array} \right. \left\{ \begin{array}{l} \mathcal{J}_1 = \left\{ (\gamma_{21}, \gamma_{12}) : \frac{\gamma_{12}}{\gamma_{21}} > \frac{\alpha_2}{\alpha_1} \right\} \\ \mathcal{J}_0 = \left\{ (\gamma_{21}, \gamma_{12}) : \frac{\alpha_2}{\alpha_1} \geq \frac{\gamma_{12}}{\gamma_{21}} > \frac{L_1}{L_2} \right\} \\ \mathcal{J}_2 = \left\{ (\gamma_{21}, \gamma_{12}) : \frac{L_1}{L_2} > \frac{\gamma_{12}}{\gamma_{21}} \right\} \\ \mathcal{J}_1 = \left\{ (\gamma_{21}, \gamma_{12}) : \frac{\gamma_{12}}{\gamma_{21}} < \frac{\alpha_2}{\alpha_1} \right\} \\ \mathcal{J}_0 = \left\{ (\gamma_{21}, \gamma_{12}) : \frac{\alpha_2}{\alpha_1} \leq \frac{\gamma_{12}}{\gamma_{21}} < \frac{L_1}{L_2} \right\} \\ \mathcal{J}_2 = \left\{ (\gamma_{21}, \gamma_{12}) : \frac{L_1}{L_2} < \frac{\gamma_{12}}{\gamma_{21}} \right\} \end{array} \right. \quad (61)$$

We have the following result which gives the conditions for which patchiness is beneficial or detrimental in model (60).

Proposition A.1. *The total equilibrium population of (60) satisfies the following properties*

1. If $r_1 = r_2$ then $X_T^*(\beta) \leq L_1 + L_2$ for all $\beta \geq 0$.
2. If $r_2 \neq r_1$, let $\mathcal{J}_0, \mathcal{J}_1$ and \mathcal{J}_2 , be defined by (61). Then we have:
 - If $(\gamma_{21}, \gamma_{12}) \in \mathcal{J}_0$ then $X_T^*(\beta) > L_1 + L_2$ for any $\beta > 0$.
 - If $(\gamma_{21}, \gamma_{12}) \in \mathcal{J}_1$ then $X_T^*(\beta) > L_1 + L_2$ for $0 < \beta < \beta_0$ and $X_T^*(\beta) < L_1 + L_2$ for $\beta > \beta_0$, where

$$\beta_0 = \frac{r_2 - r_1}{\frac{\gamma_{12}}{\alpha_2} - \frac{\gamma_{21}}{\alpha_1}} \frac{1}{\alpha_1 + \alpha_2}.$$
 - If $(\gamma_{21}, \gamma_{12}) \in \mathcal{J}_2$ then $X_T^*(\beta) < L_1 + L_2$ for any $\beta > 0$.
 - If $\frac{\gamma_{12}}{\gamma_{21}} = \frac{L_1}{L_2}$, then $x_1^*(\beta) = L_1$ and $x_2^*(\beta) = L_2$ for all $\beta \geq 0$. Therefore $X_T^*(\beta) = L_1 + L_2$ for all $\beta \geq 0$.

Proof. This result was established by Arditi et al. [2]. Part (1) is Proposition 1 of [2]. The first three items of part (2) are Proposition 2 of [2]. For the last item of part (2), see the last paragraph in page 12 of [2]. The explicit expression of β_0 was not given in [2], however, it is easy to deduce it from the formulas given in [2]. \square

B. Some useful results

We begin with a

Lemma B.1. *The matrix \mathcal{L} defined by (21) is stable, that is to say, all its eigenvalues have negative real part.*

Proof. We consider the two matrices

$$G := \begin{bmatrix} L - U & V \\ 0 & \dots & 0 & 0 \end{bmatrix}, \quad P := \begin{bmatrix} I & 0 \\ 1 & \dots & 1 & 1 \end{bmatrix},$$

where L, V , and U are defined right after (21). We prove that the two matrices Γ and G are conjugate by the matrix P , that is to say $P^{-1}GP = \Gamma$.

The inverse of matrix P is given by

$$P^{-1} = \begin{bmatrix} I & 0 \\ -1 & \dots & -1 & 1 \end{bmatrix}.$$

We have

$$P^{-1}GP = \begin{bmatrix} L & V \\ \gamma_{n1} & \dots & \gamma_{nn-1} & -\sum_{j=1, j \neq 1}^n \gamma_{jn} \end{bmatrix} = \Gamma.$$

Two conjugate matrices have the same eigenvalues. As the matrix G is block-triangular, its eigenvalues are zero and the eigenvalues of the matrix $L - U$. Therefore, since 0 is a simple eigenvalue of the matrix Γ , the eigenvalues of the matrix $L - U$ are the eigenvalues of the matrix Γ except 0. By Lemma 3.1 all non-zero eigenvalues of Γ have negative real part. \square

Lemma B.2. *Let $(u_n)_{n \geq 1}$, $(v_n)_{n \geq 1}$ and $(w_n)_{n \geq 1}$ be three real and non-negative sequences. Then,*

1. *if $(u_n)_{n \geq 1}$ and $(v_n)_{n \geq 1}$ are both non-increasing, or both non-decreasing, then we have, for all $N \geq 1$,*

$$\left(\sum_{n=1}^N w_n \right) \left(\sum_{n=1}^N w_n u_n v_n \right) \geq \left(\sum_{n=1}^N w_n u_n \right) \left(\sum_{n=1}^N w_n v_n \right), \quad (62)$$

2. *if $(u_n)_{n \geq 1}$ is non-decreasing and $(v_n)_{n \geq 1}$ is non-increasing, or if $(u_n)_{n \geq 1}$ is non-increasing and $(v_n)_{n \geq 1}$ is non-decreasing, then, we have, for all $N \geq 1$,*

$$\left(\sum_{n=1}^N w_n \right) \left(\sum_{n=1}^N w_n u_n v_n \right) \leq \left(\sum_{n=1}^N w_n u_n \right) \left(\sum_{n=1}^N w_n v_n \right). \quad (63)$$

In both items, if $(u_n)_{n \geq 1}$ is not constant, then the inequality in the conclusion is strict.

Proof. We prove Item 1 by induction on N , in the case when $(u_n)_{n \geq 1}$ and $(v_n)_{n \geq 1}$ are both non-decreasing, the other case being identical. Obviously, Equation (62) holds for $N = 1$. Now, assume that (62) holds for N , then we proceed to show that (62) holds for $N + 1$. Since

$$u_{n+1} [w_1(v_{n+1} - v_1) + \dots + w_n(v_{n+1} - v_n)] \geq u_1 w_1(v_{n+1} - v_1) + \dots + u_n w_n(v_{n+1} - v_n),$$

the inequality being strict if $(u_n)_{n \geq 1}$ is not constant, we observe that

$$\sum_{n=1}^N w_n u_n v_n + \left(\sum_{n=1}^N w_n \right) u_{N+1} v_{N+1} \geq \left(\sum_{n=1}^N w_n v_n \right) u_{N+1} + \left(\sum_{n=1}^N w_n u_n \right) v_{N+1}. \quad (64)$$

From the induction hypothesis and the Equation (64), it follows that

$$\begin{aligned} \left(\sum_{n=1}^{N+1} w_n \right) \left(\sum_{n=1}^{N+1} w_n u_n v_n \right) &= \left(\sum_{n=1}^N w_n \right) \left(\sum_{n=1}^N w_n u_n v_n \right) + w_{N+1} \left(\sum_{n=1}^N w_n u_n v_n \right) \\ &\quad + w_{N+1}^2 u_{N+1} v_{N+1} + \left(\sum_{n=1}^N w_n \right) w_{N+1} u_{N+1} v_{N+1} \\ &\geq \left(\sum_{n=1}^N w_n \right) \left(\sum_{n=1}^N w_n u_n v_n \right) + w_{N+1}^2 u_{N+1} v_{N+1} \\ &\quad + \left(\sum_{n=1}^N w_n v_n \right) u_{N+1} w_{N+1} + \left(\sum_{n=1}^N w_n u_n \right) v_{N+1} w_{N+1} \\ &\geq \left(\sum_{n=1}^N w_n u_n \right) \left(\sum_{n=1}^N w_n v_n \right) + w_{N+1}^2 u_{N+1} v_{N+1} \\ &\quad + \left(\sum_{n=1}^N w_n v_n \right) u_{N+1} w_{N+1} + \left(\sum_{n=1}^N w_n u_n \right) v_{N+1} w_{N+1} \\ &= \left(\sum_{n=1}^{N+1} w_n u_n \right) \left(\sum_{n=1}^{N+1} w_n v_n \right). \end{aligned}$$

This completes the proof of item 1. Equation (63) can then be proved by reversing all the inequalities in the proof of (62) above. \square

This result is proved by DeAngelis et al. [9, Lemma 2.6] for Part (2) and in [10, Proposition A.3] for part (1), where $w_n = 1$ for all $n \geq 1$. Here we generalize this result to any positive sequence.

References

- [1] Arditi R., Lobry C. and Sari T., “Is dispersal always beneficial to carrying capacity? New insights from the multi-patch logistic equation”, *Theor. Popul. Biol.*, 106 (2015), 45-59. doi: 10.1016/j.tpb.2015.10.001
- [2] Arditi R., Lobry C. and Sari T., “Asymmetric dispersal in the multi-patch logistic equation”, *Theor. Popul. Biol.*, 120 (2018), 11-15. doi: 10.1016/j.tpb.2017.12.006
- [3] Arino J., Bajoux N. and Kirkland S., “Number of source patches required for population persistence in a source-sink metapopulation with explicit movement”, *Bull. Math. Biol.*, 81 (2019), No. 6, 1916-1942. doi: 10.1007/s11538-019-00593-1
- [4] Arino J., “Diseases in metapopulations”, in *Modeling and Dynamics of Infectious Diseases* (eds. Ma Z., Zhou Y. and Wu J.), World Scientific (2009), 64-122. doi: 10.1142/9789814261265%_0003
- [5] Cosner C., Beier J.C., Cantrell R.S., Impoinvil D., Kapitanski L., Potts M.D., Troyo A. and Ruan S., “The effects of human movement on the persistence of vector-borne diseases”, *J. Theoret. Biol.*, 258 (2009), No. 4, 550-560. doi: 10.1016/j.jtbi.2009.02.016
- [6] Castillo C. and Thieme H.R., “Asymptotically autonomous epidemic models”, in *Mathematical Population Dynamics: Analysis of Heterogeneity 2* (eds. Arino O., Axelrod D.E., Kimmel M., Langlais M.), *Theory of Epidemics* (1995), 33-50.
- [7] Castillo G., C.W. and Castillo C., “A Tour of the Basic Reproductive Number and the Next Generation of Researchers”, in *An Introduction to Undergraduate Research in Computational and Mathematical Biology* (eds. Callender H., Capaldi A., Diaz C.), Nature Publishing Group (2020), 87-124.
- [8] DeAngelis D.L., Travis C.C. and Post W.M., “Persistence and stability of seed-dispersal species in a patchy environment”, *Theoret. Popul. Biol.*, 16 (1979), No. 2, 107-125. doi: 10.1016/0040-5809(79)90008-X
- [9] DeAngelis D.L., Ni W. and Zhang B., “Dispersal and heterogeneity: single species”, *J. Math. Biol.*, 72 (2016), No. 1-2, 239-254. doi: 10.1007/s00285-015-0879-y
- [10] DeAngelis D.L., Ni W. and Zhang B., “Effects of diffusion on total biomass in heterogeneous continuous and discrete-patch systems”, *Theor. Ecol.*, 9 (2016), No. 4, 443-453. doi: 10.1007/s12080-016-0302-3
- [11] DeAngelis D.L. and Zhang B., “Effects of dispersal in a non-uniform environment on population dynamics and competition: a patch model approach”, *Discrete Contin. Dyn. Syst. Ser. B*, 19 (2014), No. 10, 3087-3104. doi: 10.3934/dcdsb.2014.19.3087
- [12] Elbetch B., Benzekri T., Massart D. and Sari T., “The multi-patch logistic equation”, *Discrete Contin. Dyn. Syst. Ser. B*, 26 (2021), No. 12, 6405-6424. doi: 10.3934/dcdsb.2021025

- [13] Freedman H.I., Rai B. and Waltman P., “Mathematical Models of Population Interactions with Dispersal II: Differential Survival in a Change of Habitat”, *J. Math. Anal. Appl.*, 115 (1986), No. 1, 140-154. doi: 10.1016/0022-247X(86)90029-6
- [14] Freedman H.I. and Waltman P., “Mathematical Models of Population Interactions with Dispersal I: Stability of two habitats with and without a predator”, *SIAM J. Appl. Math.*, 32 (1977), No. 3, 631-648. doi: 10.1137/0132052
- [15] Gao D., “How does dispersal affect the infection size?”, *SIAM J. Appl. Math.*, 80 (2020), No. 5, 2144-2169. doi: 10.1137/19M130652X
- [16] Gao D. and Dong C.P., “Fast diffusion inhibits disease outbreaks”, *Proc. Am. Math. Soc.*, 148 (2020), No. 4, 1709-1722. doi: 10.1090/proc/14868
- [17] Guo H., Li M. and Shuai Z., “Global stability of the endemic equilibrium of multigroup SIR epidemic models”, *Can. Appl. Math. Q.*, 14 (2006), No. 3, 259-284.
- [18] Holt R.D., “Population dynamics in two patch environments: some anomalous consequences of an optimal habitat distribution”, *Theor. Popul. Biol.*, 28 (1985), No. 2, 181-208. doi: 10.1016/0040-5809(85)90027-9
- [19] Hanski I., *Metapopulation Ecology*, Oxford University Press, 1st ed., New York, 1999.
- [20] Levin S.A., “Dispersion and population interactions”, *Amer. Natur.*, 108 (1974), No. 960, 207-228. doi: 10.1086/282900
- [21] Levin S.A., “Spatial patterning and the structure of ecological communities”, in *Some Mathematical Questions in Biology* (ed. Oster G.), Lecture Notes on Mathematics in the Life Sciences (1976), 1-35.
- [22] Lobry C., Sari T. and Touhami S., “On Tykhonov’s theorem for convergence of solutions of slow and fast systems”, *Electron. J. Differential Equations*, 19 (1998), 1-22.
- [23] Markus L., “Asymptotically autonomous differential systems”, in *Contributions to the Theory of Nonlinear Oscillations III*, (ed. Lefschetz S.), Princeton University Press (1956), 17-30. doi: 10.1515/9781400882175-003
- [24] Nagahara K., Lou Y. and Yanagida E., “Maximizing the total population with logistic growth in a patchy environment”, *J. Math. Biol.*, 82 (2021), No. 1, 1-50. doi: 10.1007/s00285-021-01565-7
- [25] Poggiale J.C., Auger P., Nérini D., Manté C. and Gilbert F., “Global production increased spatial heterogeneity in a population dynamics model”, *Acta Biotheoretica*, 53 (2005), No. 4, 359-370. doi: 10.1007/s10441-005-4890-3
- [26] Thieme H.R., “Convergence results and a Poincaré-Bendixson trichotomy for asymptotically autonomous differential equations”, *J. Math. Biol.*, 30 (1992), No. 7, 755-763. doi: 10.1007/BF00173267
- [27] Thieme H.R., “Asymptotically Autonomous Differential Equations in the Plane”, *Rocky Mountain J. Math.*, 24 (1993), No. 1, 351-380. doi: 10.1216/rmj/1181072470
- [28] Tikhonov A.N., “Systems of differential equations containing small parameters in the derivatives”, *Mat. Sb. (N.S.)*, 31 (1952), No. 3, 575-586.

- [29] Van den Driessche P. and Watmough J., “Further Notes on the Basic Reproduction Number”, in *Mathematical Epidemiology* (eds. Brauer F., van den Driessche P., Wu J.), *Lecture Notes in Mathematics* (2008), 159-178. doi:10.1007/978-3-540-78911-6_6
- [30] Wu H., Wang Y., Li Y. and DeAngelis D.L., “Dispersal asymmetry in a two-patch system with source-sink populations”, *Theor. Popul. Biol.*, 131 (2020), 54-65. doi: 10.1016/j.tpb.2019.11.004
- [31] Wasow W.R., *Asymptotic Expansions for Ordinary Differential Equations*, Robert E. Krieger Publishing Company, Huntington, New York, 1976.
- [32] Yurk B.P. and Cobbold C.A., “Homogenization techniques for population dynamics in strongly heterogeneous landscapes”, *J. Biol. Dyn.*, 12 (2018), No. 1, 171-193. doi: 10.1080/17513758.2017.1410238
- [33] Zaker N., Ketchemen L. and Lutscher F., “The Effect of Movement Behavior on Population Density in Patchy Landscapes”, *Bull. Math. Biol.*, 82 (2020), No. 1, 1-24. doi: 10.1007/s11538-019-00680-3