



# On dispersal and population growth for multistate matrix models

Chi-Kwong Li<sup>\*</sup>, Sebastian J. Schreiber

*Department of Mathematics, The College of William and Mary, P.O. Box 8795, Williamsburg, VA 23187, United States*

Received 27 July 2005; accepted 17 March 2006

Available online 22 May 2006

Submitted by R.A. Brualdi

---

## Abstract

To describe the dynamics of stage-structured populations with  $m$  stages living in  $n$  patches, we consider matrix models of the form  $\mathbf{SD}$ , where  $\mathbf{S}$  is a block diagonal matrix with  $n \times n$  column substochastic matrices  $S_1, \dots, S_m$  along the diagonal and  $\mathbf{D}$  is a block matrix whose blocks are  $n \times n$  nonnegative diagonal matrices. The matrix  $\mathbf{S}$  describes movement between patches and the matrix  $\mathbf{D}$  describes growth and reproduction within the patches. Consider the multiple arc directed graph  $G$  consisting of the directed graphs corresponding to the matrices  $S_1, \dots, S_m$ , where each directed graph is drawn in a different color. We say  $G$  has a *polychromatic cycle* if  $G$  has a directed cycle that includes arcs of more than one color. We prove that  $\rho(\mathbf{SD}) \leq \rho(\mathbf{D})$  for all block matrices  $\mathbf{D}$  with nonnegative diagonal blocks if and only if  $G$  has no polychromatic cycle. Applications to ecological models are presented.

© 2006 Elsevier Inc. All rights reserved.

*AMS classification:* 15A51; 15A18; 92D25

*Keywords:* Stochastic matrices; Multistate matrix models; Population growth

---

## 1. Introduction

Denote by  $\rho(A)$  the spectral radius of a square matrix  $A$ . The column sum norm of  $A = (a_{ij})_{1 \leq i, j \leq n}$  is defined by

---

<sup>\*</sup> Corresponding author. Tel.: +1 757 221 2042; fax: +1 757 221 7400.

*E-mail addresses:* [ckli@math.wm.edu](mailto:ckli@math.wm.edu) (C.-K. Li), [sjs@math.wm.edu](mailto:sjs@math.wm.edu) (S.J. Schreiber).

$$\|A\| = \max \left\{ \sum_{i=1}^n |a_{ij}| : 1 \leq j \leq n \right\}.$$

An  $n \times n$  nonnegative matrix  $S$  is *column substochastic* (respectively, *column stochastic*) if all the column sums are bounded by one (respectively, equal to one). The well-known fact that  $\rho(A) \leq \|A\|$  immediately implies the following proposition.

**Proposition 1.1.** *Let  $D$  be a diagonal nonnegative matrices and  $S$  a column substochastic matrix. Then*

$$\rho(SD) \leq \rho(D).$$

For biological models that account for spatial structure, this fact has some important implications. For example, consider a population (e.g. viruses, animals, plants, molecules) residing in an environment consisting of  $n$  spatial locations or patches. If this population did not disperse across the environment,  $x = (x_1, \dots, x_n)^t$  denotes the vector of abundances (e.g. density, concentration, or expected size) and  $d_i$  is the per-capita growth rate of the population in the  $i$ th patch, then a matrix model of this population is given by

$$x(t + 1) = Dx(t),$$

where  $x(t)$  denotes the vector of abundances in the  $t$ th time step and  $D = \text{diag}(d_1, \dots, d_n)$ . Since  $x(t) = D^t x(0)$ , population  $i$  grows asymptotically at a geometric rate if and only if  $d_i > 1$ . In particular, the entire population  $x_1(t) + \dots + x_n(t)$  grows asymptotically at a geometric rate if and only if  $\rho(D) = \max_i d_i > 1$ . Now assume the population disperses across the environment after the reproductive or growth phase. More specifically, a fraction of individuals  $s_{ij}$  successfully moves from location  $j$  to location  $i$ . Then the matrix  $S = (s_{ij})$  is column substochastic, and the population dynamics become

$$x(t + 1) = SDx(t).$$

This dispersing population exhibits growth if and only if  $\rho(SD) > 1$ . Since  $\rho(SD) < \rho(D)$  if not all  $d_i$ 's are equal and  $S$  is irreducible, one can conclude that generically dispersal decreases the asymptotic population growth rate. Moreover, since  $\rho(SD) = \rho(DS)$ , this conclusion holds whether growth occurs before dispersal, as we have assumed, or growth occurs after dispersal.

Many biological models not only account for spatial structure but also account for stage structure. For example, in ecological models, the population may consist of individuals in different age classes (e.g. juveniles, sub-adults, adults) living in different spatial locations [1]. Similarly, epidemiological models often account for different classes of individuals (e.g. susceptible, exposed, infected, removed) as well as spatial structure [3]. For these multistate models, one can ask the following question:

**Question 1.2.** Under what conditions does dispersal decrease the asymptotic growth rate of a population?

In other words, when does the analog of Proposition 1.1 hold for these models. To address this question, consider a population with  $m$  life stages living in  $n$  spatial locations. Let  $x_j^i \in [0, \infty)$  denote the abundance of stage  $i$  individuals in location  $j$ . Then

$$x^i = \begin{pmatrix} x_1^i \\ \vdots \\ x_n^i \end{pmatrix}, \quad x_j = \begin{pmatrix} x_j^1 \\ \vdots \\ x_j^m \end{pmatrix}, \quad \text{and} \quad x = \begin{pmatrix} x^1 \\ \vdots \\ x^m \end{pmatrix}$$

are the vector of abundances in stage  $i$ , the vector of abundances in location  $j$ , and the vector of all abundances, respectively. Let us assume that the population first goes through a growth phase in which individuals within a location survive, reproduce, and change stages. For each spatial location  $j$ , let  $A_j$  be an  $m \times m$  nonnegative matrix representing the growth dynamics in location  $j$ . In the absence of spatial considerations, the population dynamics are given by

$$x_j(t + 1) = A_j x_j(t), \quad j = 1, \dots, n.$$

If  $P$  is the permutation matrix such that

$$Px = P \begin{pmatrix} x^1 \\ \vdots \\ x^m \end{pmatrix} = \begin{pmatrix} x_1 \\ \vdots \\ x_n \end{pmatrix},$$

then

$$x(t + 1) = \mathbf{D}x(t) \quad \text{with } \mathbf{D} = P^t(A_1 \oplus \dots \oplus A_n)P,$$

where  $A_1 \oplus \dots \oplus A_n$  denotes the block diagonal matrix with diagonal blocks  $A_1, \dots, A_n$ . Let  $\{E_{11}, E_{12}, \dots, E_{nn}\}$  be the standard basis for the linear space of  $n \times n$  matrices, and let  $A \otimes B = (a_{ij}) \otimes B = (a_{ij}B)$  be the Kronecker product of the two matrices  $A = (a_{ij})$  and  $B$ . Then  $\mathbf{D}$  is a block matrix whose blocks are  $n \times n$  nonnegative diagonal matrices

$$\mathbf{D} = \sum_{j=1}^n A_j \otimes E_{jj}.$$

To account for movement between locations after the growth phase, let  $S_1, \dots, S_m$  be column substochastic  $n \times n$  matrices whose  $(i, j)$ th entry corresponds to fraction of individuals in a given stage that move successfully from patch  $j$  to patch  $i$ . Including these spatial movements by setting  $\mathbf{S} = S_1 \oplus \dots \oplus S_m$ , we see that the population dynamics become

$$x(t + 1) = \mathbf{SD}x(t).$$

Hunter and Caswell [6] discuss alternative representations of the same model. The population exhibits asymptotically geometric growth if and only if  $\rho(\mathbf{SD}) > 1$ . Unlike the purely spatially structured model, the following example illustrates that the inclusion of spatial movement into a stage-structured population can enhance the asymptotic growth rate of the population.

**Example 1.3.** Consider a population of juveniles and reproductively mature adults living in two spatial locations (e.g. salmon where juveniles develop in fresh water and adults become reproductively mature in the ocean). For illustrative purposes, let us assume that in location 1 (i.e. a freshwater river), all adults produce two juveniles before dying but juveniles can not become reproductively mature adults. In location 2 (i.e. the ocean), all juveniles become reproductively mature adults but progeny produced by the adults in location 2 can not survive (i.e. salmon fry can not develop in salt water). In other words,  $m = n = 2$ ,  $A_1 = \begin{pmatrix} 0 & 2 \\ 0 & 0 \end{pmatrix}$ , and  $A_2 = \begin{pmatrix} 0 & 0 \\ 1 & 0 \end{pmatrix}$ . In which case,

$$\rho(\mathbf{D}) = \rho \left( \begin{pmatrix} 0 & 0 & 2 & 0 \\ 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \end{pmatrix} \right) = 0$$

as  $\mathbf{D}^2 = 0$ : without movement between the patches, the population goes extinct in two time steps. Alternatively, if all juveniles move to patch 2 and all adults move to patch 1, then  $S_1 = \begin{pmatrix} 0 & 0 \\ 1 & 1 \end{pmatrix}$  and  $S_2 = \begin{pmatrix} 1 & 1 \\ 0 & 0 \end{pmatrix}$ . In which case,

$$\rho(\mathbf{SD}) = \rho \left( \begin{pmatrix} 0 & 0 & 0 & 0 \\ 0 & 0 & 2 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 \end{pmatrix} \right) = \sqrt{2}$$

and the population grows asymptotically at a geometric rate.

Proposition 1.1 and Example 1.3 suggest the following general question:

**Question 1.4.** Given  $\mathbf{S}$  and  $\mathbf{D}$  is there a practical way of determining whether or not  $\rho(\mathbf{SD}) \leq \rho(\mathbf{D})$ ?

As a step to understanding this question, we provide in Section 2 an affirmative answer to the following question:

**Question 1.5.** Given  $\mathbf{S}$  is there a practical way of determining whether  $\rho(\mathbf{SD}) \leq \rho(\mathbf{D})$  for all block matrices  $\mathbf{D}$  whose blocks are  $m \times m$  nonnegative diagonal matrices?

In particular, if one does not have much information about  $\mathbf{D}$ , and yet one would like to control or change  $\mathbf{S}$  to ensure that  $\rho(\mathbf{SD}) \leq \rho(\mathbf{D})$ , our result provides useful information.

Our paper is organized as follows. In Section 2, we present the statement of our main theorem (Theorem 2.1) answering Question 1.5, and illustrate applications of the result to biological models studied by other authors. In Section 3, we give a proof of Theorem 2.1.

## 2. Statement of theorem and applications

To state our theorem, we need to introduce one definition. We say the  $n \times n$  nonnegative matrices  $S_1, \dots, S_m$  admit a *polychromatic cycle* if there exist nonzero entries of  $S_1 + \dots + S_m$  at the  $(i_1, i_2), (i_2, i_3) \dots, (i_{k-1}, i_k), (i_k, i_1)$  positions for some distinct  $i_1, \dots, i_k \in \{1, \dots, n\}$  and these nonzero entries do not come from a single matrix  $S_j$ . One can think of this definition as follows. Let  $G_j$  be the directed graph corresponding to the matrix  $S_j$  for  $j = 1, \dots, m$ . In other words,  $G_j$  has vertex set  $V(G_j) = \{1, \dots, n\}$ , and the arc set  $E(G_j)$  consists of arcs  $(r, s)$  from vertex  $r$  to vertex  $s$  if the  $(r, s)$  entry of  $S_j$  is nonzero. Here we do not consider self loops. Consider the multiple arc directed graph  $G$  consisting of the directed graphs  $G_1, \dots, G_m$ , where each directed graph is drawn in a different color. A polychromatic cycle is a (directed) cycle in  $G$  that includes arcs of more than one color. For instance, in Example 1.3, the entries  $(2, 1)$  and  $(1, 2)$  of  $S_1$  and  $S_2$ , respectively, define a polychromatic cycle for the matrices  $S_1$  and  $S_2$ .

**Theorem 2.1.** Suppose  $\mathbf{S} = S_1 \oplus \dots \oplus S_m$  so that  $S_1, \dots, S_m$  are  $n \times n$  column substochastic matrices. The following conditions are equivalent:

- (a) For every block matrix  $\mathbf{D} = (D_{ij})_{1 \leq i, j \leq m}$ , where each  $D_{ij} \in M_n$  is a diagonal matrix with nonnegative diagonal entries, we have  $\rho(\mathbf{SD}) \leq \rho(\mathbf{D})$ .
- (b)  $S_1, \dots, S_m$  do not admit a polychromatic cycle.

The proof of Theorem 2.1 will be given in Section 3. Here we provide two applications to multistate matrix models studied by other researchers.

**Example 2.2 (Patch development models).** Many species live in environments where the patches change state stochastically in time. For instance, a patch of land may be recently disturbed by a fire or have been undisturbed for an extended period of time. Alternatively, a patch of land may have recently experienced a rainfall or be going through a dry spell. Following Horvitz and Schemske [5], Caswell [1, Example 4.3.1.5 on p. 70] describes matrix models that account for changes in the state of a patch. These models assume that each patch exhibits transitions between  $n$  different states and individuals living in these patches can be in one of  $m$  stages, where  $m \geq 2$ . Let  $x_j^i$  denote the abundance of individuals of state  $j$  living in a patch in stage  $i$ . Let  $S$  be a column stochastic matrix that represents the transition probabilities between patch states i.e.  $s_{ij}$  is the probability that the patch goes from state  $j$  to state  $i$  in one time step. Let  $A_1, \dots, A_n$  be  $m \times m$  nonnegative matrices that represent the population dynamics (i.e. transitions between life stages and reproduction) for a patch in states  $1, \dots, n$ . If we define

$$\mathbf{S} = I_m \otimes S \quad \text{and} \quad \mathbf{D} = \sum_{j=1}^n A_j \otimes E_{jj}$$

then the population dynamics are given by  $x(t + 1) = \mathbf{SD}x(t)$ . Since  $\mathbf{S}$  admits a polychromatic cycle if and only if  $S$  has a cycle, Theorem 2.1 implies that  $\rho(\mathbf{SD}) \leq \rho(\mathbf{D})$  for all  $\mathbf{D}$  if and only if  $S$  admits no cycle.

**Example 2.3 (Planktonic dynamics).** Many species have a single life stage that disperses through the environment [6–8]. For matrix models of these species, Theorem 2.1 typically implies that dispersal decreases the asymptotic population growth rate. Sometimes this application of Theorem 2.1 requires augmenting the matrix model by additional state variables. For instance, Caswell [1, Example 4.3.1.3 on p. 68] describes a stage structured and spatially structured model of Davis [2] for planktonic species (e.g. copepods, water fleas, etc.) dispersing in ocean currents during their larval stage. This model assumes that the growth dynamics are independent of spatial location and determined by two nonnegative matrices: a nonnegative  $m \times m$  matrix  $T = (t_{ij})$  that represents the transitions between life stages and a nonnegative  $m \times m$  matrix  $F = (f_{ij})$  that represents reproduction. Since reproduction only contributes to first life stage (i.e. the larval stage),  $F$  has all zero entries except in the first row. To describe dispersal between locations by the larvae, let  $S$  be an  $n \times n$  column substochastic matrix whose  $(i, j)$ th entry  $s_{ij}$  corresponds to the likelihood that a newly born larva from location  $j$  ends up in location  $i$ . Then, the model for the planktonic dynamics is given by

$$x_i(t + 1) = Tx_i(t) + \sum_{j=1}^n s_{ij} Fx_j(t), \quad i = 1, \dots, n,$$

where the first term corresponds to transitions between life stages and the second term corresponds to new larvae dispersing to an  $i$ th patch. Equivalently,

$$x(t + 1) = \mathbf{A}x(t), \quad \text{where } \mathbf{A} = T \otimes I_n + (S \oplus (I_{m-1} \otimes I_n))(F \otimes I_n). \tag{2.1}$$

In particular, if the planktonic larvae do not disperse between spatial locations, then the matrices  $S$  and  $\mathbf{A}$  reduce to  $I_n$  and  $(T + F) \otimes I_n$ , respectively. We claim that

$$\rho(T \otimes I_n + (S \oplus (I_{m-1} \otimes I_n))(F \otimes I)) \leq \rho((T + F) \otimes I_n) \tag{2.2}$$

for any column substochastic matrix  $S$ . In other words, dispersal of the larva reduces the asymptotic growth rate of the population.

To prove this claim using Theorem 2.1, we need to introduce an extra variable  $\tilde{x}_j^0$  that keeps track of the newly born larval stage. Let  $\tilde{x}_j^i$  for  $i = 0, 1, \dots, m$  and  $j = 1, \dots, n$  correspond to

the abundance of life stage  $i$  in location  $j$ ,  $\tilde{x}^i = \begin{pmatrix} \tilde{x}_1^i \\ \vdots \\ \tilde{x}_n^i \end{pmatrix}$ , and  $\tilde{x} = \begin{pmatrix} \tilde{x}^0 \\ \vdots \\ \tilde{x}^m \end{pmatrix}$ . Moreover, define

$$\mathbf{D} = \begin{pmatrix} f_{11} & f_{11} & f_{12} & \dots & f_{1m} \\ t_{11} & t_{11} & t_{12} & \dots & t_{1m} \\ t_{21} & t_{21} & t_{22} & \dots & t_{2m} \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ t_{m1} & t_{m1} & t_{m2} & \dots & t_{mm} \end{pmatrix} \otimes I_n \quad \text{and} \quad \mathbf{S} = S \oplus (I_m \otimes I_n).$$

With the inclusion of this additional variable, the planktonic model in (2.1) becomes

$$\tilde{x}(t + 1) = \mathbf{SD}\tilde{x}(t).$$

Furthermore, we have

$$\begin{pmatrix} I_n & I_n & 0_{n,n(m-1)} \\ 0_{n(m-1),n} & 0_{n(m-1),n} & I_{m-1} \otimes I_n \end{pmatrix} \mathbf{SD} = \mathbf{A} \begin{pmatrix} I_n & I_n & 0_{n,n(m-1)} \\ 0_{n(m-1),n} & 0_{n(m-1),n} & I_{m-1} \otimes I_n \end{pmatrix}.$$

Hence, if  $\tilde{x} = \begin{pmatrix} \tilde{x}^0 \\ \tilde{x}^1 \\ \vdots \\ \tilde{x}^m \end{pmatrix}$  is a nonnegative right eigenvector of  $\mathbf{SD}$ , then  $x = \begin{pmatrix} \tilde{x}^0 + \tilde{x}^1 \\ \tilde{x}^2 \\ \vdots \\ \tilde{x}^m \end{pmatrix} \neq 0$  is a

nonnegative right eigenvector of  $\mathbf{A}$  with the same eigenvalue. Conversely, if  $y = (y_1, \dots, y_n)$  is a left eigenvector of  $\mathbf{A}$ , then  $\tilde{y} = (y_1, y_1, y_2, \dots, y_n) \neq 0$  is a left eigenvector of  $\mathbf{SD}$  with the same eigenvalue. It follows that that  $\rho(\mathbf{SD}) = \rho(\mathbf{A})$ . Evidently,  $\mathbf{S}$  does not admit a polychromatic cycle. By Theorem 2.1, we have  $\rho(\mathbf{A}) = \rho(\mathbf{SD}) \leq \rho(\mathbf{D})$  and thus (2.2) holds for any column substochastic matrix  $S$ .

### 3. Proof of Theorem 2.1

In this section, we give a proof of Theorem 2.1. Because our proof contains some intricate combinatorial arguments and constructions, we give several examples at the end of this section to illustrate the ideas and constructions in our proofs. In particular, Example 3.1 illustrates the idea and construction in the proof of (a)  $\Rightarrow$  (b), and the other examples illustrate the ideas and constructions in the proof of (b)  $\Rightarrow$  (a). Readers may study the examples along with the proofs to gain better insight.

Throughout this section, we will assume that  $P$  is the permutation matrix such that

$$Px = P \begin{pmatrix} x^1 \\ \vdots \\ x^m \end{pmatrix} = \begin{pmatrix} x_1 \\ \vdots \\ x_n \end{pmatrix}. \tag{3.1}$$

We often use the notation  $PS P^t = \tilde{\mathbf{S}}$  and  $PD P^t = \tilde{\mathbf{D}}$ . Note that  $\tilde{\mathbf{D}}$  will be a direct sum of  $n$  matrices of order  $m \times m$ ; and  $\tilde{\mathbf{S}}$  will be an  $n \times n$  block matrix such that each block is an  $m \times m$  diagonal matrix.

Also, we will continue to use the graph theory notation introduced at the beginning of Section 2. Note that if  $G_j$  is the directed graph of  $S_j$ , and if we relabel the vertices of  $G_j$ , it is the same as replacing  $S_j$  by  $Q^t S_j Q$  for a suitable  $n \times n$  permutation matrix  $Q$ . Note that replacing  $S_j$  by  $Q^t S_j Q$  for all  $j = 1, \dots, m$ , is the same as replacing  $\mathbf{S}$  by  $(I_m \otimes Q)^t \mathbf{S} (I_m \otimes Q)$ . If we also replace each  $\mathbf{D}$  by  $(I_m \otimes Q)^t \mathbf{D} (I_m \otimes Q)$ , conditions (a) and (b) will not be affected.

We will write  $X \geq Y$  if  $X - Y$  is nonnegative. Clearly, if  $X, Y$  and  $Z$  are nonnegative matrices satisfying  $X \geq Y$ , then  $XZ \geq YZ$  and hence  $\rho(XZ) \geq \rho(YZ)$ ; see [4, Theorem 8.4.5].

**The proof of (a)  $\Rightarrow$  (b):** We prove the contrapositive. Suppose  $\neg(b)$  holds, i.e.,  $S_1, \dots, S_m$  admit a polychromatic cycle. We will construct an  $m \times m$  block matrix  $\mathbf{D}$  such that  $\rho(\mathbf{D}) > \rho(\mathbf{SD})$  and each block is a nonnegative diagonal matrix.

Assume  $S_1, \dots, S_m$  admit a polychromatic cycle  $(i_1, \dots, i_k)$ . Replacing  $\mathbf{S}$  and  $\mathbf{D}$  by  $(I \otimes Q)^t \mathbf{S} (I \otimes Q)$  and  $(I_m \otimes Q)^t \mathbf{D} (I_m \otimes Q)$  with a suitable permutation matrix  $Q$ , we may assume that  $(i_1, \dots, i_k) = (1, \dots, k)$ . Then  $G$  has a polychromatic cycle with arcs  $(1, 2), (2, 3), \dots, (k, 1)$ . For  $i = 1, \dots, m$ , construct the  $n \times n$  zero-one matrix  $F_i$  from  $S_i$  by changing those entries of  $S_i$  which contribute to the arcs  $(1, 2), (2, 3), \dots, (k, 1)$  of the polychromatic cycle to one, and changing all other entries to zero. Let  $\mathbf{F} = d(F_1 \oplus \dots \oplus F_m)$ , where

$$d = \min\{s : s \text{ is a positive entry of } \mathbf{S}\}.$$

Then  $\mathbf{F}$  has  $k$  nonzero entries all equal to  $d$  and  $\mathbf{S} \geq \mathbf{F}$ . We will construct  $\mathbf{D}$  so that

$$\rho(\mathbf{FD}) = \rho(\mathbf{DF}) > \rho(\mathbf{D}). \tag{3.2}$$

Since  $\mathbf{S} \geq \mathbf{F}$ , it will then follow that  $\rho(\mathbf{SD}) \geq \rho(\mathbf{FD}) > \rho(\mathbf{D})$ .

Let  $P$  be the permutation matrix satisfying (3.1), and let  $\tilde{\mathbf{F}} = P \mathbf{F} P^t = d(\tilde{F}_{ij})_{1 \leq i, j \leq n}$ , where  $\tilde{F}_{ij}$  are  $m \times m$  diagonal matrices. Then  $\tilde{F}_{ij} = 0$  if  $(i, j) \notin \{(1, 2), (2, 3), \dots, (k-1, k), (k, 1)\}$ . Since the nonzero entries in  $F_1 + \dots + F_m$  do not come from a single matrix  $F_r$ , the matrices  $\tilde{F}_{12}, \dots, \tilde{F}_{k-1,k}, \tilde{F}_{k,1}$  cannot be identical and each of them has only one nonzero entry equal to 1 on the diagonal.

Let  $\{e_1, \dots, e_m\}$  be the standard basis for  $\mathbb{R}^m$ , and let  $\mu > \max\{d^{-k}, 1\}$ . Suppose

$$\tilde{\mathbf{D}} = \tilde{D}_1 \oplus \dots \oplus \tilde{D}_n,$$

where

- (i)  $\tilde{D}_j = [e_r + \mu \sum_{s \neq r} e_s] e_r^t$  if the  $(r, r)$  entry of  $\tilde{F}_{j, j+1}$  is nonzero for  $j \in \{1, \dots, k-1\}$ ,
- (ii)  $\tilde{D}_k = [e_r + \mu \sum_{s \neq r} e_s] e_r^t$  if the  $(r, r)$  entry of  $\tilde{F}_{k,1}$  is nonzero,
- (iii)  $\tilde{D}_j = 0_m$  for  $j \in \{k+1, \dots, n\}$ .

Then  $\rho(\tilde{\mathbf{D}}) = 1$ , and  $\tilde{\mathbf{D}}\tilde{\mathbf{F}} = (R_{ij})$ , where

$$(R_{12}, R_{23}, \dots, R_{k-1,k}, R_{k,1}) = (d\tilde{D}_1, \dots, d\tilde{D}_k)$$

and  $R_{ij} = 0$  for all other  $(i, j)$ . Thus,

$$(\tilde{\mathbf{D}}\tilde{\mathbf{F}})^k = T_1 \oplus \cdots \oplus T_k \oplus 0_{m(n-k)},$$

where each  $T_j$  is a cyclic product of  $d\tilde{D}_1, \dots, d\tilde{D}_k$  so that all  $T_j$  have the same eigenvalues. Let  $\tilde{D}_i = [e_{r_i} + \mu \sum_{s \neq r_i} e_s] e_{r_i}^t$  for  $i = 1, \dots, k$ . Then

$$\tilde{D}_1 \cdots \tilde{D}_k = (v_1 \cdots v_{n-1}) \left[ e_{r_1} + \mu \sum_{s \neq r_1} e_s \right] e_{r_n}^t,$$

where for  $j = 1, \dots, k - 1$ ,

$$v_j = e_{r_j}^t \left[ e_{r_{j+1}} + \mu \sum_{s \neq r_{j+1}} e_s \right] = \begin{cases} 1 & \text{if } r_j = r_{j+1}, \\ \mu & \text{if } r_j \neq r_{j+1}. \end{cases}$$

Since  $\tilde{F}_{12}, \dots, \tilde{F}_{k1}$  are not identical, the matrices  $\tilde{D}_1, \dots, \tilde{D}_k$  are not identical. So, there is  $j \in \{1, \dots, k - 1\}$  such that  $v_j = \mu$ . As a result, the matrix  $T_1 = d^k(\tilde{D}_1 \cdots \tilde{D}_k)$  has exactly one nonzero column which contains a diagonal entry of the form  $d^k \mu^s$  for some  $s \geq 1$ . It follows that  $\rho((\tilde{\mathbf{D}}\tilde{\mathbf{F}})^k) = \rho(T_1) = d^k \mu^s > 1$  because  $\mu > \max\{d^{-k}, 1\}$  and  $s \geq 1$ . Let  $\mathbf{D} = P^t \tilde{\mathbf{D}} P$ . Then

$$\rho(\mathbf{D}\mathbf{F}) = \rho(\tilde{\mathbf{D}}\tilde{\mathbf{F}}) > 1 = \rho(\tilde{\mathbf{D}}) = \rho(\mathbf{D}).$$

**The proof of (b)  $\Rightarrow$  (a):** Suppose  $\mathbf{S}$  satisfy condition (b). Using the graph theory description of condition (b) before Theorem 2.1, we see that if  $G_i$  is the directed graph associated with  $S_i$  for  $i = 1, \dots, m$ , then in the multiple arc directed graph  $G$  with vertex set  $V(G) = \{1, \dots, n\}$  and multiple arc set  $E(G) = \cup_{j=1}^m E(G_j)$  every cycle is monochromatic. We will show that

$$\rho(\mathbf{S}\mathbf{D}) \leq \rho(\mathbf{D}) \tag{3.3}$$

for all  $m \times m$  block matrices  $\mathbf{D}$  in which each block is a diagonal matrix with positive diagonal entries. By continuity, the inequality (3.3) will also hold for  $m \times m$  block matrices  $\mathbf{D}$  such that each block is a nonnegative diagonal matrix.

To prove inequality (3.3), we can impose some additional assumptions on the matrix  $\mathbf{D}$  as follows. Suppose  $P\mathbf{D}P^t = \tilde{\mathbf{D}} = D_1 \oplus \cdots \oplus D_n$ . We can replace  $\mathbf{D}$  by  $\mathbf{D}/\rho(\mathbf{D})$  on both sides of (3.3) and assume that  $\rho(\mathbf{D}) = 1$ . Furthermore, we can replace  $D_j$  by  $D_j/\rho(D_j)$  for each  $j = 1, \dots, n$ , and assume that  $\rho(D_1) = \cdots = \rho(D_n) = 1$ . Note that after such a replacement,  $\rho(\mathbf{D})$  will stay the same, but  $\rho(\mathbf{S}\mathbf{D})$  may increase.

Now, suppose  $\mathbf{D}$  is an  $m \times m$  block matrix such that each block is an  $n \times n$  diagonal matrix with positive diagonal entries, and  $P\mathbf{D}P^t = \tilde{\mathbf{D}} = D_1 \oplus \cdots \oplus D_n$  with  $\rho(D_j) = 1$  for each  $j = 1, \dots, n$ . Our strategy is to show that there exists a diagonal matrix  $\mathbf{V}$  with positive diagonal entries such that  $\mathbf{V}\mathbf{D}\mathbf{V}^{-1}$  is column stochastic and  $\mathbf{V}\mathbf{S}\mathbf{V}^{-1}$  has column sum norm at most one. It will then follow that:

$$\rho(\mathbf{S}\mathbf{D}) = \rho(\mathbf{V}\mathbf{S}\mathbf{V}^{-1}\mathbf{V}\mathbf{D}\mathbf{V}^{-1}) \leq \|\mathbf{V}\mathbf{S}\mathbf{V}^{-1}\mathbf{V}\mathbf{D}\mathbf{V}^{-1}\| \leq \|\mathbf{V}\mathbf{S}\mathbf{V}^{-1}\| \|\mathbf{V}\mathbf{D}\mathbf{V}^{-1}\| \leq 1 = \rho(\mathbf{D}).$$

To achieve our goal, let  $Q$  be an  $n \times n$  permutation matrix such that

$$Q^t(S_1 + \cdots + S_m)Q = (T_{ij})_{1 \leq i, j \leq k} \tag{3.4}$$

is in block upper triangular form (Frobenius normal form) and each diagonal block  $T_{jj}$  is an irreducible square matrix. We may assume that  $Q = I_n$ ; otherwise, replace  $\mathbf{S}$  by  $(I_m \otimes Q)\mathbf{S}(I_m \otimes Q)$ , and replace  $\mathbf{D}$  by  $(I_m \otimes Q)^t \mathbf{D} (I_m \otimes Q)$ , accordingly.



Suppose  $C_1, \dots, C_k$  are the strongly connected components of the directed graph  $G$  with multiple arcs corresponding to the matrices  $T_{11}, \dots, T_{kk}$ . For  $r \in \{1, \dots, m\}$ , if  $(p, q)$  is an arc of  $G_r$  in  $C_i$ , then  $(p, q)$  belongs to a cycle in  $G_r$  lying entirely in  $C_i$ . Otherwise, the arc  $(p, q)$  will lie in a polychromatic cycle in the strongly connected component  $C_i$ . So, each arc of  $G_r$  in  $C_i$  must belong to a strongly connected component of  $G_r$  lying entirely in  $C_i$ . Consequently, every  $C_i$  is a directed graph consisting of strongly connected sub-components  $C_{i,1}, \dots, C_{i,p_i}$  from the directed graphs  $G_1, \dots, G_m$ .

To construct the desired diagonal matrix  $\mathbf{V}$ , we need to relabel the vertices of  $G$ . We do this in two steps.

**Step 1:** We relabel the sub-components  $C_{i,1}, \dots, C_{i,p_i}$  of  $C_i$  for each  $i$  as follows.

If  $p_i = 1$ , then no relabeling is needed. If  $p_i > 1$  then  $C_{i,1}$  has a common vertex with one of directed graphs  $C_{i,2}, \dots, C_{i,p_i}$ . Since  $C_1$  is strongly connected, permuting the subscripts  $2, \dots, p_i$  if necessary, we may assume that  $C_{i,1}$  and  $C_{i,2}$  have common vertices. Observe that  $C_{i,1}$  and  $C_{i,2}$  cannot have two or more common vertices; otherwise, there will be a two-color cycle in  $C_i$ . If  $p_i > 2$ , then  $C_{i,1} \cup C_{i,2}$  will have a common vertex with one of the sub-components  $C_{i,3}, \dots, C_{i,p_i}$ , say,  $C_{i,3}$ . Here the union of two directed graphs means the union of vertex sets as well as the arc sets of the two directed graphs. Again,  $C_{i,1} \cup C_{i,2}$  and  $C_{i,3}$  cannot have two common vertices; otherwise, there will be a polychromatic cycle in  $C_i$ . We can repeat this argument until we are done with  $C_{i,p_i}$ .

**Step 2:** We relabel the vertex set  $V(G) = \{1, \dots, n\}$  as follows.

Assume that  $C_1$  is a union of  $C_{1,1}, C_{1,2}, \dots, C_{1,p_1}$ , and  $C_1$  has  $q_1$  vertices. Relabel these vertices by the indices  $1, \dots, q_1$  as follows. Arrange the vertices of  $C_{1,1}$  in a sequence (in any order); then continue with the vertices in  $C_{1,2}$ , till we get to the vertices of  $C_{1,p_1}$ . Assume  $C_2$  is a union of  $C_{2,1}, \dots, C_{2,p_2}$  and has  $q_2$  vertices. Then use a similar procedure to relabel the vertices of  $C_2$  by the indices  $q_1 + 1, \dots, q_1 + q_2$ . Continue this process till we are done with  $C_k$ .

Suppose  $Q$  is the  $n \times n$  permutation matrix corresponding to the relabeling of vertices of  $G$  in Step 2. Again, we may assume that  $Q = I_n$ . Otherwise, replace  $\mathbf{S}$  by  $(I_m \otimes Q)^t \mathbf{S} (I_m \otimes Q)$ , and replace  $\mathbf{D}$  by  $(I_m \otimes Q)^t \mathbf{D} (I_m \otimes Q)$  accordingly.

We are now ready to construct the desired diagonal matrix  $\mathbf{V}$ . Since  $D_j$  is a positive matrix and  $\rho(D_j) = 1$ , there is a left Perron vector  $\mathbf{v}_j = (v_{j1}, \dots, v_{jm})$  with positive entries such that  $v_{j1} = 1$  and  $\mathbf{v}_j D_j = \mathbf{v}_j$  for  $j = 1, \dots, n$ ; see [4, Theorem 8.2.11]. If  $V_j$  is the diagonal matrix with diagonal entries  $v_{j1}, \dots, v_{jm}$ , then  $V_j D_j V_j^{-1}$  is column stochastic. Let  $\mu_1, \dots, \mu_n$  be positive numbers, and  $\mathbf{V}$  be such that  $P \mathbf{V} P^t = \mu_1 V_1 \oplus \dots \oplus \mu_n V_n$ . Then

$$P(\mathbf{V} \mathbf{D} \mathbf{V}^{-1}) P^t = (V_1 D_1 V_1^{-1} \oplus \dots \oplus V_n D_n V_n^{-1})$$

is column stochastic. Consider the diagonal entries of  $\mathbf{V}$  arranged as follows:

$$\begin{array}{cccc}
 \mu_1 v_{11} & \mu_1 v_{12} & \cdots & \mu_1 v_{1m} \\
 \mu_2 v_{21} & \mu_2 v_{22} & \cdots & \mu_2 v_{2m} \\
 \vdots & \vdots & \dots & \vdots \\
 \mu_n v_{n1} & \mu_n v_{n2} & \cdots & \mu_n v_{nm}
 \end{array} \tag{3.5}$$

If  $U_i = \text{diag}(\mu_1 v_{1i}, \mu_2 v_{2i}, \dots, \mu_n v_{ni})$  for  $i = 1, \dots, m$ , then  $\mathbf{V} = U_1 \oplus \dots \oplus U_m$ . We will select positive numbers  $\mu_1, \dots, \mu_n$  so that the column sum norm of  $U_i S_i U_i^{-1}$  is at most one for each  $i$ . Then  $\mathbf{V}$  will satisfy the desired property.

If  $C_1$  has only one vertex, set  $\mu_1 = 1$ . Otherwise, suppose the sub-component  $C_{1,1}$  of  $C_1$  is a strongly connected component of  $G_r$  and has  $\alpha_1$  vertices. Then in the matrix  $\mathbf{S} = S_1 \oplus \cdots \oplus S_m$ , only  $S_r$  has a leading  $\alpha_1 \times \alpha_1$  irreducible principal submatrix, and all other  $S_j$  will have a diagonal leading  $\alpha_1 \times \alpha_1$  principal submatrix. For  $j = 1, \dots, \alpha_1$ , select  $\mu_j$  so that  $\mu_j v_{jr} = 1$ . Then for any choices of other  $\mu_j$  for  $j > \alpha_1$ , if  $\mathbf{VSV}^{-1} = \widehat{S}_1 \oplus \widehat{S}_2 \oplus \cdots \oplus \widehat{S}_m$  then the  $\alpha_1 \times \alpha_1$  leading submatrix of  $\widehat{S}_j$  will be the same as that of  $S_j$  for  $j = 1, \dots, m$ . Note that at this point, we have selected the first  $\alpha_1$  rows in list (3.5).

Next, consider the sub-component  $C_{1,2}$  in  $C_1$ . By the discussion in the Step 1 relabeling procedure,  $C_{1,1}$  and  $C_{1,2}$  has exactly one common vertex say,  $s \in \{1, \dots, \alpha_1\}$ . Suppose  $C_{1,2}$  comes from  $G_t \neq G_r$  and has  $\alpha_2 - \alpha_1 + 1$  vertices (including  $s$ ). Then  $\mu_s v_{st}$  is determined in the preceding paragraph, and we can choose  $\mu_{\alpha_1+1}, \dots, \mu_{\alpha_2}$  so that  $\mu_j v_{jt} = \mu_s v_{st}$  for  $j = \alpha_1 + 1, \dots, \alpha_2$ . Then for any choices of other  $\mu_j$  for  $j > \alpha_2$ , if  $\mathbf{VSV}^{-1} = \widehat{S}_1 \oplus \widehat{S}_2 \oplus \cdots \oplus \widehat{S}_m$  then the  $\alpha_2 \times \alpha_2$  leading submatrix of  $\widehat{S}_j$  will be the same as that of  $S_j$  for  $j = 1, \dots, m$ . Note that at this point, we have selected the first  $\alpha_2$  rows in list (3.5).

In the labeling of the sub-components of  $C_1$  in the Step 1 relabeling procedure, every additional sub-component would have exactly one common vertex with the union of the previously labeled sub-components. So, we can repeat the above process to select  $\mu_j$  until we are done with all the sub-components of  $C_1$ . Since  $C_1$  has  $q_1$  vertices, for any choices of other  $\mu_j$  with  $j > q_1$ , if  $\mathbf{VSV}^{-1} = \widehat{S}_1 \oplus \widehat{S}_2 \oplus \cdots \oplus \widehat{S}_m$  then the  $q_1 \times q_1$  leading submatrix of  $\widehat{S}_j$  will be the same as that of  $S_j$  for  $j = 1, \dots, m$ . Note that at this point, we have selected the first  $q_1$  rows in list (3.5).

Now we move to the second strongly connected component  $C_2$  of  $G$ . Let us identify the smallest constant  $\eta$  such that

$$\mu_i v_{i\ell} \leq \eta v_{j\ell} \quad \text{if } j > q_1, \text{ and } \ell \in \{1, \dots, n\} \tag{3.6}$$

for each number  $\mu_i v_{i\ell}$  in the first  $q_1$  rows of list (3.5). In the future selection of  $\mu_j$  for  $j > q_1$ , we will insist that  $\mu_j \geq \eta$ . Then

$$(\mu_i v_{i\ell})(\mu_j v_{j\ell})^{-1} \leq 1 \quad \text{for all } i \leq q_1 < j, \text{ and } \ell \in \{1, \dots, n\}. \tag{3.7}$$

In other words, all diagonal entries of  $(\mu_i V_i)(\mu_j V_j)^{-1}$  are less than or equal to one for  $i \leq q_1 \leq j$ . If  $C_2$  has only one vertex, set  $\mu_{q_1+1}$  to be any number larger than  $\eta$ . Otherwise, consider the first sub-component  $C_{2,1}$  in  $C_2$ . Assume  $C_{2,1}$  belongs to  $G_\ell$  and has  $\beta_1$  vertices. Then for  $j = q_1 + 1, \dots, q_1 + \beta_1$ , choose  $\mu_j \geq \eta$  so that  $\mu_j v_{j\ell}$  are all equal. Then for any choices of other  $\mu_j$  with  $j > q_1 + \beta_2$ , if  $\mathbf{VSV}^{-1} = \widehat{S}_1 \oplus \widehat{S}_2 \oplus \cdots \oplus \widehat{S}_m$  the  $q_1 \times q_1$  leading submatrix and the following  $\beta_1 \times \beta_1$  principal submatrix of  $\widehat{S}_j$  will be the same as that of  $S_j$  for  $j = 1, \dots, m$ . Note that at this point, we have selected the first  $q_1 + \beta_1$  rows in list (3.5).

Applying similar arguments as those to  $C_1$  with the precaution that  $\mu_j \geq \eta$  for  $j = q_1 + 1, \dots, q_1 + q_2$ , we can select the first  $q_1 + q_2$  rows of list (3.5) so that for any choices of other  $\mu_j$  with  $j > q_1 + q_2$ , if  $\mathbf{VSV}^{-1} = \widehat{S}_1 \oplus \widehat{S}_2 \oplus \cdots \oplus \widehat{S}_m$  then the  $q_1 \times q_1$  leading submatrix and the following  $q_2 \times q_2$  principal submatrix of  $\widehat{S}_j$  will be the same as that of  $S_j$  for  $j = 1, \dots, m$ .

Note that each  $S_j$  has an upper triangular block form according to the Frobenius normal form  $(T_{ij})$  in (3.4). If  $\mathbf{VSV}^{-1} = \widehat{S}_1 \oplus \widehat{S}_2 \oplus \cdots \oplus \widehat{S}_m$  and there is a nonzero  $(i, j)$  entry  $\xi$  in  $S_\ell$  such that  $1 \leq i \leq q_1 < j \leq q_1 + q_2$ , then it is obtained from the original entry by multiplying the quantity  $(\mu_i v_{i\ell})(\mu_j v_{j\ell})^{-1}$ , and hence it is not larger than the original entry by (3.7). So, the first  $q_1 + q_2$  columns of  $\mathbf{VSV}^{-1}$  have column sums bounded above by one.

Now, update  $\eta$  in (3.6) so that

$$\mu_i v_{i\ell} \leq \eta v_{j\ell} \quad \text{if } i \leq q_1 + q_2 < j, \ell \in \{1, \dots, n\}$$

for all numbers  $\mu_i v_{i\ell}$  in the first  $q_1 + q_2$  rows of list (3.5). Then we can proceed to consider the strongly connected component  $C_3$  of  $G$  and determine  $\mu_j$  for  $j = q_1 + q_2 + 1, \dots, q_1 + q_2 + q_3$ .

Repeating the above argument, we can determine  $\mu_1, \dots, \mu_n$  and construct the diagonal matrix  $\mathbf{V} = U_1 \oplus \dots \oplus U_m$  so that  $U_i S_i U_i^{-1}$  is still in upper block triangular form, whose diagonal blocks are the same as those of  $S_i$ , and  $S_i - U_i S_i U_i^{-1}$  is nonnegative. Hence,

$$\|\mathbf{V}\mathbf{S}\mathbf{V}^{-1}\| \leq \|\mathbf{S}\| \leq 1. \quad \square$$

The following example illustrates the construction in our proof of the implication (a)  $\Rightarrow$  (b).

**Example 3.1.** Let  $\mathbf{S} = S_1 \oplus S_2$ , where

$$S_1 = \frac{1}{2} \begin{pmatrix} 2 & 1 & 0 \\ 0 & 1 & 2 \\ 0 & 0 & 0 \end{pmatrix} \quad \text{and} \quad S_2 = \frac{1}{2} \begin{pmatrix} 0 & 0 & 2 \\ 0 & 2 & 0 \\ 2 & 0 & 0 \end{pmatrix}.$$

Then the directed graph  $G$  of  $S_1 + S_2$  admits a polychromatic cycle with arcs (1, 2), (2, 3), (3, 1). Let  $\mathbf{F} = F_1 \oplus F_2$  with

$$F_1 = \begin{pmatrix} 0 & 1/2 & 0 \\ 0 & 0 & 1/2 \\ 0 & 0 & 0 \end{pmatrix} \quad \text{and} \quad F_2 = \begin{pmatrix} 0 & 0 & 0 \\ 0 & 0 & 0 \\ 1/2 & 0 & 0 \end{pmatrix}.$$

Then  $\mathbf{S} \geq \mathbf{F}$  and  $\tilde{\mathbf{F}} = \mathbf{P}\mathbf{F}\mathbf{P}^t = (\tilde{F}_{ij})_{1 \leq i, j \leq 3}$ , where

$$\tilde{F}_{12} = \tilde{F}_{23} = \frac{1}{2} \begin{pmatrix} 1 & 0 \\ 0 & 0 \end{pmatrix}, \quad \tilde{F}_{31} = \frac{1}{2} \begin{pmatrix} 0 & 0 \\ 0 & 1 \end{pmatrix}, \quad \text{and} \quad \tilde{F}_{ij} = \begin{pmatrix} 0 & 0 \\ 0 & 0 \end{pmatrix} \text{ otherwise.}$$

Suppose  $\mu > 0$  satisfies  $\mu^2 > 8$ . Construct  $\mathbf{D}$  so that  $\tilde{\mathbf{D}} = \mathbf{P}\mathbf{D}\mathbf{P}^t = \tilde{D}_1 \oplus \tilde{D}_2 \oplus \tilde{D}_3$  with

$$\tilde{D}_1 = \tilde{D}_2 = \begin{pmatrix} 1 & 0 \\ \mu & 0 \end{pmatrix} \quad \text{and} \quad \tilde{D}_3 = \begin{pmatrix} 0 & \mu \\ 0 & 1 \end{pmatrix}.$$

Then

$$\tilde{\mathbf{D}}\tilde{\mathbf{F}} = \frac{1}{2} \begin{pmatrix} 0_2 & \tilde{D}_1 & 0_2 \\ 0_2 & 0_2 & \tilde{D}_2 \\ \tilde{D}_3 & 0_2 & 0_2 \end{pmatrix}$$

and

$$(\tilde{\mathbf{D}}\tilde{\mathbf{F}})^3 = \frac{1}{8} \{ \tilde{D}_1 \tilde{D}_2 \tilde{D}_3 \oplus \tilde{D}_2 \tilde{D}_3 \tilde{D}_1 \oplus \tilde{D}_3 \tilde{D}_1 \tilde{D}_2 \} = \frac{1}{8} \left\{ \begin{pmatrix} 0 & \mu \\ 0 & \mu^2 \end{pmatrix} \oplus \begin{pmatrix} \mu^2 & 0 \\ \mu^3 & 0 \end{pmatrix} \oplus \begin{pmatrix} \mu^2 & 0 \\ \mu & 0 \end{pmatrix} \right\}$$

has spectral radius  $\mu^2/8 > 1$ , and hence  $\rho(\tilde{\mathbf{D}}\tilde{\mathbf{F}}) > 1$ . Since  $\mathbf{S} \geq \mathbf{F}$ , we have

$$\rho(\mathbf{S}\mathbf{D}) \geq \rho(\mathbf{F}\mathbf{D}) = \rho(\mathbf{D}\mathbf{F}) = \rho(\tilde{\mathbf{D}}\tilde{\mathbf{F}}) > 1 = \rho(\mathbf{D}).$$

The next three examples illustrate the idea and construction in our proof for implication (b)  $\Rightarrow$  (a).

**Example 3.2.** Suppose  $\mathbf{S} = S_1 \oplus S_2 \oplus S_3$ , and  $Q$  is a permutation matrix such that  $Q^t S_1 Q$ ,  $Q^t S_2 Q$  and  $Q^t S_3 Q$ , are the column stochastic matrices

$$\frac{1}{3} \begin{pmatrix} 3 & 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 0 & 0 \\ 0 & 1 & 1 & 0 & 0 \\ 0 & 0 & 0 & 2 & 0 \\ 0 & 0 & 0 & 0 & 2 \end{pmatrix}, \quad \frac{1}{3} \begin{pmatrix} 3 & 1 & 1 & 1 & 1 \\ 0 & 1 & 0 & 1 & 0 \\ 0 & 0 & 2 & 0 & 0 \\ 0 & 1 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 2 \end{pmatrix}, \quad \frac{1}{3} \begin{pmatrix} 3 & 1 & 1 & 1 & 1 \\ 0 & 1 & 0 & 0 & 1 \\ 0 & 0 & 2 & 0 & 0 \\ 0 & 0 & 0 & 2 & 0 \\ 0 & 1 & 0 & 0 & 1 \end{pmatrix}.$$

We may relabel the vertices of  $G$  and assume that  $Q = I_5$ . Then  $G$  has two strongly connected components  $C_1$  and  $C_2$ , where  $C_1$  has vertex set  $\{1\}$  and  $C_2$  has vertex set  $\{2, 3, 4, 5\}$ . Furthermore,  $C_2$  has three sub-components  $C_{21}, C_{22}, C_{23}$  with vertex sets  $\{2, 3\}, \{2, 4\}, \{2, 5\}$ , respectively. Now, suppose  $\mathbf{D}$  is given such that  $P^t \mathbf{D} P = D_1 \oplus \dots \oplus D_5$  with

$$D_1 = \frac{1}{12} \begin{pmatrix} 1 & 3 & 8 \\ 1 & 3 & 8 \\ 1 & 3 & 8 \end{pmatrix}, \quad D_2 = D_3 = \frac{1}{6} \begin{pmatrix} 1 & 2 & 3 \\ 1 & 2 & 3 \\ 1 & 2 & 3 \end{pmatrix}, \quad \text{and} \quad D_4 = D_5 = \frac{1}{4} \begin{pmatrix} 1 & 1 & 2 \\ 1 & 1 & 2 \\ 1 & 1 & 2 \end{pmatrix}.$$

Then  $\rho(D_j) = 1$  for each  $j$ ;  $(1, 3, 8)D_1 = (1, 3, 8)$ ,  $(1, 2, 3)D_j = (1, 2, 3)$  for  $j = 2, 3$ ,  $(1, 1, 2)D_j = (1, 1, 2)$  for  $j = 4, 5$ . Consider the list

$$\begin{matrix} \mu_1 1 & \mu_1 3 & \mu_1 8 \\ \mu_2 1 & \mu_2 2 & \mu_2 3 \\ \mu_3 1 & \mu_3 2 & \mu_3 3 \\ \mu_4 1 & \mu_4 1 & \mu_4 2 \\ \mu_5 1 & \mu_5 1 & \mu_5 2. \end{matrix}$$

We follow the construction in the proof and choose  $\mu_1 = 1$ , and insist that:

$$\mu_j \geq \eta = \max\{v_{1j}/v_{\ell j} : 1 \leq j \leq 3; 2 \leq \ell \leq 5\} = 4, \quad j = 2, 3, 4, 5.$$

Now,  $C_{21}$  comes from  $G_1$  and has vertex set  $\{2, 3\}$ ; so, we choose  $\mu_2 = \mu_3 = 4$ . Next,  $C_{22}$  comes from  $S_2$  and has vertex set  $\{2, 4\}$ ; so, we choose  $\mu_4$  such that  $\mu_2 v_{22} = \mu_4 v_{42}$ , i.e.,  $\mu_4 = 8$ . Finally,  $C_{23}$  comes from  $S_3$  and has vertex set  $\{2, 5\}$ ; so, we choose  $\mu_5$  such that  $\mu_2 v_{32} = \mu_5 v_{35}$ , i.e.,  $\mu_5 = 6$ . Consequently, we have

$$\mathbf{V} = \text{diag}(1, 4, 4, 8, 6) \oplus \text{diag}(3, 8, 8, 8, 6) \oplus \text{diag}(8, 12, 12, 16, 12).$$

Then  $\mathbf{V} \mathbf{D} \mathbf{V}^{-1}$  is column stochastic, and  $\|\mathbf{V} \mathbf{S} \mathbf{V}^{-1}\| \leq 1$ .

**Example 3.3.** Suppose  $\mathbf{S}$  is such that  $S_2 = \dots = S_m = I_n$ . Let  $Q$  be a permutation matrix such that  $Q^t S_1 Q$  is in the block upper triangular form. Then  $Q^t(S_1 + \dots + S_m)Q = (T_{ij})_{1 \leq i, j \leq k}$  is in block upper triangular form (3.4) as in our proof. We can relabel the vertices of  $G$  and assume that  $Q = I_n$ . Now,  $G$  has  $k$  connected components, and each comes from  $G_1$ . Suppose  $T_{jj}$  is  $n_j \times n_j$  for  $j = 1, \dots, k$ . Given any  $\mathbf{D}$ , we can follow the construction in our proof and obtain the desired  $\mathbf{V} = U_1 \oplus \dots \oplus U_m$  such that  $U_1 = \mu_1 I_{n_1} \oplus \dots \oplus \mu_k I_{n_k}$  with  $\mu_1 < \dots < \mu_k$ . Then  $\mathbf{V} \mathbf{D} \mathbf{V}^{-1}$  is column stochastic,  $U_1 S_1 U_1^{-1}$  is in block upper triangular form such that  $S_1 \geq U_1 S_1 U_1^{-1}$ , and  $U_i S_i U_i^{-1} = I_n$  for  $i = 2, \dots, m$ ; thus,  $\|\mathbf{V} \mathbf{S} \mathbf{V}^{-1}\| \leq 1$ .

**Example 3.4.** Suppose there is an  $n \times n$  permutation matrix  $Q$  such that  $Q^t(S_1 + \dots + S_n)Q$  is in upper triangular form. Thus, each connected component of  $G$  has only one vertex. We may relabel the vertices of  $G$  and assume that  $Q = I_n$ . Given any  $\mathbf{D}$ , we can follow the construction

in our proof and obtain the desired  $\mathbf{V} = U_1 \oplus \cdots \oplus U_m$  such that each  $U_i$  is a diagonal matrix with diagonal entries arranged in ascending order. Since each  $S_i$  is in upper triangular form, we see that  $S_i - U_i S_i U_i^{-1}$  is a nonnegative matrix in strictly upper triangular form. So,  $\mathbf{V}\mathbf{D}\mathbf{V}^{-1}$  is column stochastic and  $\|\mathbf{V}\mathbf{S}\mathbf{V}^{-1}\| \leq 1$ .

## Acknowledgments

Research of the authors were partially supported by NSF. The first author was also supported by a HK RCG grant. The authors would like to thank Professor Eva Czabarka for her helpful comments. They are also indebted to the referee for the thorough reviews.

## References

- [1] H. Caswell, *Matrix Population Models*, Sinauer, Sunderland, Massachusetts, 2001.
- [2] C.S. Davis, Interaction of a copepod population with the mean circulation on Georges Bank, *J. Marine Res.* 42 (1984) 573–590.
- [3] O. Diekmann, J.A.P. Heesterbeek, *Mathematical epidemiology of infectious diseases*, Wiley Series in Mathematical and Computational Biology, John Wiley & Sons Ltd., Chichester, 2000.
- [4] R.A. Horn, C.R. Johnson, *Matrix Analysis*, Cambridge University Press, Cambridge, 1985.
- [5] C.C. Horvitz, D.W. Schemske, Chapter: Seed dispersal and environmental heterogeneity in a neotropical herb: a model of population and patch dynamics, *Frugivores and seed dispersal*, Dr. W. Junk Publishers, Dordrecht, Netherlands, 1986.
- [6] C.M. Hunter, H. Caswell, The use of the vec-permutation matrix in spatial matrix population models, *Ecol. Model.* 188 (2005) 15–21.
- [7] J.D. Lebreton, Demographic models for subdivided populations: the renewal equation approach, *Theoret. Popul. Biol.* 49 (1996) 291–313.
- [8] J.T. Wootton, D.A. Bell, A metapopulation model of the peregrine falcon in California: viability and management strategies, *Ecol. Appl.* 2 (1992) 307–321.