



Stability criteria for a multi-city epidemic model with travel delays and infection during travel

Dedicated to Professor Tibor Krisztin on the occasion of his 60th birthday

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Abstract. We present a compartmental SIR (susceptible–infected–recovered) model to describe the propagation of an infectious disease in a human population, when individuals travel between p different cities. The time needed for travel between any two locations is incorporated, and we assume that disease progression is possible during travel. The model is equivalent to an autonomous system of differential equations with multiple delays, and each delayed term is defined through a system of ordinary differential equations. We establish some necessary and sufficient conditions for the disease-free equilibrium of the model to be asymptotically stable.

Keywords: stability, functional differential equations, dynamically defined delayed term, epidemic model.

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1 Introduction

In recent years there has been an increasing interest in the mathematical modelling community for the spatial spread of infectious diseases. Historical examples like the 1918 influenza pandemic, as well as recent threats like the 2002–2003 SARS epidemic, the 2009 A(H1N1) influenza pandemic, the 2012 MERS coronavirus outbreak, and the 2015 extensive Ebola virus (EBOV) outbreak in West Africa exemplify that national boundaries or oceans have never prevented infectious diseases to reach distant territories. Differential equation-based models for spatial epidemic spread have been discussed for an array of infectious diseases including measles, influenza, malaria and SARS; most recent literature includes the work of Arino and coauthors [1,2], Gao and coauthors [7], Hsieh and coauthors [10], Li and coauthors [14], Ruan

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and coauthors [21], and Wang and coauthors [28,29]. These studies are mainly concerned with the spatial dispersal of infected individuals in connected regions, and do not take it into account that long distance travel also provides a platform for disease dynamics. A recent report by the European Centre for Disease Prevention and Control [5] confirms that some infectious diseases, such as tuberculosis, measles and seasonal influenza, are transmissible during commercial flights, even on those with a duration of less than eight hours. There are some documented cases of influenza transmission occurring during train journeys, including when seven healthy people caught Tamiflu-resistant A(H1N1) influenza while travelling for 42 hours on a train in Vietnam (Mai et al. [17]).

In their disease transmission models, Cui and coauthors [3], Lui and Takeuchi [16], and Takeuchi and coauthors [25] examined the possibility that individuals may contract a disease while travelling. They used the SIS (susceptible–infected–susceptible) framework to describe disease progression, and implicitly used the assumption that transportation between regions takes place instantaneously. This approach ignores that the delay in the transportation can lead to a significant underestimation of the roles of global travel in the spread of some diseases of major public health concern, such as SARS and influenza. These diseases are rapidly progressive (typically, a few days), so even a short delay (a fraction of a day) can be significant. This consideration motivated the work of Liu, Wu and Zhou [15], where they introduced the time delay needed to complete the travel into the SIS-type epidemic model and considered the possibility of infections during this time. The global dynamics was described by Nakata [18] for two identical regions, and then by Nakata and Röst [19,20] for multiple regions with different characteristics and general travel networks.

For most diseases with a potential to pose a threat for a pandemic (like influenza, SARS and measles), the SIS-type model is inadequate. This was first noted by Knipl and Röst [12] when they suggested a general framework for SIR (susceptible–infected–recovered)-based disease progression with a general incidence term. This concept was further elaborated in an SEAIR (susceptible–exposed–asymptomatic infected–symptomatic infected–recovered)-framework initiated by Knipl, Röst and Wu [13], closely mimicking the spatiotemporal evolution of the H1N1 pandemic in multiple regions connected by long distance travel. In these models the framework is required to include a subsystem structured by age, to incorporate the possibility of disease transmission during travel. In the subsystem, age represents the time elapsed since the start of the travel. Following the assumption that transmitting the infection is possible on-board, the model setup leads to a system of delay differential equations with delay representing travel time. The two systems, describing the dynamics in the regions and during transportation, are interconnected; initial values of the system for disease spread during travel depend on the state of the system in the regions, while the inflow term of arrivals to the regions after being in transportation for a fixed time arises as the solution of the subsystem for travel. If the subsystem can be solved analytically (this was the case for the age-structured SI model used in [15,18,20]), then the system for disease spread in the regions decouple from the subsystem. Recalling that initial data of the system during travel comes from the equations for the regions, the inflow term of travellers completing a trip appears as a delayed feedback term. On the other hand, in case of choosing SIR-type models as an epidemic building block when the subsystem does not admit a closed form solution, the delayed term in the system for the regions cannot be expressed explicitly (as exemplified in [12,13]), but is defined dynamically, via the solutions of another system.

The general form of initial value problems for nonautonomous functional differential equations with dynamically defined delayed feedback function were studied by Knipl [11]. Fun-

damental properties were obtained such as the usual existence, uniqueness and continuous dependence result for the solution, and some further, biologically relevant properties were also studied. Although the paper [11] provides mathematical framework for studying the SIR model in [12] and the SEAIR model in [13], it remains a major theoretical challenge to classify the global dynamics in such systems. In particular, no general method has been developed to our knowledge to investigate stability in delayed systems with dynamically defined delayed feedback function. To classify local stability of the disease-free steady state is extremely important as it characterizes whether the disease will become extinct or persist. This was attempted in [13] in the special case when the subsystem structured by travel time did obtain a closed form solution. In this paper, we present a general, delayed SIR model for the spread of an infectious disease when individuals travel between p cities. The model setup leads to a differential system with multiple delays, and transport-related infection is incorporated; more precisely the corresponding subsystems obtain SIR dynamics therefore cannot be solved analytically. We describe local stability properties of the disease-free equilibrium. The approach presented in the paper has the potential to be adopted to other delayed differential models where the delayed feedback term is defined via the solution of another system.

2 Model formulation

We describe the spread of an infectious disease in a population of individuals who travel between different locations. Considering p locations (cities), we divide the entire populations into the disjoint classes I_j , S_j , and R_j , where the letters I , S , and R represent the compartments of infected, susceptible, and recovered individuals respectively, and lower index j ($1 \leq j \leq p$) specifies the current city. Let $I_j(t)$, $S_j(t)$, and $R_j(t)$ be the number of individuals belonging to I_j , S_j , and R_j respectively, at time t . We assume constant recruitment terms A_j , while d_j denotes natural mortality in city j . Disease transmission is modelled by standard incidence and the transmission rate between an infected individual and a susceptible individual in city j is denoted by β_j . Infected individuals in city j recover at rate μ_j . For the total population currently being in city j at time t , we use the notation $N_j(t) = I_j(t) + S_j(t) + R_j(t)$.

We denote by $m_{k,j}$ the movement rate from city j to city k ($1 \leq j, k \leq p, j \neq k$), furthermore, we let $m_{j,j} = 0$ for $j = 1, \dots, p$. The total travel outflow rate from city j is given by $M_j = \sum_{k=1}^p m_{k,j}$, and the movement matrix \mathcal{M} is defined as $\mathcal{M} = (m_{j,k})_{j,k=1}^p$. We assume that \mathcal{M} is an irreducible matrix. We introduce the terms $\mathcal{I}_{j,k}^\tau$, $\mathcal{S}_{j,k}^\tau$, and $\mathcal{R}_{j,k}^\tau$ for the inflow of infected, susceptible, and recovered individuals respectively, into city j from k ($j \neq k$). These terms will be precisely described later. Based on the assumptions above, the following system is formulated for the dynamics of an infectious disease in p cities:

$$\left\{ \begin{array}{l} \frac{d}{dt} I_j(t) = \beta_j S_j(t) \frac{I_j(t)}{N_j(t)} - (\mu_j + d_j) I_j(t) - \left(\sum_{k=1}^p m_{k,j} \right) I_j(t) + \sum_{\substack{k=1 \\ k \neq j}}^p \mathcal{I}_{j,k}^\tau(t), \\ \frac{d}{dt} S_j(t) = A_j - d_j S_j(t) - \beta_j S_j(t) \frac{I_j(t)}{N_j(t)} - \left(\sum_{k=1}^p m_{k,j} \right) S_j(t) + \sum_{\substack{k=1 \\ k \neq j}}^p \mathcal{S}_{j,k}^\tau(t), \\ \frac{d}{dt} R_j(t) = \mu_j I_j(t) - d_j R_j(t) - \left(\sum_{k=1}^p m_{k,j} \right) R_j(t) + \sum_{\substack{k=1 \\ k \neq j}}^p \mathcal{R}_{j,k}^\tau(t), \\ j = 1, \dots, p. \end{array} \right. \quad (2.1)$$

The time needed to complete travel from city k to j is denoted by $\tau_{j,k}$. It is assumed that $\tau_{j,k} > 0$ whenever $m_{j,k} > 0$; however, for convenience we let $\tau_{j,k} = 0$ for all j, k such that $m_{j,k} = 0$, that is, there is no movement from city k to j .

A principal assumption of our modelling approach is that we do not neglect the possibility of infection and recovery during the travel between the cities. To describe these processes during a travel to city j started from city k at any time t_* , we divide the population during the travel into the classes $i_{j,k}$, $s_{j,k}$, and $r_{j,k}$ for infected, susceptible, and recovered travellers, respectively. Then $i_{j,k}(\theta)$, $s_{j,k}(\theta)$, and $r_{j,k}(\theta)$ give the density of infected, susceptible, and recovered individuals respectively, who started travel at time t_* , with respect to θ that denotes the time elapsed since the beginning of the travel. With

$$n_{j,k}(\theta) = i_{j,k}(\theta) + s_{j,k}(\theta) + r_{j,k}(\theta),$$

and by denoting by $\beta_{j,k}^T$ and $\mu_{j,k}^T$ the transmission rate and the recovery rate respectively, we consider the ODE system for the dynamics of the disease during a travel from city k to j :

$$\begin{cases} \frac{d}{d\theta} i_{j,k}(\theta) = \beta_{j,k}^T s_{j,k}(\theta) \frac{i_{j,k}(\theta)}{n_{j,k}(\theta)} - \mu_{j,k}^T i_{j,k}(\theta), \\ \frac{d}{d\theta} s_{j,k}(\theta) = -\beta_{j,k}^T s_{j,k}(\theta) \frac{i_{j,k}(\theta)}{n_{j,k}(\theta)}, \\ \frac{d}{d\theta} r_{j,k}(\theta) = \mu_{j,k}^T i_{j,k}(\theta). \end{cases} \quad (2.2)$$

This system can be obtained in a compact form as

$$\frac{d}{d\theta} y_{j,k}(\theta) = g_{j,k}(y_{j,k}(\theta)), \quad (2.3)$$

where $y_{j,k} = (i_{j,k}, s_{j,k}, r_{j,k})$ and $g_{j,k} : \mathbb{R}^3 \rightarrow \mathbb{R}^3$ is defined by the right hand sides in (2.2). Standard results from the theory of ordinary differential equations guarantee that for any initial data $y_* \in \mathbb{R}^3$ there is a unique solution to (2.3) on $[0, \infty)$. We denote the solution by $\tilde{y}_{j,k}(\theta) = \tilde{y}_{j,k}(\theta; 0, y_*)$.

Let $h_{j,k} : \mathbb{R}^3 \rightarrow \mathbb{R}^3$, $h_{j,k}(v) = m_{j,k}v$, and let $\mathcal{I}_{j,k}, \mathcal{S}_{j,k}, \mathcal{R}_{j,k} : \mathbb{R}^3 \rightarrow \mathbb{R}$ such that

$$\begin{aligned} \mathcal{I}_{j,k}(v) &= (\tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v)))_1, \\ \mathcal{S}_{j,k}(v) &= (\tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v)))_2, \\ \mathcal{R}_{j,k}(v) &= (\tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v)))_3. \end{aligned} \quad (2.4)$$

For $v = (I_k(t_*), S_k(t_*), R_k(t_*))$, note that $h_{j,k}(v)$ gives the number of infected, susceptible, and recovered individuals who leave city k at time t_* to travel to city j . Therefore $\tilde{y}_{j,k}(\theta; 0, h_{j,k}(v))$ describes the density of individuals in the three disease classes θ units of time after the beginning of their travel from city k to j , that started at t_* . In particular, for $\theta = \tau_{j,k}$ and using the notations in (2.4), $(\mathcal{I}_{j,k}(v), \mathcal{S}_{j,k}(v), \mathcal{R}_{j,k}(v))$ gives the density of infected, susceptible, and recovered individuals at the end of the travel.

We are now in the position to define the inflow terms $\mathcal{I}_{j,k}^\tau, \mathcal{S}_{j,k}^\tau, \mathcal{R}_{j,k}^\tau$ in system (2.1). The individuals who arrive at time t to city j from k are precisely those who started their travel at time $t - \tau_{j,k}$; hence, letting $t_* = t - \tau_{j,k}$ we derive by the above arguments that the density of infected, susceptible, and recovered individuals arriving at time t to city j from k is given

as $\mathcal{I}_{j,k}(v)$, $\mathcal{S}_{j,k}(v)$, and $\mathcal{R}_{j,k}(v)$ respectively, with $v = (I_k(t - \tau_{j,k}), S_k(t - \tau_{j,k}), R_k(t - \tau_{j,k}))$. It is therefore meaningful to define the inflow terms as

$$\begin{aligned}\mathcal{I}_{j,k}^\tau(t) &:= \mathcal{I}_{j,k}(I_k(t - \tau_{j,k}), S_k(t - \tau_{j,k}), R_k(t - \tau_{j,k})), \\ \mathcal{S}_{j,k}^\tau(t) &:= \mathcal{S}_{j,k}(I_k(t - \tau_{j,k}), S_k(t - \tau_{j,k}), R_k(t - \tau_{j,k})), \\ \mathcal{R}_{j,k}^\tau(t) &:= \mathcal{R}_{j,k}(I_k(t - \tau_{j,k}), S_k(t - \tau_{j,k}), R_k(t - \tau_{j,k})).\end{aligned}$$

Note that these terms depend on some past states of the system and therefore they are delayed terms; however they are not expressed explicitly by the variables of the system (2.1) but are defined dynamically, through the solution of another system (2.2).

Let $\sigma_j = \max\{\tau_{k,j} : 1 \leq k \leq p\}$ for $j = 1, \dots, p$. We define

$$C^\tau = \prod_{j=1}^p C([- \sigma_j, 0], \mathbb{R}) \times \prod_{j=1}^p C([- \sigma_j, 0], \mathbb{R}) \times \prod_{j=1}^p C([- \sigma_j, 0], \mathbb{R}),$$

where for $\sigma > 0$ we denote by $C([- \sigma, 0], \mathbb{R})$ the space of continuous functions on $[- \sigma, 0]$, which is a Banach space with the usual uniform norm $|\phi| = \sup\{\phi(\theta) : - \sigma \leq \theta \leq 0\}$. The generic element of C^τ is denoted by $\phi = (\phi_1, \dots, \phi_{3p})$, and C^τ is a Banach space with the norm $|\phi| = \sum_{n=1}^{3p} |\phi_n|$. The system (2.1) in a compact form reads as

$$\frac{d}{dt}x(t) = f(x_t),$$

where $x = (I_1, \dots, I_p, S_1, \dots, S_p, R_1, \dots, R_p)$, and $f : C^\tau \rightarrow \mathbb{R}^{3p}$, $f = (f_{I1}, \dots, f_{I_p}, f_{S1}, \dots, f_{S_p}, f_{R1}, \dots, f_{R_p})$ where $f_{Ij}, f_{Sj}, f_{Rj} : C^\tau \rightarrow \mathbb{R}$ are defined as the right hand side of the corresponding equation in system (2.1), for $1 \leq j \leq p$. For the segment of the solution we use the usual notation x_t where $x_t = (x_1^t, \dots, x_{3p}^t)$, and $x_j^t = x_j(t + \theta)$, $x_{j+p}^t = x_{j+p}(t + \theta)$ and $x_{j+2p}^t = x_{j+2p}(t + \theta)$ for $\theta \in [- \sigma_j, 0]$, $1 \leq j \leq p$.

The feasible phase space in our model is defined as the nonnegative cone C_+^τ of C^τ . The general form of initial value problems for nonautonomous functional differential equations with dynamically defined delayed feedback function were studied in [11]. Fundamental properties were obtained such as the usual existence, uniqueness and continuous dependence result for the solution, and some further, biologically relevant properties were also studied.

3 Stability analysis

In Appendix A we show that the system (2.1) has a unique disease-free equilibrium (DFE) \hat{E} , where $\hat{E} \in C^\tau$ is the constant function equal to $(0, \dots, 0, \bar{S}_1, \dots, \bar{S}_p, 0, \dots, 0) \in \mathbb{R}^{3p}$ for all values of its argument and

$$(\bar{S}_1, \dots, \bar{S}_p)^T = (\text{diag}(d_1 + M_1, \dots, d_p + M_p) - \mathcal{M})^{-1}(A_1, \dots, A_p)^T.$$

For $1 \leq k \leq p$ we denote by \hat{E}_k the constant function equal to $(0, \bar{S}_k, 0)$ for all values of its argument. To study the stability of \hat{E} , we investigate the linear variational system corresponding to \hat{E} :

$$\frac{d}{dt}X(t) = FX_t, \quad F = Df(\hat{E}). \quad (3.1)$$

$F = Df(\hat{E})$ is a bounded linear operator from C_+^τ to \mathbb{R}^{3p} , that is, $F \in L(C_+^\tau, \mathbb{R}^{3p})$. For $\phi \in C_+^\tau$ we give $(F\phi)_n$ by computing $Df_n(\hat{E})\phi$ for $1 \leq n \leq 3p$. Note that $f_j = f_{I_j}$, $f_{j+p} = f_{S_j}$, $f_{j+2p} = f_{R_j}$, therefore

$$\begin{aligned} (Df_j(\hat{E})\phi) &= (\beta_j - \mu_j - d_j - M_j)\phi_j(0) \\ &\quad + \sum_{k=1, j \neq k}^p \text{grad}(\mathcal{I}_{j,k}(\hat{E}_k)) \cdot (\phi_k(-\tau_{j,k}), \phi_{k+p}(-\tau_{j,k}), \phi_{k+2p}(-\tau_{j,k})), \\ (Df_{j+p}(\hat{E})\phi) &= (-d_j - M_j)\phi_{j+p}(0) \\ &\quad + \sum_{k=1, j \neq k}^p \text{grad}(\mathcal{S}_{j,k}(\hat{E}_k)) \cdot (\phi_k(-\tau_{j,k}), \phi_{k+p}(-\tau_{j,k}), \phi_{k+2p}(-\tau_{j,k})), \\ (Df_{j+2p}(\hat{E})\phi) &= (-d_j - M_j)\phi_{j+2p}(0) \\ &\quad + \sum_{k=1, j \neq k}^p \text{grad}(\mathcal{R}_{j,k}(\hat{E}_k)) \cdot (\phi_k(-\tau_{j,k}), \phi_{k+p}(-\tau_{j,k}), \phi_{k+2p}(-\tau_{j,k})) \end{aligned} \quad (3.2)$$

for $1 \leq j \leq p$, where we use the usual notation $a \cdot b$ for the dot product of two vectors a and b . In Appendix B we show that the gradients on the right hand side are obtained as

$$\begin{pmatrix} \text{grad}(\mathcal{I}_{j,k}(\hat{E}_k)) \\ \text{grad}(\mathcal{S}_{j,k}(\hat{E}_k)) \\ \text{grad}(\mathcal{R}_{j,k}(\hat{E}_k)) \end{pmatrix} = \frac{\partial}{\partial v} \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(0, \bar{S}_k, 0)) = m_{j,k} \begin{pmatrix} e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} & 0 & 0 \\ -\beta_{j,k}^T \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} & 1 & 0 \\ \mu_{j,k}^T \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} & 0 & 1 \end{pmatrix}. \quad (3.3)$$

The stability of the trivial solution of the linear variational system (3.1) is determined by the characteristic equation $\Delta(\lambda) = 0$, which arises by seeking solutions of the form $X(t) = e^{\lambda t} u$ where $u \in \mathbb{R}^{3p}$. If the stability modulus, defined as

$$s(F) = \max\{\text{Re } \lambda : \Delta(\lambda) = 0\}$$

is negative then the trivial solution of the linear variational system is locally asymptotically stable (LAS) whereas it is unstable if $s(F) > 0$. These stability properties extend to the stability of the DFE \hat{E} in the system (2.1) by the principle of linearised stability.

Using the equations (3.2) and (3.3) it is possible to rewrite the linear variational system (3.1) as

$$\frac{d}{dt} X(t) = F^0 X(t) + \sum_{\substack{j,k=1 \\ j \neq k}}^p F^{j,k} X(t - \tau_{j,k}),$$

where $F^0, F^{j,k} \in \mathbb{R}^{3p \times 3p}$,

$$F^0 = \begin{pmatrix} \text{diag}(\beta_j - \mu_j - d_j - M_j)_{j=1}^p & \mathcal{O} & \mathcal{O} \\ \star & \text{diag}(-d_j - M_j)_{j=1}^p & \mathcal{O} \\ \star & \mathcal{O} & \text{diag}(-d_j - M_j)_{j=1}^p \end{pmatrix},$$

and

$$(F^{j,k})_{j,j+p,j+2p}^{k,k+p,k+2p} = \begin{pmatrix} \text{grad}(\mathcal{I}_{j,k}(\hat{E}_k)) \\ \text{grad}(\mathcal{S}_{j,k}(\hat{E}_k)) \\ \text{grad}(\mathcal{R}_{j,k}(\hat{E}_k)) \end{pmatrix} = m_{j,k} \begin{pmatrix} e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} & 0 & 0 \\ -\beta_{j,k}^T \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} & 1 & 0 \\ \mu_{j,k}^T \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} & 0 & 1 \end{pmatrix}$$

with all other elements of $F^{j,k}$ being zero. It is useful to note that for any j and k it holds that $(F^{j,k})_{u,v} = 0$ whenever $u \leq p$ and $v > p$. This implies that the first p equations (the infection subsystem) in the linear variational system are independent from the other equations. Moreover, we also note that for any j and k it holds that $(F^{j,k})_{u,v} = 0$ whenever u, v are such that either $p + 1 \leq u \leq 2p$ and $v > 2p$, or $u > 2p$ and $p + 1 \leq v \leq 2p$ holds. It follows that the two systems, formed by equations $p + 1, \dots, 2p$ (the susceptible subsystem) and by equations $2p + 1, \dots, 3p$ (the recovered subsystem) respectively, are also independent from one another. These remarks imply that the characteristic equation factorizes as the product of the characteristic equations of the three subsystems, hence the stability in (3.1) is determined by the stability properties in the three subsystems. In particular, both the susceptible and recovered subsystems can be obtained in the form

$$\frac{d}{dt}w_j(t) = (-d_j - M_j)w_j(t) + \sum_{\substack{k=1 \\ k \neq j}}^p m_{j,k}w_k(t - \tau_{j,k}), \quad j = 1, \dots, p, \quad (3.4)$$

and the infection subsystem reads

$$\frac{d}{dt}z_j(t) = (\beta_j - \mu_j - d_j - M_j)z_j(t) + \sum_{\substack{k=1 \\ k \neq j}}^p m_{j,k}e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)}z_k(t - \tau_{j,k}), \quad j = 1, \dots, p. \quad (3.5)$$

We define the following ODE system, that can be associated with (3.5) by ignoring the delays:

$$\frac{d}{dt}z_j(t) = (\beta_j - \mu_j - d_j - M_j)z_j(t) + \sum_{\substack{k=1 \\ k \neq j}}^p m_{j,k}e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)}z_k(t), \quad j = 1, \dots, p. \quad (3.6)$$

We present now one of our most important results, which states that the stability analysis of the DFE in the nonlinear model (2.1) with dynamically defined delayed feedback function reduces to studying the non-delayed infection subsystem of the linear variational system.

Theorem 3.1. *The DFE of the model (2.1) is LAS if and only if the zero solution of system (3.6) is LAS.*

Proof. By the above arguments, the stability analysis of the DFE in the model (2.1) reduces to studying stability in the systems (3.4) and (3.5). It follows from [20, Theorem 4.2] that the zero solution is LAS in system (3.5) (see system (4.4) in [20] and [9, Theorem 1]).

The principal result of Section 5, Chapter 5 in [24] is that the stability of an equilibrium of a cooperative and irreducible system of delay differential equations is the same as that of an associated system of cooperative ordinary differential equations. To study these properties for system (3.5), we recall some definitions and results from [24] in Appendix C. Then we prove in Appendix D that the system (3.5) is cooperative and irreducible. \square

To study the stability of the linear system (3.6), we define

$$\begin{aligned} T &= (c_{j,k}m_{j,k})_{j,k=1}^p + \text{diag}(\beta_1, \dots, \beta_p), \\ \Sigma &= \text{diag}(\mu_1 + d_1 + M_1, \dots, \mu_p + d_p + M_p), \end{aligned}$$

and obtain the matrix of the linear system as $T - \Sigma$, where we used the short hand notation $c_{j,k} = e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)}$ ($j \neq k$) and $c_{j,j} = 0$. The stability of the zero solution is determined by the eigenvalues of the coefficient matrix; the solution is LAS if $s(T - \Sigma) < 0$ and it is unstable

if $s(T - \Sigma) > 0$, where $s(A)$ denotes the maximum real part of all eigenvalues of any square matrix A .

We say that a square matrix A is a non-singular M-matrix if it has the Z-sign pattern (all entries are non-positive except possibly those in the diagonal) and $A^{-1} \geq 0$ holds (several definitions exist for M-matrices, see [6, Theorem 5.1]). It is easy to see that T is a nonnegative matrix and Σ is a non-singular M-matrix. The proof of the next result follows by similar arguments as those in the proof of [26, Theorem 2], where we denote by $\rho(A)$ the dominant eigenvalue of any square matrix A .

Proposition 3.2. *The zero solution of (3.6) is LAS ($s(T - \Sigma) < 0$) if and only if $\rho(T \cdot \Sigma^{-1}) < 1$, it is unstable ($s(T - \Sigma) > 0$) if and only if $\rho(T \cdot \Sigma^{-1}) > 1$, and it also holds that $s(T - \Sigma) = 0$ if and only if $\rho(T \cdot \Sigma^{-1}) = 1$.*

In the mathematical epidemiology literature, it is common to relate stability properties with the next-generation matrix (NGM) and the basic reproduction number (\mathcal{R}_0) (for description of the next-generation method and computation of this matrix and number, see [4] and the references therein). If we define reproduction (the emergence of new infections) by the matrix T and transition between the infection classes by Σ then the NGM of system (3.6) arises as

$$K := T \cdot \Sigma^{-1} = \begin{pmatrix} \frac{\beta_1}{\mu_1 + d_1 + M_1} & \frac{c_{1,2}m_{1,2}}{\mu_2 + d_2 + M_2} & \frac{c_{1,3}m_{1,3}}{\mu_3 + d_3 + M_3} & \cdots & \frac{c_{1,p}m_{1,p}}{\mu_p + d_p + M_p} \\ \frac{c_{2,1}m_{2,1}}{\mu_1 + d_1 + M_1} & \frac{\beta_2}{\mu_2 + d_2 + M_2} & \frac{c_{2,3}m_{2,3}}{\mu_3 + d_3 + M_3} & \cdots & \frac{c_{2,p}m_{2,p}}{\mu_p + d_p + M_p} \\ \frac{c_{3,1}m_{3,1}}{\mu_1 + d_1 + M_1} & \frac{c_{3,2}m_{3,2}}{\mu_2 + d_2 + M_2} & \frac{\beta_3}{\mu_3 + d_3 + M_3} & \cdots & \frac{c_{3,p}m_{3,p}}{\mu_p + d_p + M_p} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ \frac{c_{p,1}m_{p,1}}{\mu_1 + d_1 + M_1} & \frac{c_{p,2}m_{p,2}}{\mu_2 + d_2 + M_2} & \frac{c_{p,3}m_{p,3}}{\mu_3 + d_3 + M_3} & \cdots & \frac{\beta_p}{\mu_p + d_p + M_p} \end{pmatrix}.$$

The dominant eigenvalue of the NGM is denoted by \mathcal{R}_0 (that is, $\mathcal{R}_0 := \rho(K)$), that is of particular importance as by Theorem 3.1 and Proposition 3.2 it is a threshold quantity for the stability of the DFE in the model (2.1). However, due to the complexity of the NGM it is not possible to derive the formula of \mathcal{R}_0 and therefore we cannot explicitly obtain the stability condition.

The next section is devoted to addressing this issue as we consider two special cases for the movement network: first we investigate the case of two cities, then we obtain more general results as we assume that p cities are organized in a ring-like structure.

4 Stability of system (3.6)

4.1 Two cities

For $p = 2$, the system (3.6) reduces to

$$\begin{aligned} \frac{d}{dt}z_1(t) &= (\beta_1 - \mu_1 - d_1 - m_{2,1})z_1(t) + m_{1,2}e^{\tau_{1,2}(\beta_{1,2}^T - \mu_{1,2}^T)}z_2(t), \\ \frac{d}{dt}z_2(t) &= (\beta_2 - \mu_2 - d_2 - m_{1,2})z_2(t) + m_{2,1}e^{\tau_{2,1}(\beta_{2,1}^T - \mu_{2,1}^T)}z_1(t). \end{aligned} \tag{4.1}$$

We define

$$T_2 = \begin{pmatrix} \beta_1 & m_{1,2}e^{\tau_{1,2}(\beta_{1,2}^T - \mu_{1,2}^T)} \\ m_{2,1}e^{\tau_{2,1}(\beta_{2,1}^T - \mu_{2,1}^T)} & \beta_2 \end{pmatrix}, \quad \Sigma_2 = \begin{pmatrix} \mu_1 + d_1 + m_{2,1} & 0 \\ 0 & \mu_2 + d_2 + m_{1,2} \end{pmatrix}$$

and obtain the coefficient matrix of the linear system as $T_2 - \Sigma_2$. We derive that

$$K_2 := T_2(\Sigma_2)^{-1} = \begin{pmatrix} \frac{\beta_1}{\mu_1 + d_1 + m_{2,1}} & \frac{m_{1,2}e^{\tau_{1,2}(\beta_{1,2}^T - \mu_{1,2}^T)}}{\mu_2 + d_2 + m_{1,2}} \\ \frac{m_{2,1}e^{\tau_{2,1}(\beta_{2,1}^T - \mu_{2,1}^T)}}{\mu_1 + d_1 + m_{2,1}} & \frac{\beta_2}{\mu_2 + d_2 + m_{1,2}} \end{pmatrix},$$

and observe that

$$K_2 \geq K_2^0 := \text{diag} \left(\frac{\beta_1}{\mu_1 + d_1 + m_{2,1}}, \frac{\beta_2}{\mu_2 + d_2 + m_{1,2}} \right),$$

where the relation is meant entry-wise. We introduce the notation

$$\begin{aligned} \mathcal{R}_1 &= \frac{\beta_1}{\mu_1 + d_1}, & \mathcal{R}_2 &= \frac{\beta_2}{\mu_2 + d_2}, \\ \mathcal{P}_1 &= \frac{\beta_1}{\mu_1 + d_1 + m_{2,1}}, & \mathcal{P}_2 &= \frac{\beta_2}{\mu_2 + d_2 + m_{1,2}}, \end{aligned}$$

where \mathcal{R}_j and \mathcal{P}_j are the basic reproduction numbers in city j in the absence of movement and in the absence of inflow into city j , respectively. With these definitions, it holds that $\mathcal{P}_1 < \mathcal{R}_1$ and $\mathcal{P}_2 < \mathcal{R}_2$. We arrive at the following result.

Theorem 4.1. *If $\max \{ \mathcal{P}_1, \mathcal{P}_2 \} \geq 1$ then the zero solution of (4.1) is unstable for all choices of delays.*

Proof. With the definitions above, K_2 and K_2^0 are nonnegative matrices, K_2 is irreducible, and $\rho(K_2^0) = \max \{ \mathcal{P}_1, \mathcal{P}_2 \}$. It follows from the Perron–Frobenius theorem that $\rho(K_2) > \rho(K_2^0)$, therefore $\rho(K_2) > 1$ holds and the assertion of the theorem follows by Proposition 3.2. \square

It remains to investigate stability in the case when $\mathcal{P}_1 < 1$ and $\mathcal{P}_2 < 1$. In this case, $\rho(K_2^0) < 1$ holds so it is meaningful to define

$$\mathcal{T}_2 := \rho((K_2 - K_2^0)(\text{Id}_2 - K_2^0)^{-1}),$$

that can be explicitly calculated as

$$\begin{aligned} \mathcal{T}_2 &= \rho \left(\left(\begin{pmatrix} 0 & \frac{m_{1,2}e^{\tau_{1,2}(\beta_{1,2}^T - \mu_{1,2}^T)}}{\mu_2 + d_2 + m_{1,2}} \\ \frac{m_{2,1}e^{\tau_{2,1}(\beta_{2,1}^T - \mu_{2,1}^T)}}{\mu_1 + d_1 + m_{2,1}} & 0 \end{pmatrix} \left(1 - \frac{\beta_1}{\mu_1 + d_1 + m_{2,1}} \quad 0 \right)^{-1} \right) \right) \\ &= \rho \left(\begin{pmatrix} 0 & \frac{m_{1,2}e^{\tau_{1,2}(\beta_{1,2}^T - \mu_{1,2}^T)}}{\mu_2 + d_2 + m_{1,2} - \beta_2} \\ \frac{m_{2,1}e^{\tau_{2,1}(\beta_{2,1}^T - \mu_{2,1}^T)}}{\mu_1 + d_1 + m_{2,1} - \beta_1} & 0 \end{pmatrix} \right) \\ &= \sqrt{\frac{m_{1,2}m_{2,1}e^{\tau_{1,2}(\beta_{1,2}^T - \mu_{1,2}^T) + \tau_{2,1}(\beta_{2,1}^T - \mu_{2,1}^T)}}{(\mu_1 + d_1 + m_{2,1} - \beta_1)(\mu_2 + d_2 + m_{1,2} - \beta_2)}}. \end{aligned}$$

Theorem 4.2. *The zero solution of (4.1) is LAS if $\mathcal{T}_2 < 1$ whereas it is unstable if $\mathcal{T}_2 > 1$. A threshold for stability in the system (4.1) is given by $\mathcal{T}_2 = 1$.*

Proof. We use Theorem 2.1 from [22, 23] to establish the relationship between the dominant eigenvalue of $K_2 = T_2(\Sigma_2)^{-1}$ and the number \mathcal{T}_2 . Note that K_2 is an irreducible matrix. By this result, it holds that $\mathcal{T}_2 < 1$ if and only if $\rho(K_2) < 1$, $\mathcal{T}_2 = 1$ if and only if $\rho(K_2) = 1$, and $\mathcal{T}_2 > 1$ if and only if $\rho(K_2) > 1$. Proposition 3.2 completes the proof. \square

We use \mathcal{T}_2 to give stability conditions in terms of the reproduction numbers and movement rates. We assume that $\beta_{1,2}^T - \mu_{1,2}^T > 0$ and $\beta_{2,1}^T - \mu_{2,1}^T > 0$; such conditions are reasonable as evidence ([5, 27]) supports that the transmission rate of an infectious disease can be much higher than usual when a large number of passengers are sharing the same cabin during travel. Furthermore, we consider $\beta_{1,2}^T - \mu_{1,2}^T = \beta_{2,1}^T - \mu_{2,1}^T =: \beta^T - \mu^T$ that enables us to give critical values for the delays. We define

$$\tau_c := \frac{1}{\beta^T - \mu^T} \log \frac{(\mu_1 + d_1 + m_{2,1} - \beta_1)(\mu_2 + d_2 + m_{1,2} - \beta_2)}{m_{1,2}m_{2,1}},$$

and note that $\tau_c > 0$ if and only if the following condition holds:

$$H(m_{1,2}, m_{2,1}) := \frac{(\mu_1 + d_1 + m_{2,1} - \beta_1)(\mu_2 + d_2 + m_{1,2} - \beta_2)}{m_{1,2}m_{2,1}} > 1. \quad (c)$$

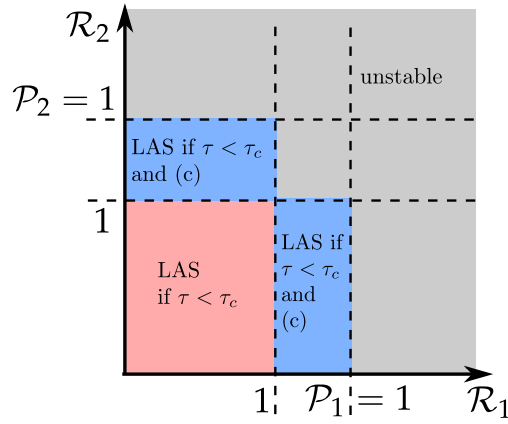


Figure 4.1: Stability of the infection subsystem (4.1).

Theorem 4.3. Assume that $\mathcal{P}_1 < 1$ and $\mathcal{P}_2 < 1$ hold. If $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 > 1$ then the zero solution of (4.1) is unstable for all choices of delays. If $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 < 1$ then the zero solution is LAS for $\tau_{1,2} + \tau_{2,1} < \tau_c$ whereas it is unstable for $\tau_{1,2} + \tau_{2,1} > \tau_c$. If $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 < 1$, or if $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 > 1$, then the zero solution is unstable for all choices of delays if the condition (c) does not hold, it is also unstable if (c) holds and $\tau_{1,2} + \tau_{2,1} > \tau_c$, and it is LAS if (c) holds and $\tau_{1,2} + \tau_{2,1} < \tau_c$.

Proof. If $\mathcal{R}_1 > 1$ then $\mu_1 + d_1 + m_{2,1} - \beta_1 < m_{2,1}$ hence $m_{2,1}/(\mu_1 + d_1 + m_{2,1} - \beta_1) > 1$ holds. Similarly, $\mathcal{R}_2 > 1$ yields $m_{1,2}/(\mu_2 + d_2 + m_{1,2} - \beta_2) > 1$, so it follows that

$$\frac{m_{1,2}m_{2,1}}{(\mu_1 + d_1 + m_{2,1} - \beta_1)(\mu_2 + d_2 + m_{1,2} - \beta_2)} > 1.$$

The assumption $\beta^T - \mu^T > 0$ implies that the square root of the left hand side above is a lower bound for \mathcal{T}_2 , therefore $\mathcal{T}_2 > 1$ for all $\tau_{1,2}$ and $\tau_{2,1}$, that is, the zero solution is unstable in the case when $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 > 1$.

If $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 < 1$ then by similar arguments as in the previous case, we derive that

$$\begin{aligned} \frac{m_{1,2}m_{2,1}}{(\mu_1 + d_1 + m_{2,1} - \beta_1)(\mu_2 + d_2 + m_{1,2} - \beta_2)} &< 1, \\ \iff \\ \sqrt{\frac{m_{1,2}m_{2,1}}{(\mu_1 + d_1 + m_{2,1} - \beta_1)(\mu_2 + d_2 + m_{1,2} - \beta_2)}} &< 1. \end{aligned}$$

The last inequality is equivalent to $\mathcal{T}_2 < 1$ for $\tau_{1,2} = 0$, $\tau_{2,1} = 0$, that is, the zero solution is stable in the model without delays. \mathcal{T}_2 is monotonically increasing in the delays, and the condition $\mathcal{T}_2 < 1$ for stability is equivalent to

$$e^{(\beta^T - \mu^T)(\tau_{1,2} + \tau_{2,1})} < \frac{(\mu_1 + d_1 + m_{2,1} - \beta_1)(\mu_2 + d_2 + m_{1,2} - \beta_2)}{m_{1,2}m_{2,1}} \Leftrightarrow \tau_{1,2} + \tau_{2,1} < \tau_c.$$

It can also be shown that $\mathcal{T}_2 > 1$ is equivalent to $\tau_{1,2} + \tau_{2,1} > \tau_c$ and $\mathcal{T}_2 = 1$ is equivalent to $\tau_{1,2} + \tau_{2,1} = \tau_c$, that is, τ_c gives stability threshold for the delays. Note that $\mu_1 + d_1 + m_{2,1} - \beta_1 > m_{2,1}$ and $\mu_2 + d_2 + m_{1,2} - \beta_2 > m_{1,2}$ by $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 < 1$, respectively, therefore τ_c is positive.

In the two cases when $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 > 1$, and when $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 < 1$, the derivation of the stability condition $\mathcal{T}_2 < 1 \Leftrightarrow \tau_{1,2} + \tau_{2,1} < \tau_c$ goes analogously as above; however τ_c might be non-positive in which case the zero solution is unstable for all choices of delays. A necessary and sufficient condition for $\tau_c > 0$ is that (c) holds, and in this case the zero solution is stable for $\tau_{1,2} + \tau_{2,1} < \tau_c$ but it loses its stability when the sum of the delays exceeds τ_c . \square

4.2 Ring of cities

For a movement network where all movement rates are zero except

$$m_{1,p} > 0, m_{2,1} > 0, \dots, m_{p,p-1} > 0,$$

the system (3.6) reduces to

$$\begin{aligned} \frac{d}{dt} z_1(t) &= (\beta_1 - \mu_1 - d_1 - m_{2,1})z_1(t) + m_{1,p}e^{\tau_{1,p}(\beta_{1,p}^T - \mu_{1,p}^T)}z_p(t) \\ \frac{d}{dt} z_2(t) &= (\beta_2 - \mu_2 - d_2 - m_{3,2})z_2(t) + m_{2,1}e^{\tau_{2,1}(\beta_{2,1}^T - \mu_{2,1}^T)}z_1(t), \\ &\vdots \\ \frac{d}{dt} z_j(t) &= (\beta_j - \mu_j - d_j - m_{j+1,j})z_j(t) + m_{j,j-1}e^{\tau_{j,j-1}(\beta_{j,j-1}^T - \mu_{j,j-1}^T)}z_{j-1}(t), \\ &\vdots \\ \frac{d}{dt} z_p(t) &= (\beta_p - \mu_p - d_p - m_{1,p})z_p(t) + m_{p,p-1}e^{\tau_{p,p-1}(\beta_{p,p-1}^T - \mu_{p,p-1}^T)}z_{p-1}(t). \end{aligned} \tag{4.2}$$

Using the short hand notations $c_1 = e^{\tau_{1,p}(\beta_{1,p}^T - \mu_{1,p}^T)}$ and $c_j = e^{\tau_{j,j-1}(\beta_{j,j-1}^T - \mu_{j,j-1}^T)}$ for $j = 2, \dots, p$, we define

$$T_p = \begin{pmatrix} \beta_1 & 0 & 0 & \dots & 0 & m_{1,p}c_1 \\ m_{2,1}c_2 & \beta_2 & 0 & \dots & 0 & 0 \\ 0 & m_{3,2}c_3 & \beta_3 & \dots & 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & 0 & \dots & m_{p,p-1}c_{p-1} & \beta_p \end{pmatrix},$$

$$\Sigma_p = \text{diag}(\mu_1 + d_1 + m_{2,1}, \mu_2 + d_2 + m_{3,2}, \dots, \mu_p + d_p + m_{1,p}),$$

and obtain the matrix of the linear system as $T_p - \Sigma_p$, where T_p is a nonnegative matrix and Σ_p is a non-singular M-matrix. Therefore it is meaningful to define

$$K_p := T_p(\Sigma_p)^{-1} = \begin{pmatrix} \frac{\beta_1}{\mu_1 + d_1 + m_{2,1}} & 0 & 0 & \cdots & 0 & \frac{m_{1,p}c_1}{\mu_p + d_p + m_{1,p}} \\ \frac{m_{2,1}c_2}{\mu_1 + d_1 + m_{2,1}} & \frac{\beta_2}{\mu_2 + d_2 + m_{3,2}} & 0 & \cdots & 0 & 0 \\ 0 & \frac{m_{3,2}c_3}{\mu_2 + d_2 + m_{3,2}} & \frac{\beta_3}{\mu_3 + d_3 + m_{4,3}} & \cdots & 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & 0 & \cdots & \frac{m_{p,p-1}c_{p-1}}{\mu_{p-1} + d_{p-1} + m_{p,p-1}} & \frac{\beta_p}{\mu_p + d_p + m_{1,p}} \end{pmatrix},$$

and similarly as in the two-city case, a relationship between $s(T_p - \Sigma_p)$ and $\rho(K_p)$ can be established by Proposition 3.2, as follows: the zero solution of (4.2) is LAS ($s(T_p - \Sigma_p) < 0$) if and only if $\rho(K_p) < 1$, it is unstable ($s(T_p - \Sigma_p) > 0$) if and only if $\rho(K_p) > 1$, and it holds that $s(T_p - \Sigma_p) = 0$ if and only if $\rho(K_p) = 1$. We introduce the notation

$$\mathcal{R}_j = \frac{\beta_j}{\mu_j + d_j}, \quad 1 \leq j \leq p,$$

$$\mathcal{P}_j = \frac{\beta_j}{\mu_j + d_j + m_{j+1,j}}, \quad 1 \leq j \leq p-1, \quad \mathcal{P}_p = \frac{\beta_p}{\mu_p + d_p + m_{1,p}},$$

and observe that

$$K_p \geq K_p^0 := \text{diag} \left(\frac{\beta_1}{\mu_1 + d_1 + m_{2,1}}, \dots, \frac{\beta_p}{\mu_p + d_p + m_{1,p}} \right).$$

The proof of the following result is omitted due to its similarity to Theorem 4.1.

Theorem 4.4. *If $\max_{1 \leq j \leq p} \mathcal{P}_j \geq 1$ then the zero solution of (4.2) is unstable for all choices of delays.*

If $\mathcal{P}_j < 1$ for all $j = 1, 2, \dots, p$ then $\rho(K_p^0) < 1$ clearly holds, so it is meaningful to define

$$\mathcal{T}_p := \rho((K_p - K_p^0)(\text{Id}_p - K_p^0)^{-1}),$$

that can be explicitly calculated as

$$\mathcal{T}_p = \rho \left(\begin{pmatrix} 0 & 0 & 0 & \cdots & 0 & \frac{m_{1,p}c_1}{\mu_p + d_p + m_{1,p} - \beta_p} \\ \frac{m_{2,1}c_2}{\mu_1 + d_1 + m_{2,1} - \beta_1} & 0 & 0 & \cdots & 0 & 0 \\ 0 & \frac{m_{3,2}c_3}{\mu_2 + d_2 + m_{3,2} - \beta_2} & 0 & \cdots & 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & 0 & \cdots & \frac{m_{p,p-1}c_{p-1}}{\mu_{p-1} + d_{p-1} + m_{p,p-1} - \beta_{p-1}} & 0 \end{pmatrix} \right)$$

$$= \sqrt[p]{\frac{m_{1,p}c_1}{\mu_p + d_p + m_{1,p} - \beta_p} \frac{m_{2,1}c_2}{\mu_1 + d_1 + m_{2,1} - \beta_1} \cdots \frac{m_{p,p-1}c_{p-1}}{\mu_{p-1} + d_{p-1} + m_{p,p-1} - \beta_{p-1}}}.$$

By the assertion of Proposition 3.2 and [22, 23, Theorem 2.1] we obtain that the zero solution is LAS if and only if $\mathcal{T}_p < 1$, and the stability threshold $\mathcal{T}_p = 1$ can be used to derive explicit conditions for the delays. To this end, it is assumed that $\beta_{1,p}^T - \mu_{1,p}^T = \beta_{2,1}^T - \mu_{2,1}^T = \cdots = \beta_{p,p-1}^T - \mu_{p,p-1}^T =: \beta^T - \mu^T$ and $\beta^T - \mu^T > 0$, that implies that $c_j \geq 1$ holds for all j . Let

$$\tau_c^p := \frac{1}{\beta^T - \mu^T} \log \frac{(\mu_p + d_p + m_{1,p} - \beta_p) \prod_{j=1}^{p-1} (\mu_j + d_j + m_{j+1,j} - \beta_j)}{m_{1,p} \prod_{j=1}^{p-1} m_{j+1,j}},$$

$$\check{\tau} := \tau_{1,p} + \sum_{j=1}^{p-1} \tau_{j+1,j},$$

where $\check{\tau}$ gives the sum of all delays, and note that $\tau_c^p > 0$ is equivalent with the following condition:

$$\frac{(\mu_p + d_p + m_{1,p} - \beta_p) \prod_{j=1}^{p-1} (\mu_j + d_j + m_{j+1,j} - \beta_j)}{m_{1,p} \prod_{j=1}^{p-1} m_{j+1,j}} > 1. \quad (\text{c}^*)$$

Theorem 4.5. *If $\mathcal{R}_j > 1$ for all j then the zero solution of (4.2) is unstable for all choices of delays. If $\mathcal{R}_j < 1$ for all j then the zero solution is LAS for $\check{\tau} < \tau_c^p$ whereas it is unstable for $\check{\tau} > \tau_c^p$. If there are j and k such that $\mathcal{R}_j > 1$ and $\mathcal{R}_k < 1$ then the zero solution is unstable for all choices of delays if the condition (c*) does not hold, it is also unstable if (c*) holds and $\check{\tau} > \tau_c^p$, and it is LAS if (c*) holds and $\check{\tau} < \tau_c^p$.*

Proof. If $\mathcal{R}_j > 1$ for all j then it follows by

$$(\mu_p + d_p + m_{1,p} - \beta_p) < m_{1,p}, \quad (\mu_j + d_j + m_{j+1,j} - \beta_j) < m_{j+1,j}, \quad j = 1, \dots, p-1,$$

that

$$\frac{m_{1,p} \prod_{j=1}^{p-1} m_{j+1,j}}{(\mu_p + d_p + m_{1,p} - \beta_p) \prod_{j=1}^{p-1} (\mu_j + d_j + m_{j+1,j} - \beta_j)} > 1,$$

that implies instability for all choices of delays by $\mathcal{T}_p > 1$. One can easily show that the above inequality is reversed if $\mathcal{R}_j < 1$ for all j , and it follows that $\tau_c^p > 0$. Straightforward calculations yield that the zero solution is LAS in the model without delays because $\mathcal{T}_p < 1$ holds, and stability is preserved in the delayed system as long as $\check{\tau} < \tau_c^p$, where τ_c^p is the critical threshold for the total sum of delays, derived from $\mathcal{T}_p = 1$.

The condition (c*) is equivalent to $\mathcal{T}_p < 1$ in the special case when all delays are zero, moreover, \mathcal{T}_p is increasing in the delays hence the condition (c*) is necessary for stability in the case when there are j and k such that $\mathcal{R}_j > 1$ and $\mathcal{R}_k < 1$. The stability threshold τ_c^p for $\check{\tau}$ is calculated using $\mathcal{T}_p = 1$, and (c*) guarantees that $\tau_c^p > 0$. The proof is complete. \square

5 Discussion on the impact of travel on stability

In this last section we return to system (4.1) to comment on the sensitivity of stability in the movement rates. The following computations are useful:

$$\begin{aligned} \frac{dH}{dm_{1,2}} &= - \frac{(\mu_1 + d_1 + m_{2,1} - \beta_1)(\mu_2 + d_2 - \beta_2)}{(m_{1,2})^2 m_{2,1}}, \\ \frac{dH}{dm_{2,1}} &= - \frac{(\mu_2 + d_2 + m_{1,2} - \beta_2)(\mu_1 + d_1 - \beta_1)}{m_{1,2}(m_{2,1})^2}, \\ \frac{d\tau_c}{dm_{1,2}} &= \frac{1}{\beta^T - \mu^T} \frac{1}{H(m_{1,2}, m_{2,1})} \frac{dH}{dm_{1,2}} \\ &= - \frac{1}{\beta^T - \mu^T} \frac{(\mu_2 + d_2 - \beta_2)}{m_{1,2}(\mu_2 + d_2 + m_{1,2} - \beta_2)} \begin{cases} < 0 & \text{if } \mathcal{R}_2 < 1, \\ > 0 & \text{if } \mathcal{R}_2 > 1, \end{cases} \\ \frac{d\tau_c}{dm_{2,1}} &= \frac{1}{\beta^T - \mu^T} \frac{1}{H(m_{1,2}, m_{2,1})} \frac{dH}{dm_{2,1}} \\ &= - \frac{1}{\beta^T - \mu^T} \frac{(\mu_1 + d_1 - \beta_1)}{m_{2,1}(\mu_1 + d_1 + m_{2,1} - \beta_1)} \begin{cases} < 0 & \text{if } \mathcal{R}_1 < 1, \\ > 0 & \text{if } \mathcal{R}_1 > 1. \end{cases} \end{aligned}$$

We learnt from Theorem 4.3 that if $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 < 1$ then for all choices of the movement rates there is a stability threshold $\tau_c > 0$. The above calculations reveal that the derivative of τ_c is negative with respect to both movement parameters, which means that increasing the movement rates shrinks the stability region and therefore it can destabilize the zero solution.

Stability is also possible if the condition (c) holds and either $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 > 1$ or $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 < 1$. We investigate now the former case, and find that τ_c increases in $m_{1,2}$ whereas it decreases in $m_{2,1}$, thereby the two movement rates have opposite impact on τ_c . The first conclusion that we can draw is that encouraging movement from the endemic city 2 to the non-endemic city 1 expands the stability region and helps stabilizing the DFE. On the other hand, increasing the movement rate in the other direction can lead to stability loss; there is a critical value for $m_{2,1}$, given by

$$m_{2,1}^u = \frac{(\mu_1 + d_1 - \beta_1)(\mu_2 + d_2 + m_{1,2} - \beta_2)}{-(\mu_2 + d_2 - \beta_2)}$$

such that τ_c becomes negative when $m_{2,1}$ exceeds $m_{2,1}^u$ (note that the condition (c) ceases to hold). An important observation to make is that $m_{2,1}^u$ depends on the other movement rate $m_{1,2}$. Lastly, we note that the conditions $\mathcal{P}_1 < 1$ and $\mathcal{P}_2 < 1$ in Theorem 4.3 give lower bounds for $m_{2,1}$ and $m_{1,2}$, respectively, as $m_{2,1} \geq 0 > \beta_1 - \mu_1 - d_1$ and $m_{1,2} > \beta_2 - \mu_2 - d_2 > 0$. Summarizing, in the case when $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 > 1$ hold, stability is possible only if the movement rates are chosen such that $m_{1,2} > \beta_2 - \mu_2 - d_2$ and $m_{2,1} < m_{2,1}^u(m_{1,2})$ hold. Similar phenomena are observed in the case when $\mathcal{R}_1 > 1$ and $\mathcal{R}_2 < 1$.

The above analysis revealed some non-trivial dependence of the stability region on the movement rates when $\mathcal{R}_1 < 1$ and $\mathcal{R}_2 > 1$. To further investigate this question, we let $m_{1,2} = m_{2,1} =: m$ and study the dependence of τ_c on the unified movement rate m . In this special case the function H arises as $H(m) = (\mu_1 + d_1 + m - \beta_1)(\mu_2 + d_2 + m - \beta_2)/m^2$, and τ_c is calculated as $\tau_c = \log [(\mu_1 + d_1 + m - \beta_1)(\mu_2 + d_2 + m - \beta_2)/m^2] / (\beta^T - \mu^T)$.

We assume that $\mathcal{R}_1 < 1$, $\mathcal{R}_2 > 1$ and the condition (c) hold, or equivalently $\mu_1 + d_1 - \beta_1 > 0$, $\mu_2 + d_2 - \beta_2 < 0$, and m is chosen such that $H(m) > 1$. To satisfy the last inequality, it follows by

$$H(m) > 1 \quad \Leftrightarrow \quad m(\mu_1 + d_1 - \beta_1 + \mu_2 + d_2 - \beta_2) > -(\mu_1 + d_1 - \beta_1)(\mu_2 + d_2 - \beta_2)$$

that \mathcal{R}_1 and \mathcal{R}_2 must be chosen such that $(\mu_1 + d_1 - \beta_1 + \mu_2 + d_2 - \beta_2) > 0$ holds, and therefore $H(m) > 1$ is equivalent to

$$m > -\frac{(\mu_1 + d_1 - \beta_1)(\mu_2 + d_2 - \beta_2)}{(\mu_1 + d_1 - \beta_1 + \mu_2 + d_2 - \beta_2)} =: m^H.$$

By $\mathcal{P}_2 < 1$, another condition $m > -(\mu_2 + d_2 - \beta_2)$ arises for m , that is weaker than the one derived above from $H(m) > 1$. Straightforward calculations yield

$$\frac{d\tau_c}{dm} = \frac{1}{\beta^T - \mu^T} \left(\frac{\mu_1 + d_1 - \beta_1 + \mu_2 + d_2 - \beta_2 + 2m}{(\mu_1 + d_1 - \beta_1 + m)(\mu_2 + d_2 - \beta_2 + m)} - \frac{2}{m} \right),$$

and we obtain that

$$\begin{aligned} \frac{d\tau_c}{dm} > 0 &\Leftrightarrow 0 > 2(\mu_1 + d_1 - \beta_1)(\mu_2 + d_2 - \beta_2) + m(\mu_1 + d_1 - \beta_1 + \mu_2 + d_2 - \beta_2), \\ &\Leftrightarrow m^u := \frac{-2(\mu_1 + d_1 - \beta_1)(\mu_2 + d_2 - \beta_2)}{(\mu_1 + d_1 - \beta_1 + \mu_2 + d_2 - \beta_2)} > m, \end{aligned}$$

where we used that \mathcal{R}_1 and \mathcal{R}_2 are chosen such that $(\mu_1 + d_1 - \beta_1 + \mu_2 + d_2 - \beta_2) > 0$. It is easy to see that $m^u > m^H > 0$, therefore we arrive at the following conclusions. If \mathcal{R}_1 and \mathcal{R}_2 are chosen such that $(\mu_1 + d_1 - \beta_1 + \mu_2 + d_2 - \beta_2) > 0$ does not hold then the zero solution is unstable for all movement rates and delays. If the last inequality is reversed, then

$$\begin{aligned}\tau_c > 0, \frac{d\tau_c}{dm} > 0 &\Leftrightarrow m^u > m > m^H, \\ \tau_c > 0, \frac{d\tau_c}{dm} = 0 &\Leftrightarrow m = m^u, \\ \tau_c > 0, \frac{d\tau_c}{dm} < 0 &\Leftrightarrow m > m^u,\end{aligned}$$

with words, for all movement rates m such that $m > m^H$ stability is possible for small delays, however increasing m expands and then shrinks the stability region.

Appendix A DFE of the system (2.1)

A DFE of the system (2.1) is a constant function \hat{E} in C^τ , equal to $(0, \dots, 0, S_1, \dots, S_p, R_1, \dots, R_p) \in \mathbb{R}^{3p}$ for all values of its argument for some $S_1, R_1, \dots, S_p, R_p \in \mathbb{R}$. Note that when the system (2.1) is at such an equilibrium then the system (2.2) is at the steady state $(0, m_{j,k}S_k, m_{j,k}R_k)$. It thus follows that at a DFE, $\mathcal{I}_{j,k}^\tau$, $\mathcal{S}_{j,k}^\tau$ and $\mathcal{R}_{j,k}^\tau$ are obtained as

$$\begin{aligned}\mathcal{I}_{j,k}^\tau &= \mathcal{I}_{j,k}(0, S_k, R_k) = 0, \\ \mathcal{S}_{j,k}^\tau &= \mathcal{S}_{j,k}(0, S_k, R_k) = m_{j,k}S_k, \\ \mathcal{R}_{j,k}^\tau &= \mathcal{R}_{j,k}(0, S_k, R_k) = m_{j,k}R_k.\end{aligned}$$

By $\mathcal{I}_{j,k}^\tau + \mathcal{S}_{j,k}^\tau + \mathcal{R}_{j,k}^\tau = m_{j,k}(S_k + R_k) = m_{j,k}N_k$ it follows that the steady state (N_1, \dots, N_p) for the total populations in each city can be calculated by solving the algebraic linear system

$$A_j = (d_j + M_j)N_j - \sum_{\substack{k=1 \\ k \neq j}}^p m_{j,k}N_k, \quad j = 1, \dots, p. \quad (\text{A.1})$$

The coefficient matrix on the right hand side is obtained as $\text{diag}(d_1 + M_1, \dots, d_p + M_p) - \mathcal{M}$, where $\mathcal{M} = (m_{j,k})$ is the movement matrix and $M_j = \sum_{i=1, i \neq j}^p m_{i,j}$. We note that the sum of all non-diagonal elements in column j is M_j , which means that the coefficient matrix is diagonally dominant. This matrix also has the Z-sign pattern: as defined in [6], all entries are non-positive except possibly those in the diagonal. Theorem 5.1 in [6] gives necessary and sufficient conditions for the non-singularity of matrices with the Z-sign pattern; in particular the equivalence of properties 3 and 11 in [6, Theorem 5.1] implies by the diagonal dominant property that the coefficient matrix is invertible. We can thus give the unique solution to the system (A.1), as

$$(\bar{N}_1, \dots, \bar{N}_p)^T = (\text{diag}(d_1 + M_1, \dots, d_p + M_p) - \mathcal{M})^{-1}(A_1, \dots, A_p)^T.$$

The components of the DFE $(0, \dots, 0, S_1, \dots, S_p, R_1, \dots, R_p)$ for the recovered classes arise as the solution of the system

$$0 = (d_j + M_j)R_j - \sum_{\substack{k=1 \\ k \neq j}}^p m_{j,k}R_k, \quad j = 1, \dots, p.$$

Similar arguments to those for the system (A.1) imply that $\bar{R}_1 = 0, \dots, \bar{R}_p = 0$ must hold. It follows that $\bar{S}_1 = \bar{N}_1, \dots, \bar{S}_p = \bar{N}_p$, and therefore there is a unique DFE, given by $(0, \dots, 0, \bar{S}_1, \dots, \bar{S}_p, 0, \dots, 0)$.

Appendix B Calculation of the gradients in the system (3.2)

To derive the gradients on the right hand side of the system (3.2), we recall the definition $(\mathcal{I}_{j,k}(v), \mathcal{S}_{j,k}(v), \mathcal{R}_{j,k}(v)) = \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v))$ from (2.4), and it follows that

$$\begin{pmatrix} \text{grad}(\mathcal{I}_{j,k}(v)) \\ \text{grad}(\mathcal{S}_{j,k}(v)) \\ \text{grad}(\mathcal{R}_{j,k}(v)) \end{pmatrix} = \frac{\partial}{\partial v} \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v)).$$

Proposition B.1. *It holds that*

$$\frac{\partial}{\partial v} \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v)) = e^{\int_0^{\tau_{j,k}} Dg_{j,k}(\tilde{y}_{j,k}(v; 0, h_{j,k}(v))) \, dv} Dh_{j,k}(v).$$

If $h_{j,k}(v)$ is an equilibrium of the system (2.3) then

$$\frac{\partial}{\partial v} \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v)) = e^{\tau_{j,k} Dg_{j,k}(h_{j,k}(v))} Dh_{j,k}(v).$$

Proof. Theorem 3.3 in Chapter I in [8] states that as $g_{j,k}$ has continuous first derivative, the solution $\tilde{y}_{j,k}(\theta; 0, y_*)$ of system (2.3) is continuously differentiable with respect to θ and y_* on its domain of definition. The matrix $(\frac{\partial \tilde{y}_{j,k}(\theta; 0, y_*)}{\partial y_*}) \in \mathbb{R}^{3 \times 3}$ satisfies the linear variational equation

$$\frac{d}{d\theta} Y(\theta) = Dg_{j,k}(\tilde{y}_{j,k}(\theta; 0, y_*)) Y(\theta),$$

where $Y: \mathbb{R} \rightarrow \mathbb{R}^{3 \times 3}$ and $\frac{\partial \tilde{y}_{j,k}(0; 0, y_*)}{\partial y_*} = \text{Id}_3$. By solving the linear variational equation, we derive that $Y(\theta) = e^{\int_0^\theta Dg(\tilde{y}_{j,k}(v; 0, y_*)) \, dv} Y(0)$, that yields

$$\frac{\partial \tilde{y}_{j,k}(\tau_{j,k}; 0, y_*)}{\partial y_*} = e^{\int_0^{\tau_{j,k}} Dg_{j,k}(\tilde{y}_{j,k}(v; 0, y_*)) \, dv} Y(0) \quad (\text{B.1})$$

for $Y(0) = \text{Id}_3$. Hence the derivative of the solution $\tilde{y}_{j,k}$ with respect to $v \in \mathbb{R}^3$ arises as

$$\begin{aligned} \frac{\partial}{\partial v} \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v)) &= \frac{\partial \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v))}{\partial y_*} Dh_{j,k}(v) \\ &= e^{\int_0^{\tau_{j,k}} Dg_{j,k}(\tilde{y}_{j,k}(v; 0, h_{j,k}(v))) \, dv} Dh_{j,k}(v). \end{aligned}$$

If $h_{j,k}(v)$ is an equilibrium of the system (2.3) then $\tilde{y}_{j,k}(v; 0, h_{j,k}(v)) = h_{j,k}(v)$ holds for all $v \in [0, \tau_{j,k}]$, and therefore the above formula reduces to

$$\frac{\partial}{\partial v} \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v)) = e^{\tau_{j,k} Dg_{j,k}(h_{j,k}(v))} Dh_{j,k}(v).$$

□

In order to calculate $\text{grad}(\mathcal{I}_{j,k}(\hat{E}_k))$, $\text{grad}(\mathcal{S}_{j,k}(\hat{E}_k))$ and $\text{grad}(\mathcal{R}_{j,k}(\hat{E}_k))$, we need to evaluate $\frac{\partial}{\partial v} \tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(v))$ at $v = \hat{E}_k$. Note that $h_{j,k}(\hat{E}_k) = m_{j,k}(0, \bar{S}_k, 0)$ is an equilibrium of the system (2.3). First, we obtain that $Dh_{j,k}(v) = m_{j,k} \text{Id}_3$, moreover, $Dg_{j,k}(h_{j,k}(\hat{E}_k))$ is derived as

$$Dg_{j,k}(h_{j,k}(\hat{E}_k)) = \begin{pmatrix} \beta_{j,k}^T - \mu_{j,k}^T & 0 & 0 \\ -\beta_{j,k}^T & 0 & 0 \\ \mu_{j,k}^T & 0 & 0 \end{pmatrix}.$$

We use the short hand notation G for this matrix. Observe that $G^2 = (\beta_{j,k}^T - \mu_{j,k}^T)G$, and $G^n = (\beta_{j,k}^T - \mu_{j,k}^T)^{n-1}G$ by analogy for $n \geq 2$. We deduce that

$$\begin{aligned} e^{\tau_{j,k}G} &= \text{Id}_3 + (\tau_{j,k}G) + \frac{(\tau_{j,k}G)^2}{2!} + \dots + \frac{(\tau_{j,k}G)^n}{n!} + \dots \\ &= \text{Id}_3 + (\tau_{j,k}G) + \frac{\tau_{j,k}^2(\beta_{j,k}^T - \mu_{j,k}^T)G}{2!} + \dots + \frac{\tau_{j,k}^n(\beta_{j,k}^T - \mu_{j,k}^T)^{n-1}G}{n!} + \dots \\ &= \text{Id}_3 + \frac{G}{(\beta_{j,k}^T - \mu_{j,k}^T)} \left(-1 + 1 + \tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T) + \frac{\tau_{j,k}^2(\beta_{j,k}^T - \mu_{j,k}^T)^2}{2!} + \dots \right. \\ &\quad \left. + \frac{\tau_{j,k}^n(\beta_{j,k}^T - \mu_{j,k}^T)^n}{n!} + \dots \right) \\ &= \text{Id}_3 + \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} G \\ &= \begin{pmatrix} e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} & 0 & 0 \\ -\beta_{j,k}^T \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} & 1 & 0 \\ \mu_{j,k}^T \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} & 0 & 1 \end{pmatrix}. \end{aligned}$$

Using the definition of $\tilde{y}_{j,k}(\tau_{j,k}; 0, h_{j,k}(0, \bar{S}_k, 0))$ and Proposition B.1 we obtain that

$$\begin{pmatrix} \text{grad}(\mathcal{I}_{j,k}(\hat{E}_k)) \\ \text{grad}(\mathcal{S}_{j,k}(\hat{E}_k)) \\ \text{grad}(\mathcal{R}_{j,k}(\hat{E}_k)) \end{pmatrix} = m_{j,k} \begin{pmatrix} e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} & 0 & 0 \\ -\beta_{j,k}^T \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} & 1 & 0 \\ \mu_{j,k}^T \frac{e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} - 1}{(\beta_{j,k}^T - \mu_{j,k}^T)} & 0 & 1 \end{pmatrix}.$$

Appendix C Cooperative and irreducible FDE

We recall some definitions and theorems from [24]. Let $n \in \mathbb{Z}_+$ and $r_1, \dots, r_n > 0$. For $r > 0$, we denote by $C([-r, 0], \mathbb{R})$ the space of continuous functions on $[-r, 0]$, which is a Banach space with the usual uniform norm $|\phi| = \sup\{\phi(\theta) : -r \leq \theta \leq 0\}$. Let

$$C_r = \prod_{i=1}^n C([-r_i, 0], \mathbb{R}),$$

that is a Banach space with the norm $|\phi| = \sum |\phi_i|$. We denote the generic element of C_r by $\phi = (\phi_1, \dots, \phi_n)$, and by $L(C_r, \mathbb{R}^n)$ the space of bounded linear maps from C_r to \mathbb{R}^n .

Consider the general nonautonomous linear system

$$\frac{d}{dt}z(t) = L(t)z_t, \quad (\text{C.1})$$

where $z: \mathbb{R} \rightarrow \mathbb{R}^n$, $L: \mathbb{R} \rightarrow L(C_r, \mathbb{R}^n)$ is continuous. For the segment of the solution we use the usual notation z_t where $z_t = (z_1^t, \dots, z_n^t)$, $z_i^t = z_i(t + \theta)$ for $\theta \in [-r_i, 0]$. Let $L_i(t)\phi$ denote the i -th component of $L(t)\phi$. We introduce the following definition for the property (K).

Definition C.1. (K) Whenever $\phi \geq 0$ and $\phi_i(0) = 0$ holds for some i , then $L_i(t)\phi \geq 0$.

Lemma C.2. (K) holds if and only if there exists $a_i(t) \in \mathbb{R}$ for $1 \leq i \leq n$ and positive Borel measures $\eta_{ij}(t)$ for $1 \leq i, j \leq n$ on $[-r_j, 0]$ such that $L(t)$ has a representation in terms of Borel measures, as

$$L_i(t)\phi = a_i(t)\phi_i(0) + \sum_{j=1}^n \int_{-r_j}^0 \phi_j(\theta) d\theta \eta_{ij}(t, \theta)$$

and $\eta_{ii}(t)\{0\} = 0$. Moreover, if (K) holds then the representation is unique and $a_i(t)$ and $\eta_{ij}(t)$ are continuous functions of t .

For $1 \leq i \leq n$ let e_i denote the standard basis vectors for \mathbb{R}^n , and let $\hat{e}_i \in C_r$ be the constant function equal to e_i for all values of its argument. We need two further conditions.

Definition C.3. (R) For each j for which $r_j > 0$, there exists i such that for all t , $\eta_{ij}(t)([-r_j, -r_j + \epsilon]) > 0$ for all small $\epsilon > 0$.

Definition C.4. (I) The matrix $A(L)(t)$ defined by

$$A(L)(t) = \text{col}(L(t)\hat{e}_1, \dots, L(t)\hat{e}_n)$$

is irreducible.

If (K) holds then by Lemma C.2 we obtain that

$$A(L)_{ij}(t) = L_i(t)\hat{e}_j = a_i(t)(e_j)_i + \eta_{ij}(t)([-r_j, 0]) = \begin{cases} a_i(t) + \eta_{ii}(t)([-r_i, 0]) & \text{if } i = j, \\ \eta_{ij}(t)([-r_j, 0]) & \text{if } i \neq j. \end{cases}$$

The condition (K) implies that if a component of a solution of the differential equation gets turned on, then it stays on. We need the condition (R) to ensure that some component actually gets turned on if the initial condition is nontrivial. (I) is a kind of irreducibility assumption; it implies that once one component gets turned on then all components eventually get turned on.

A general autonomous functional differential equation (FDE) is denoted by

$$\frac{d}{dt}z(t) = f(z_t), \quad (\text{C.2})$$

where $f: U \rightarrow \mathbb{R}^n$ is continuously differentiable and U is an open subset of C_r . For $\psi \in U$, $Df(\psi)$ gives a bounded linear operator, that is, $Df(\psi)\phi \in \mathbb{R}^n$ for $\phi \in C_r$, and the corresponding linear system arises as

$$\frac{d}{dt}z(t) = Df(\psi)z_t.$$

For any $\phi, \psi \in C_r$ we say that $\phi \leq \psi$ ($\phi < \psi$) holds if and only if $\phi(s) \leq \psi(s)$ ($\phi(s) < \psi(s)$) holds in \mathbb{R}^n for every $s \in \prod[-r_i, 0]$.

Definition C.5. For a set X , let $u, v \in X$, $u < v$. Then the order interval generated by u, v is defined as $[u, v] \equiv \{x \in X : u \leq x \leq v\}$. The set X is said to be order convex if $[u, v] \subset X$ whenever $u, v \in X$ satisfy $u < v$. In particular, for a subset U of C_r , $[\phi_1, \phi_2] \equiv \{\psi \in U : \phi_1 \leq \psi \leq \phi_2\}$ where $\phi_1, \phi_2 \in U$, and U is order convex if $[\phi_1, \phi_2] \subset U$ whenever $\phi_1, \phi_2 \in U$ satisfy $\phi_1 < \phi_2$.

Definition C.6. The FDE is cooperative if U is order convex and if $Df(\psi)$ satisfies (K) for each $\psi \in U$.

If $Df(\psi)$ satisfies (K) then as a consequence of Lemma C.2, $df(\psi)$ can be represented as in Lemma C.2, that is, $a_i = a_i(\psi)$ and $\eta_{ij} = \eta_{ij}(\psi)$ exist such that for $\phi \in U$

$$(Df(\psi))\phi = a_i\phi_i(0) + \sum_{j=1}^n \int_{-r_j}^0 \phi_j(\theta) d\theta \eta_{ij}(\theta).$$

Definition C.7. The FDE is cooperative and irreducible if it is cooperative and the following hold:

- (i) $Df(\psi)$ satisfies (I) for each $\psi \in U$;
- (ii) for each j such that $r_j > 0$, there exists i such that for all $\psi \in U$, $\eta_{ij}(\psi)([-r_j, -r_j + \epsilon]) > 0$ for all small $\epsilon > 0$, where $\eta_{ij}(\psi)$ is the Borel measure uniquely defined for $Df(\psi)$.

The equilibria of the autonomous FDE (C.2) are those $\phi \in U$ such that $z_t(\phi) = \phi$ for all $t \geq 0$. Evaluating this equality at $\theta = 0$ gives $z(t) = \phi(0)$ for all $t \geq 0$, hence ϕ must be a constant function and $f(\phi) = 0$. For $v \in \mathbb{R}^n$ we denote by \hat{v} an element of C_r that satisfies $\hat{v}(\theta) \equiv v$, and \hat{v} is an equilibrium of (C.2) if and only if $\hat{v} \in U$ and $f(\hat{v}) = 0$. To investigate the stability of an equilibrium \hat{v} , by the principle of linearized stability it is sufficient to study the stability of the trivial equilibrium in the linear variational system

$$\frac{d}{dt}z(t) = Lz_t, \tag{C.3}$$

where $L = Df(\hat{v}) \in L(C_r, \mathbb{R}^n)$. Note that (C.3) is a special case of the general nonautonomous linear system (C.1). The stability of the trivial solution of the linear variational system (3.1) is determined by the characteristic equation $\Delta(\lambda) = 0$, which arises by seeking solutions of the form $z(t) = e^{\lambda t}u$ where $u \in \mathbb{R}^n$. More precisely, if the stability modulus, defined as

$$s(L) = \max\{\operatorname{Re} \lambda : \Delta(\lambda) = 0\},$$

is negative then the trivial solution of (C.3) is locally asymptotically stable (LAS) whereas it is unstable if $s(L) > 0$.

Assume that the nonlinear FDE (C.2) is cooperative and irreducible. Then a cooperative and irreducible system of ordinary differential equations (ODE) can be associated with (C.2), by ignoring the delays which appear in (C.2):

$$\frac{d}{dt}z(t) = F(z), \quad F(z) = f(\hat{z}), \quad F: \mathbb{R}^n \rightarrow \mathbb{R}^n. \tag{C.4}$$

The ODE (C.4) has the same equilibria as the FDE (C.2), more precisely, v is an equilibrium of (C.4) if and only if \hat{v} is an equilibrium of (C.2). We quote here Corollary 5.2 of Chapter 5, Section 5 of [24]:

Corollary C.8. $s(L) < 0$ ($s(L) > 0$) if and only if $s(DF(v)) < 0$ ($s(DF(v)) > 0$).

Summarizing, the stability of an equilibrium \hat{v} of the cooperative and irreducible FDE (C.2) can be investigated by determining the stability of the equilibrium v in the linear approximation of the associated ODE (C.4). The stability properties of the linear systems extend to the nonlinear equations by the principle of linearized stability.

Appendix D The system (3.5) is cooperative and irreducible

For the theory of cooperative and irreducible functional-differential equations we refer to the Appendix C, where we recalled some definitions and results from [24].

The appropriate domain for (3.5) is $D = \prod_{j=1}^p C([- \sigma_j, 0], \mathbb{R}_+)$, and we recall that σ_j is defined as $\sigma_j = \max\{\tau_{k,j} : 1 \leq k \leq p\}$ for $j = 1, \dots, p$. We obtain system (3.5) in the FDE form as

$$\frac{d}{dt}z(t) = \mathcal{B}z_t,$$

where $\mathcal{B} \in L(D, \mathbb{R}^p)$ and

$$(\mathcal{B}\phi)_j = (\beta_j - \mu_j - d_j - M_j)\phi_j(0) + \sum_{\substack{k=1 \\ k \neq j}}^p m_{j,k} e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} \phi_k(-\tau_{j,k})$$

for $1 \leq j \leq p$. It follows from $m_{j,k} e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} \geq 0$ and the definition of \mathcal{B} that the condition (K) holds, that is, the system is cooperative. A unique representation of $(\mathcal{B}\phi)_j$ in terms of Borel measures arises, as

$$(\mathcal{B}\phi)_j = a_j \phi_j(0) + \sum_{\substack{j,k=1 \\ j \neq k}}^p \int_{-\tau_{j,k}}^0 \phi_k(-\tau_{j,k}) d\eta_{j,k}(\theta)$$

with $a_j = (\beta_j - \mu_j - d_j - M_j)$ and $\eta_{j,k}(\theta) \equiv m_{j,k} e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)}$.

For every k such that $\sigma_k > 0$ there is a j such that $\sigma_k = \tau_{j,k}$, hence the condition (R) follows from $\eta_{j,k}([- \tau_{j,k}, -\tau_{j,k} + \epsilon]) = m_{j,k} e^{\tau_{j,k}(\beta_{j,k}^T - \mu_{j,k}^T)} > 0$ for $0 < \epsilon < \tau_{j,k}$. To check the condition (I), note that

$$\text{col}(\mathcal{B}\hat{e}_1, \dots, \mathcal{B}\hat{e}_p) = B^0 + \sum_{\substack{j,k=1 \\ j \neq k}}^p B^{j,k}.$$

It follows from the irreducibility of \mathcal{M} that $B^0 + \sum_{j,k=1, j \neq k}^p B^{j,k}$ is irreducible and hence (I) is satisfied. The conditions (R) and (I) ensure that the system (3.5) is irreducible, which completes the proof.

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References

- [1] J. ARINO, C. SUN, W. YANG, Revisiting a two-patch SIS model with infection during transport, *Math. Med. Biol.* **33**(2015), No. 1, 29–55. [MR3483090](#); [url](#)
- [2] J. ARINO, P. VAN DEN DRIESSCHE, Disease spread in metapopulations, in: *Fields Inst. Commun.*, Vol. 48., Amer. Math. Soc., Providence, RI, 2006, 1–12. [MR2223345](#)
- [3] J. CUI, Y. TAKEUCHI, Y. SAITO, Spreading disease with transport-related infection, *J. Theoret. Biol.* **239**(2006), No. 3, 376–390. [MR2224621](#); [url](#)
- [4] O. DIEKMANN, J. A. P. HEESTERBEEK, M. G. ROBERT, The construction of next-generation matrices for compartmental epidemic models, *J. R. Soc. Interface* **7**(2010), 873–885. [url](#)
- [5] EUROPEAN CENTRE FOR DISEASE PREVENTION AND CONTROL, Risk assessment guidelines for infectious diseases transmitted on aircraft, http://ecdc.europa.eu/en/publications/Publications/0906_TER_Risk_Assessment_Guidelines_for_Infectious_Diseases_Transmitted_on_Aircraft.pdf, 2009.
- [6] M. FIEDLER, *Special matrices and their applications in numerical mathematics*, Martinus Nijhoff Publishers, Dordrecht, The Netherlands, 1986. [MR1105955](#)
- [7] D. GAO, S. RUAN, A multipatch malaria model with logistic growth populations, *SIAM J. Appl. Math.* **72**(2012), No. 3, 819–841. [MR2968751](#); [url](#)
- [8] J. K. HALE, *Ordinary differential equations*, Robert E. Krieger Publishing Co., Inc., Huntington, N.Y., 1980. [MR0587488](#)
- [9] J. HOFBAUER, J. SO, Diagonal dominance and harmless off-diagonal delays, *Proc. Amer. Math. Soc.* **128**(2000), No. 9, 2675–2682. [MR1707519](#)
- [10] Y. H. HSIEH, P. VAN DEN DRIESSCHE, L. WANG, Impact of travel between patches for spatial spread of disease, *Bull. Math. Biol.* **69**(2007), No. 4, 1355–1375. [MR2312219](#); [url](#)
- [11] D. H. KNIPL, Fundamental properties of differential equations with dynamically defined delayed feedback, *Electron. J. Qual. Theory Differ. Equ.* **2013**, No. 17, 1–18. [MR3033802](#); [url](#)
- [12] D. H. KNIPL, G. RÖST, Multiregional SIR model with infection during transportation *Biomath* **1**(2012), No. 1, 1209255, 7 pp. [MR3105481](#); [url](#)
- [13] D. H. KNIPL, G. RÖST, J. WU, Epidemic spread and variation of peak times in connected regions due to travel related infections—dynamics of an anti-gravity type delay differential model, *SIAM J. Appl. Dyn. Syst.* **12**(2013), No. 4, 1722–1762. [MR3116637](#); [url](#)
- [14] J. LI, X. ZOU, Dynamics of an epidemic model with non-local infections for diseases with latency over a patchy environment, *J. Math. Biol.* **60**(2010), No. 5, 645–686. [MR2606520](#); [url](#)
- [15] J. LIU, J. WU, Y. ZHOU, Modeling disease spread via transport-related infection by a delay differential equation, *Rocky Mountain J. Math.* **38**(2008), No. 5, 1525–1540. [MR2457374](#); [url](#)
- [16] X. LIU, Y. TAKEUCHI, Spread of disease with transport-related infection and entry screening, *J. Theoret. Biol.* **242**(2006), No. 2, 517–528. [MR2272571](#); [url](#)

- [17] L. Q. MAI, H. F. WERTHEIM, T. N. DUONG, H. R. VAN DOORN, N. T. HIEN, P. HORBY, A community cluster of oseltamivir-resistant cases of 2009 H1N1 influenza, *N. Engl. J. Med.* **362**(2010), No. 1, 86–87. [url](#)
- [18] Y. NAKATA, On the global stability of a delayed epidemic model with transport-related infection, *Nonlinear Anal. Real World Appl.* **12**(2011), No. 6, 3028–3034. [MR2832945](#); [url](#)
- [19] Y. NAKATA, G. RÖST, Global dynamics of a delay differential system of a two-patch SIS-model with transport-related infections, *Math. Bohem.* **140**(2015), No. 2, 171–193. [MR3368492](#)
- [20] Y. NAKATA, G. RÖST, Global analysis for spread of infectious diseases via transportation networks, *J. Math. Biol.* **70**(2015), No. 6, 1411–1456. [MR3323601](#); [url](#)
- [21] S. RUAN, W. WANG, S. A. LEVIN, The effect of global travel on the spread of SARS, *Math. Biosci. Eng.* **3**(2006), No. 1, 205–218. [MR2192135](#); [url](#)
- [22] Z. SHUAL, J. A. P. HEESTERBEEK, P. VAN DEN DRIESSCHE, Extending the type reproduction number to infectious disease control targeting contacts between types, *J. Math. Biol.* **67**(2013), No. 5, 1067–1082. [MR3111982](#); [url](#)
- [23] Z. SHUAL, J. A. P. HEESTERBEEK, P. VAN DEN DRIESSCHE, Erratum to: Extending the type reproduction number to infectious disease control targeting contacts between types, *J. Math. Biol.* **71**(1)(2015), 1–3. [MR3345968](#); [url](#)
- [24] H. L. SMITH, *Monotone dynamical systems. An introduction to the theory of competitive and cooperative systems*, AMS, Providence, RI, 1995. [MR1319817](#)
- [25] Y. TAKEUCHI, X. LIU, J. CUI, Global dynamics of SIS models with transport-related infection, *J. Math. Anal. Appl.* **329**(2007), No. 2, 1460–1471. [MR2296936](#); [url](#)
- [26] P. VAN DEN DRIESSCHE, J. WATMOUGH, Reproduction numbers and sub-threshold endemic equilibria for compartmental models of disease transmission, *Math. Biosci.* **180**(2002), No. 1, 29–48. [MR1950747](#); [url](#)
- [27] B. WAGNER, B. COBURN, S. BLOWER, Calculating the potential for within-flight transmission of influenza A (H1N1), *BMC Med.* **7**(2009), No. 81, 7 pp. [url](#)
- [28] W. WANG, Epidemic models with population dispersal, in: Y. Takeuchi, Y. Iwasa, K. Sato (eds.), *Mathematics for life science and medicine*, Biological and Medical Physics, Biomedical Engineering, Springer, Berlin, 2007, pp. 67–95. [MR2309364](#)
- [29] W. WANG, X. ZHAO, An age-structured epidemic model in a patchy environment, *SIAM J. Appl. Math.* **65**(2005), No. 5, 1597–1614. [MR2177716](#); [url](#)