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3 **A new method for choosing the computational cell in stochastic**  
4 **reaction-diffusion systems**

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**Abstract.** How to choose the computational compartment or cell size for the stochastic simulation of a reaction-diffusion system is still an open problem, and a number of criteria have been suggested. A generalized measure of the noise for finite-dimensional systems based on the largest eigenvalue of the covariance matrix of the number of molecules of all species has been suggested as a measure of the overall fluctuations in a multivariate system, and we apply it here to a discretized reaction-diffusion system. We show that for a broad class of first-order reaction networks this measure converges to the square root of the reciprocal of the smallest mean species number in a compartment at the steady state. We show that a suitably re-normalized measure stabilizes as the volume of a cell approaches zero, which leads to a criterion for the maximum volume of the compartments in a computational grid. We then derive a new criterion based on the sensitivity of the entire network, not just of the fastest step, that predicts a grid size that assures that the concentrations of all species converge to a spatially-uniform solution. This criterion applies for all orders of reactions and for reaction rate functions derived from singular perturbation or other reduction methods, and encompasses both diffusing and non-diffusing species. We show that this predicts the maximal volume found in the linear problems, and we illustrate our results with an example motivated by anterior-posterior pattern formation in *Drosophila*, and with several other examples.

## 1. Introduction

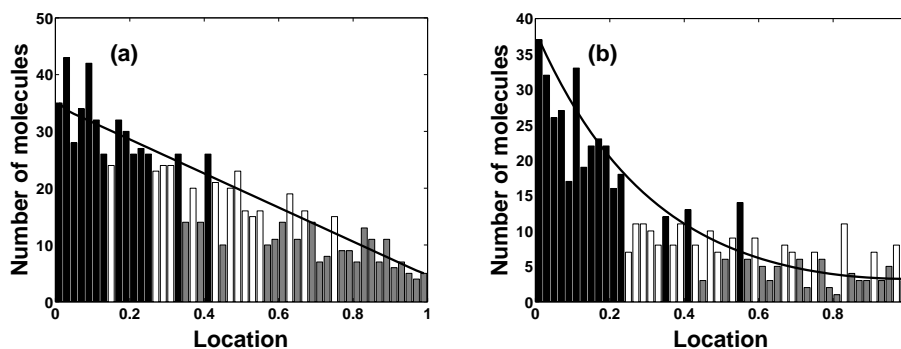
It is now widely-recognized that stochastic effects can play an important role in diverse processes such as gene expression and spatial pattern formation in development because many key biological molecules are present in low copy numbers. For example, gene transcription in some bacteria involves interactions between 1-3 promoter elements, 10-20 polymerase holoenzyme units, 10-20 copies of repressor proteins, 3000 RNA polymerase molecules, and approximately 1000 ribosomes [26]. Since chemical reactions occur in discrete steps at the molecular level, the processes are inherently stochastic and the inherent “irreproducibility” in these dynamics has been demonstrated experimentally for single cell gene expression events [43, 36, 29]. In some contexts stochastic effects simply add noise to an output, but have no beneficial role, but in others, such as asymmetric cell division, their role is essential. In general organisms show a remarkable degree of resilience or robustness in the face of noise, and thus understanding the time-dependent behavior of a system of interacting species and how noise influences the outcome is important in numerous contexts, including temporal gene expression profiles, signal transduction, and other biochemical processes. Of course when the numbers of molecules of all species are large enough, a ‘law of large numbers’ argument shows that for finite times the stochastic formulation described later converges to the mass-action based deterministic description commonly used, at least in well-mixed systems [25].

One of the earliest investigations of stochastic effects in reactions is due to Delbrück [13], who studied the distribution of the number of molecules for a single reacting species in a one-component, enzyme-catalyzed system. It was assumed there that the substrate is in excess and thus the process is effectively first-order. There are many other examples of first-order reaction networks that involve a small number of molecules, including transcription and translation modeled as first-order catalytic reactions [46], for which stochastic analysis is necessary. The evolution of the surface morphology during epitaxial growth involves the nucleation and growth of atomic islands, and these processes may be described by first-order adsorption and desorption reactions coupled with diffusion along the surface. Proteins exist in distinct conformational states, and the reversible transitions between states can be described as first-order conversion processes [30]. Fluctuating protein conformations are important in the movement of small molecules through proteins such as myoglobin; hence it is important to understand the distribution of these states [20, 4]. RNA also exists in several conformations, and the transitions between various folding states follow first-order kinetics [9]. In Gadgil *et. al.* [17] the linear problem for an arbitrary number of components is more or less completely solved, in that it is shown there how to obtain the evolution equations for the mean and variance in closed form. These results also address the problem of understanding how the interplay between the nature of the individual steps and the connectivity or topology of the entire network affects the dynamics of the system, irrespective of whether a deterministic or a stochastic description is the most appropriate, but this problem remains unsolved for general nonlinear reaction schemes.

In the context of biological pattern formation, robustness or resilience is frequently defined with respect to the precision and sensitivity of the determination of boundaries between different cell types in a developing tissue [48]. A classical paradigm for this process is the French flag model, in which a one-dimensional domain is to be divided into three equal-size sub-domains [48]. In the simplest deterministic version of this model, either specialized source and

1 sink cells located at the boundary of the developmental field maintain the concentration of a signaling molecule, called  
 2 a morphogen, at appropriate fixed levels, or boundary cells produce the morphogen at a fixed rate. In the former case,  
 3 when there is no degradation of the morphogen in the interior of the domain a linear distribution can be established  
 4 in a one-dimensional system of about 1 mm in length in the time that is normally available for commitment to  
 5 differentiation [52,11]. Given fixed thresholds between different cell types, the tissue can be proportioned into any  
 6 number of cell types and the determination of the boundaries is robust with respect to changes in the size of the system.  
 7 In the latter case we consider a simple version in which the flux at the boundary is specified, and degradation by  
 8 first-order reaction occurs throughout the domain - generalizations will be discussed later in the context of *Drosophila*  
 9 patterning.

10 However a deterministic description of either of these models ignores the possible effects of stochastic fluctuations  
 11 in the signaling and gene control networks and the effect stochastic fluctuations may have on the precision of pattern-  
 12 ing. Figure 1(a) shows one realization of a stochastic model of a linear chain of compartments with fixed numbers of  
 13 molecules at the endpoints of the domain, and Figure 1(b) shows one realization for the second scheme, in which the  
 14 input flux is fixed at the left. In both panels the solid line shows the mean of the distribution, which can be computed  
 15 directly since the equations are linear [17]. These curves also represent the steady-state distribution for the corre-  
 sponding deterministic system. Since each developing embryo represents one realization of the stochastic patterning



**Fig. 1.** Two examples of the French flag model with stochastic dynamics. In each case the system is  $1 \times 0.01 \times 0.01 \text{ mm}^3$ , the x-length is discretized into 50 compartments, the diffusion coefficient is  $1000 \mu\text{m}^2 \text{ min}^{-1}$ , and one realization is shown at 100 mins. **(a)** The distribution for fixed concentrations at the boundaries: 35 molecules at the left and 5 at the right, with diffusion and no degradation throughout. The color indicates the thresholds: black – greater than 25 and gray – greater than 15 molecules. **(b)** The distribution when there is an influx from a source at the left end at a rate  $0.1 \text{ nM } \mu\text{m} \text{ min}^{-1}$ , and diffusion and degradation at a rate  $0.01 \text{ min}^{-1}$  throughout. Initially, each compartment has 10 molecules of morphogen. The color indicates the thresholds: black – greater than 12 and gray – greater than 6 molecules.

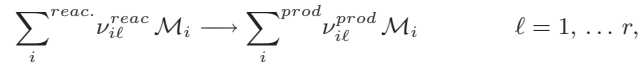
16 process, the results illustrate the difficulty in determining the location of the boundaries between cell types in the face  
 17 of such fluctuations. In embryonic patterning in *Drosophila*, the primary morphogen Dpp has signaling activity in the  
 18 range of  $10^{-10} \text{ M}$  to  $10^{-9} \text{ M}$  [39], and at these concentrations there are very few Dpp signaling molecules available to  
 19 the receptors. Thus fluctuations will be significant, and how the embryos cope with such noise to pattern reliably is  
 20 still not fully understood. Preliminary work has shown that a postulated positive feedback mechanism [38] increases  
 21 the reliability of spatial patterning [53]. In any case, spatially-distributed systems add a new level of complexity to  
 22 the problem of understanding the importance of noise in development.  
 23

### 24 1.1. The description of reaction networks

25 Throughout we deal with reacting systems that are not diffusion-limited, and thus a stochastic analysis and simulation  
 26 of reaction and diffusion can be built around a discretized or compartmentalized model of the system in which each  
 27 compartment is well-mixed. We discuss the rationale for this approach and the method for choosing the compartment  
 28 size in more detail later, but we begin by developing the chemical master equation that can be used to describe  
 29 such systems. We begin with some background on a deterministic description of reacting systems, then derive the

1 master equation for an arbitrary network of reacting species, and finally add diffusion. The formulation of the reaction  
 2 component is described in more detail elsewhere [33, 17, 28].

Suppose that a reacting mixture contains a set  $\mathcal{M}$  of  $s$  chemical species  $\mathcal{M}_i$  that participate in a total of  $r$  reactions. Let  $\nu_{i\ell}$  be the stoichiometric coefficient of the  $i^{\text{th}}$  species in the  $\ell^{\text{th}}$  reaction. The  $\nu_{i\ell}$  are non-negative integers that represent the normalized molar proportions of the species in a reaction. Each reaction is written in the form



where the sums are over reactants and products, respectively in the  $\ell^{\text{th}}$  reaction. Here the forward and reverse reaction of a reversible pair are treated as two irreversible reactions. There may be other species that do not react, but they play no role here. Once the reactants and products are specified the network topology of the associated reaction graph is defined. The linear combinations of species that appear as reactants or products in the various elementary steps are called complexes, and the relation defined by which complexes are connected by reaction gives rise to a directed graph  $\mathcal{G}$  in which each complex is identified with a vertex  $V_j$  in  $\mathcal{G}$  and a directed edge  $E_\ell$  is introduced into  $\mathcal{G}$  for each reaction. Each edge carries a nonnegative weight  $\hat{\mathcal{R}}_\ell(c)$  given by the intrinsic rate of the corresponding reaction. The topology of  $\mathcal{G}$  is in turn represented in its vertex-edge incidence matrix  $\mathcal{E}$ , which is defined as follows.

$$\mathcal{E}_{j\ell} = \begin{cases} +1 & \text{if } E_\ell \text{ is incident at } V_j \text{ and is directed toward it} \\ -1 & \text{if } E_\ell \text{ is incident at } V_j \text{ and is directed away from it} \\ 0 & \text{otherwise} \end{cases}$$

3 If there are  $p$  complexes and  $r$  reactions, then  $\mathcal{E}$  has  $p$  rows and  $r$  columns and every column has exactly one +1 and  
 4 one -1.

5 Once the complexes and reactions are fixed, the stoichiometry of the complexes is determined, and we let  $\nu$  denote  
 6 the  $s \times p$  matrix whose  $j^{\text{th}}$  column encodes the stoichiometric amounts of the reacting species in the  $j^{\text{th}}$  complex.  
 7 Then the deterministic temporal evolution of the composition of a spatially-uniform reacting mixture is governed by

$$\frac{dc}{dt} = \nu \mathcal{E} \hat{\mathcal{R}}(c). \quad (1)$$

A special but important class of rate functions is that in which the rate of the  $\ell^{\text{th}}$  reaction can be written as

$$\hat{\mathcal{R}}_\ell(c) = k_\ell R_\ell(c).$$

9 This includes ideal mass action rate laws, in which the rate is given by

$$R_\ell = \prod_{i=1}^n (c_i)^{\nu_{i\ell}}. \quad (2)$$

11 As it stands, (1) includes all reacting species, but those whose concentration is constant on the time scale of  
 12 interest can be deleted from each of the complexes in which it appears, and its concentration or mole fraction can be  
 13 absorbed into the rate constant for reactions in which it participates as a reactant<sup>1</sup>. Furthermore, some complexes  
 14 may not comprise any time-dependent species; these will be called null complexes and denoted by  $\mathcal{M}^0$ , and each  
 15 null complex gives rise to a column of zeroes in  $\nu$ . The rate of any reaction in which the reactant complex is a null  
 16 complex is usually constant. For instance, any transport reaction of the form  $\mathcal{M}^0 \rightarrow \mathcal{M}_i$  introduces a null complex  
 17 and the corresponding flux of  $\mathcal{M}_i$  represents a constant input to the reaction network, provided that the rate of the  
 18 transport step does not depend on the concentration of a time-dependent species.

One can also base the description of a reacting system on the number of molecules, and to connect the deterministic and stochastic descriptions, we let  $n = (n_1, n_2, \dots, n_s)$  denote the discrete composition vector whose  $i^{\text{th}}$  component  $n_i$  is the number of molecules of species  $\mathcal{M}_i$  present in the volume  $\mathbf{V}$ . This is the discrete version of the composition vector  $c$ , and they are related by  $n = \mathcal{N}_A \mathbf{V} c$ , where  $\mathcal{N}_A$  is Avogadro's number. From (1) we obtain the deterministic

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<sup>1</sup> Hereafter  $s$  will denote the number of species whose concentration may be time-dependent.

evolution for  $n$  as

$$\frac{dn}{dt} = \nu \mathcal{E} \tilde{\mathcal{R}}(n)$$

where  $\tilde{\mathcal{R}}(n) \equiv \mathcal{N}_A \mathbf{V} \hat{\mathcal{R}}(n / \mathcal{N}_A \mathbf{V})$ . In particular, for mass-action kinetics

$$\tilde{\mathcal{R}}_\ell(n) = \mathcal{N}_A \mathbf{V} k_\ell \mathcal{R}_\ell(n / \mathcal{N}_A \mathbf{V}) = \mathcal{N}_A \mathbf{V} k_\ell \prod_{i=1}^s \left( \frac{n_i}{\mathcal{N}_A \mathbf{V}} \right)^{\nu_{i\ell}} = \frac{k_\ell}{(\mathcal{N}_A \mathbf{V})^{\sum_i \nu_{i\ell} - 1}} \prod_{i=1}^s (n_i)^{\nu_{i\ell}} = \hat{k}_\ell \prod_{i=1}^s (n_i)^{\nu_{i\ell}}.$$

- 1 Care is needed in accounting for volume factors when one species in a bimolecular reaction is confined to a surface  
2 and the other to the adjacent fluid, as occurs in receptor-ligand interactions.

### 3 1.2. The stochastic description of reactions

In the stochastic description the evolution of the number of molecules of a species is modeled as a continuous-time Markov jump process. Let  $N_i(t)$  be a random variable that represents the number of molecules of species  $\mathcal{M}_i$  at time  $t$ , and let  $N$  denote the vector of  $N_i$ s. The state of the system at any time is a point in  $\mathcal{Z}_0^s$ , where  $\mathcal{Z}_0$  is the set of nonnegative integers. Let  $P(n, t)$  be the joint probability that  $N(t) = n$ , i.e.,  $N_1 = n_1, N_2 = n_2, \dots, N_s = n_s$ ; then the master equation for the evolution of  $P$  is

$$\frac{d}{dt} P(n, t) = \sum_{m \in \mathcal{S}(n)} \mathcal{R}(m, n) \cdot P(m, t) - \sum_{m \in \mathcal{T}(n)} \mathcal{R}(n, m) \cdot P(n, t)$$

- 4 where  $\mathcal{R}(i, j)$  is the probability per unit time of a transition from state  $i$  to state  $j$ , the ‘source’ set  $\mathcal{S}(n)$  is the set  
5 of all states that can terminate at  $n$  after one reaction step, and  $\mathcal{T}(n)$ , the ‘target’ set, is the set all states reachable  
6 from  $n$  in one step of the feasible reactions. The sets  $\mathcal{S}(n)$  and  $\mathcal{T}(n)$  are easily determined using the underlying graph  
7 structure. It follows from the definition of  $\nu$  and  $\mathcal{E}$  that the  $\ell^{\text{th}}$  reaction  $C(k) \rightarrow C(k')$  induces a change  $\Delta n^{(\ell)} = \nu \mathcal{E}_{(\ell)}$   
8 in the number of molecules of all species after one step of the reaction, where subscript  $\ell$  denotes the  $\ell^{\text{th}}$  column of  $\mathcal{E}$   
9 and  $C(k)$  denotes the  $k^{\text{th}}$  complex of species. Therefore the state  $m = n - \nu \mathcal{E}_{(\ell)}$  is a source or predecessor to  $n$  under  
10 one step of the  $\ell^{\text{th}}$  reaction. Similarly, states of the form  $m = n + \nu \mathcal{E}_{(\ell)}$  are reachable from  $n$  in one step of the  $\ell^{\text{th}}$   
11 reaction.

Once the graph of the network and the stoichiometry are fixed, we can sum over reactions rather than sources and targets, and consequently the master equation takes the form

$$\frac{d}{dt} P(n, t) = \sum_{\ell} \mathcal{R}_\ell(n - \nu \mathcal{E}_{(\ell)}) \cdot P(n - \nu \mathcal{E}_{(\ell)}, t) - \sum_{\ell} \mathcal{R}_\ell(n) \cdot P(n, t).$$

However, the transition probabilities  $\mathcal{R}_\ell(n)$  are not simply the macroscopic rates  $\tilde{\mathcal{R}}$  if the reactions are second-order (or higher), but are given by [19,17]

$$\mathcal{R}_\ell = c_{j(\ell)} h_{j(\ell)}(n).$$

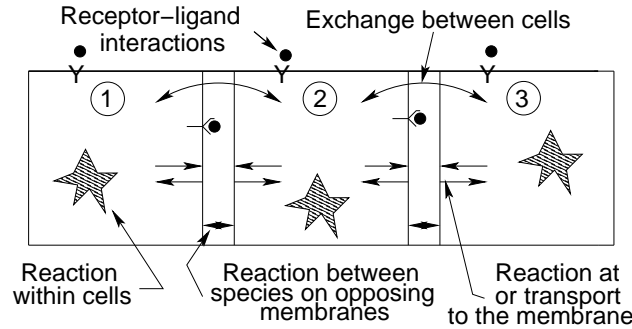
Here  $j(\ell)$  denotes the reactant complex for the  $\ell^{\text{th}}$  reaction, the factor  $c_{j(\ell)}$  is the probability per unit time that the molecular species in the  $j^{\text{th}}$  complex react via the  $\ell^{\text{th}}$  reaction, and  $h_{j(\ell)}(n)$  is the number of independent combinations of the molecular components in this complex. Thus

$$c_{j(\ell)} = \frac{k_\ell}{(\mathcal{N}_A \mathbf{V})^{\sum_i \nu_{ij(\ell)} - 1}} = \hat{k}_\ell \quad \text{and} \quad h_{j(\ell)} = \prod_i \prod_{m_i=0}^{\nu_{ij(\ell)} - 1} (n_i - m_i).$$

- 12 For a first-order reaction  $\hat{k}_\ell$  is the probability per unit time per molecule of a transition, for a bimolecular step of the  
13 form  $A + A \rightarrow C$  it represents the probability per unit time of a transition per ordered AA pair, and for a bimolecular  
14 step of the form  $A + B \rightarrow C$  it represents the probability per unit time of a transition per AB pair. For a first-order  
15 reaction  $1/\hat{k}_\ell$  represents a *bona fide* time scale, as in the deterministic case, whereas for bimolecular reactions it does  
16 not. We return to this point later.

1 1.3. The master equation for a general system of reaction and diffusion

2 Many biological problems involve more than just one well-mixed compartment, and this provides the motivation  
 3 for describing a general framework for treating networks of cells or compartments that can communicate by various  
 4 mechanisms, some of which involve material transport, and others of which merely involve signaling by, for instance, a  
 5 ligand binding to a receptor. A schematic of an array of discrete cells that interact by several mechanisms is shown in  
 6 Figure 2. An example of a system that can be described by such a network is the developing wing disc in *Drosophila*  
 7 *melanogaster*. We illustrate the types of interactions within and between cells, but there may also be transport by  
 8 diffusion within the gaps between cells and in the surrounding fluid. We allow reaction within cells, reaction between  
 9 species at the membranes with either species in the associated cell or on the membrane of the adjacent cell, and  
 transport between cells.



**Fig. 2.** A schematic of a 1D network of coupled cells in which cells can interact either at their opposing membranes or via transport between cells.

10

If we associate a graph with only one class of nodes to the array, then ordering them so that the first  $N_c$  are cell labels and the remaining  $2N_c$  (or smaller, depending on boundary conditions) are pairs of opposed membranes produces the cleanest separation, for then the incidence matrix has the form

$$\mathcal{E} = \begin{bmatrix} \mathcal{E}_{cc} & \mathcal{E}_{cm} \\ \mathcal{E}_{mc} & \mathcal{E}_{mm} \end{bmatrix}$$

11 where  $\mathcal{E}_{cc}$ ,  $\mathcal{E}_{cm}$ ,  $\mathcal{E}_{mc}$ , and  $\mathcal{E}_{mm}$  denote the cell-cell, the cell-membrane, the membrane-cell and the membrane-membrane  
 12 connections, respectively.

13 It is not difficult to formulate a master equation for the general network in Figure 2, since all the steps shown can  
 14 be treated as chemical reactions, but the details depend on the specific steps involved. When there is no interaction  
 15 between membranes the matrix collapses to  $\mathcal{E}_{cc}$ , which encodes the connectivity of the cell network. It is clear that  
 16 we can still allow receptor-ligand interactions, and examples of this are given later. In the remainder of the section we  
 17 assume that transport is only by diffusion and that the network arises from the discretization of a reaction-diffusion  
 18 problem.

19 In a stochastic description of reaction-diffusion systems in which the reactions are not diffusion-limited, the  
 20 domain  $\Omega$  containing the mixture can be discretized into spatially-uniform compartments, and diffusion can be  
 21 treated as a jump process between compartments. How to correctly choose the computational cell or compartment  
 22 size is a significant problem, and we deal with this in the following sections – here we simply derive the master  
 23 equation for a reaction-diffusion system. In fact the compartments need not be computational cells that arise from  
 24 the discretization of a domain; they could be compartments in the wider sense used above, as long a transport is  
 25 diffusion-like, and we describe all cases as compartments hereafter. However, for simplicity we assume that transport  
 26 between compartments is symmetric and linear in the concentration differences, so as to describe diffusion, and that  
 27 all compartments are cubes of side-length  $h$ . We will comment later on how to generalize the resulting equation for  
 28 other transport mechanisms and unequal compartment sizes.

1 Earlier we introduced the reaction graph  $\mathcal{G}$  and we indicated above that one can associate a graph with the cellular  
 2 network as well. To distinguish between these we denote the former as  $\mathcal{G}_r$  and the latter as  $\mathcal{G}_c$ . We suppose that there  
 3 are  $N_c$  nodes in  $\mathcal{G}_c$ , we define the incidence matrix  $\mathcal{E}_c$ , which was  $\mathcal{E}_{cc}$  in the general case, and the discrete Laplacian  
 4 as  $\Delta_c = -\mathcal{E}_c \mathcal{E}_c^T / 2$ . When the network stems from a regular grid, the Laplacian is the discretization, modulo a scale  
 5 factor of  $h^{-2}$ , of the spatial Laplacian [35].

Let  $N(t) = (N^1(t), N^2(t), \dots, N^{N_c}(t))$  be the vector of random variables whose  $k^{th}$  vector component represents  
 the numbers of molecules of species in the  $k^{th}$  compartment. Let  $P(n, t)$  be the probability that  $N(t) = n$ , *i.e.*,  
 the joint probability that  $N^1(t) = n^1 = (n_1^1, n_2^1, \dots, n_s^1)$ ,  $N^2(t) = n^2 = (n_1^2, n_2^2, \dots, n_s^2)$ ,  $\dots$ ,  $N^{N_c}(t) = n^{N_c} =$   
 $(n_1^{N_c}, n_2^{N_c}, \dots, n_s^{N_c})$ . Then  $P(n, t) \equiv P(n^1, n^2, \dots, n^k, \dots, n^{N_c}, t)$ , and in the absence of coupling between the com-  
 partments the master equation for  $P(n, t)$  is

$$\frac{d}{dt}P(n, t) = \sum_{k=1}^{N_c} \left[ \sum_{\ell} \mathcal{R}_{\ell}(n^k - \nu \mathcal{E}_{(\ell)}) \cdot P(n^1, \dots, n^k - \nu \mathcal{E}_{(\ell)}, \dots, n^{N_c}, t) - \sum_{\ell} \mathcal{R}_{\ell}(n^k) \cdot P(n^k, t) \right]$$

6 since this just involves the sum over all compartments of the changes in state of the individual compartments. Since  
 7 the compartments evolve independently in the absence of coupling, the joint distribution can be factored into a  
 8 product of  $N_c$  distributions, but this plays no role here.

In the absence of reaction but in the presence of diffusive coupling between compartments, the flux of the  $i^{th}$   
 species from  $k$  to one of its neighbors  $k'$  is assumed to be given by

$$J_i^{kk'} = \frac{D_i}{h^2} n_i^k,$$

while the reverse flux is

$$J_i^{k'k} = \frac{D_i}{h^2} (n_i^{k'} + 1).$$

The transfer of one molecule in the first step involves the change  $n_i^k \rightarrow n_i^k - 1$ ,  $n_i^{k'} \rightarrow n_i^{k'} + 1$  whereas the reverse  
 flux involves the change  $n_i^{k'} + 1 \rightarrow n_i^{k'}$ ,  $n_i^k - 1 \rightarrow n_i^k$ . Of course both steps conserve the particle number. Thus the  
 evolution equation for  $P(n, t)$  when diffusion alone is considered is

$$\frac{d}{dt}P(n, t) = \sum_{k=1}^{N_c} \sum_{k' \in \mathcal{N}(k)} \left[ \sum_{i=1}^s \frac{D_i}{h^2} (n_i^{k'} + 1) P(n^1, \dots, n^k - e_i, \dots, n^{k'} + e_i, \dots, n^{N_c}, t) - \frac{D_i}{h^2} n_i^k P(n^k, t) \right]$$

9 where  $\mathcal{N}(k)$  is the set of all neighbors of  $k$  in  $\mathcal{G}_c$  and  $e_i = (0, 0, \dots, 1, \dots, 0)^T$  has a 1 in the  $i^{th}$  position and zeroes  
 10 elsewhere. To obtain the full equation we simply add the reaction and diffusion contributions; thus

$$\begin{aligned} 11 \quad \frac{d}{dt}P(n, t) = \sum_{k=1}^{N_c} \left\{ \sum_{k' \in \mathcal{N}(k)} \left[ \sum_{i=1}^s \frac{D_i}{h^2} (n_i^{k'} + 1) P(n^1, \dots, n^k - e_i, \dots, n^{k'} + e_i, \dots, n^{N_c}, t) - \frac{D_i}{h^2} n_i^k P(n^k, t) \right] \right. \\ 12 \quad \left. + \sum_{\ell} \mathcal{R}_{\ell}(n^k - \nu \mathcal{E}_{(\ell)}) \cdot P(n^1, \dots, n^k - \nu \mathcal{E}_{(\ell)}, \dots, n^{N_c}, t) - \sum_{\ell} \mathcal{R}_{\ell}(n^k) \cdot P(n^k, t) \right\}. \end{aligned} \quad (3)$$

14 The formulation at (3) is based on the assumption that the domain is decomposed into equal-size compartments  
 15 defined by a Cartesian grid. However this is not necessary, and a general formulation based on the compartment graph  
 16 that allows for unequal volumes goes as follows. A first step is to generalize the foregoing to arbitrary topologies,  
 17 albeit with equal-size compartments. This can be done using the Laplacian and the adjacency matrix  $A$ , which is  
 18 defined via  $\Delta = -d(\Delta) + A$ , where  $d(\Delta)$  is the diagonal matrix whose  $k^{th}$  entry is the degree of the  $k^{th}$  node. In  
 19 the second step one incorporates unequal volumes of compartments and differences in the area for transfer between  
 20 compartments. The latter is equivalent to allowing the diffusion coefficients to depend on the pair of compartments  
 21 involved in the exchange. Finally, one has to scale the reaction rates differently in different compartments to account  
 22 for the fact that the volumes of the compartments are not equal. We leave the details to the interested reader.

1 *1.4. Overview of the paper*

2 If the compartment size used to develop the master equation (3) arises as a basic unit in the system, for example, a  
 3 biological cell size, then no further analysis is needed, and a stochastic simulation based on (3) using the Gillespie's  
 4 algorithm [19] or one of its many modifications [14] is appropriate. However, if the system is described initially as a  
 5 continuum and a reaction-diffusion equation of the form

$$6 \quad \frac{\partial u(\mathbf{x}, t)}{\partial t} = \mathcal{D}\Delta u(\mathbf{x}, t) + \mathcal{R}(u(\mathbf{x}, t)) \quad (4)$$

7 is used, then the conversion to the master equation (3) requires a choice of compartment size. A numerical algorithm  
 8 for the solution of the deterministic equation (4) would involve a discretization of space, and finer discretizations would  
 9 produce more accurate solutions under suitable conditions. However this assumes that the solution is continuous in  
 10  $\mathbf{x}$ , but this is clearly not true when there are few molecules of any species present. One expects that for a fixed total  
 11 number of molecules in a system, smaller compartments will produce larger variation in the number of molecules  
 12 within a compartment, and the following example makes this precise.

Consider a closed system containing  $N$  molecules distributed in  $N_c$  compartments that are connected by diffusion.  
 The steady-state distribution is spatially uniform, and it is known that it is multinomial [17] with mean and variance  
 given by

$$M_i = \frac{N}{N_c} \quad \sigma_i^2 = M_i \left(1 - \frac{1}{N_c}\right) = \frac{N}{N_c} \left(1 - \frac{1}{N_c}\right).$$

13 Therefore, if we adopt the coefficient of variation,  $CV = \sigma_i/M_i$  as a suitable measure of the noise, then one has

$$14 \quad CV = \sqrt{\frac{N_c - 1}{N}}, \quad (5)$$

15 which is zero for  $N_c = 1$  and which grows as  $\sqrt{N_c}$  for large  $N_c$ . Thus choosing a very small compartment size leads to  
 16 large fluctuations, as measured by the  $CV$ , in the amounts in various compartments. When reactions also occur the  
 17 interaction between reaction and diffusion must be taken into account, and this will be done in later sections. Here we  
 18 introduce some of the limitations of a compartmental analysis and then discuss previous work aimed at determining  
 19 a suitable compartment size.

20 One problem that presents difficulties, both for a continuum description such as (4), as well as for the master  
 21 equation approach, arises when one or more reactions are diffusion-limited. This applies only for bimolecular and  
 22 higher-order (should they ever occur) reactions, and refers to reactions of the type  $A + A \rightarrow P$  or  $A + B \rightarrow P$  in  
 23 which the reaction occurs instantaneously when the reactants are in sufficiently close proximity. Thus formation of  
 24 homo- and heterodimers, polymerization reactions, ligand-receptor interactions, and enzyme-catalyzed reactions are  
 25 all potentially diffusion-limited. In this context neither (4) nor the conventional framework of a compartmentalized  
 26 system can be used directly, and both require some modifications. The classical work of Smoluchowski [42] dealt with  
 27 coagulation reactions, but since then it has been extended by many others (for a review see [5]). Suppose that the  
 28 molecules are assigned a radius  $r_A$  and  $r_B$ , respectively, and assume that the molecules react with a probability  $k_0$   
 29 when the distance between their centers is  $r_A + r_B$ . In a coordinate frame in which  $B$  is fixed, the concentration of  
 30  $A$  satisfies

$$31 \quad \frac{\partial c}{\partial t} = \frac{D}{r^2} \frac{\partial}{\partial r} \left( r^2 \frac{\partial c}{\partial r} \right) \quad \text{for } r \in (r_A + r_B, \infty)$$

$$32 \quad 4\pi r^2 D \frac{\partial c}{\partial r} = k_0 c \quad \text{at } r = r_A + r_B$$

$$33 \quad \lim_{r \rightarrow \infty} c(r) = c_0.$$

In the strictly diffusion-limited case  $k_0$  is infinite, and one finds that as  $t \rightarrow \infty$ , the effective reaction rate  $k_e$  reduces  
 to

$$k_e = 4\pi(D_A + D_B)(r_A + r_B) = 4\pi DR$$



where  $D = D_A + D_B$  and  $R = r_A + r_B$  [5]. The units of  $k_e$  appear formally to be volume/time, but this is the rate per molecule of  $B$ , and therefore the units of  $k_e$  are

$$\left( \frac{\text{molecule of } B}{\text{volume}} \text{time} \right)^{-1}$$

1 as is necessary for a bimolecular rate constant. An associated time scale for a diffusion-limited reaction can be defined  
2 as

$$3 \quad \tau_{dl} = (k_e \cdot c^*)^{-1} \quad (6)$$

4 where  $c^*$  is a characteristic concentration of  $B$  in molecules/unit volume.

5 An estimate obtained via a stochastic analysis begins with the problem of computing the mean first passage time  
6 for a random walker searching for a specified target. Suppose that the walker is confined to a spherical shell of inner  
7 radius  $r_0$  and outer radius  $r_1$ , that it cannot escape through the outer boundary and is annihilated upon hitting  
8 the inner boundary. The mean first passage time  $\tau(r)$  for annihilation beginning on a spherical surface of radius  
9  $r \in (r_0, r_1)$  is given by the solution of

$$10 \quad \frac{D}{r^2} \frac{d}{dr} \left( r^2 \frac{d\tau}{dr} \right) = -1$$

$$11 \quad \tau(r_0) = 0$$

$$12 \quad \tau'(r_1) = 0. \quad (7)$$

One finds that the solution is

$$\tau(r) = \frac{1}{6D} \left( r_0^2 - r^2 + 2r_1^3 \left( \frac{1}{r_0} - \frac{1}{r} \right) \right)$$

and the average of this over the spherical shell is

$$\overline{\tau(r)} = \frac{r_1^2}{2D(1-\xi^3)} \left[ -\frac{2\xi^5}{15} + \frac{2\xi^2}{3} + \frac{2}{3\xi} - \frac{6}{5} \right]$$

where  $\xi \equiv r_0/r_1$ . If this ratio is sufficiently small, then to leading order

$$\overline{\tau(r)} = \frac{r_1^2}{3D} \frac{r_1}{r_0} = \frac{V_s}{4\pi D r_0}$$

13 where  $V_s$  is, to lowest order, the volume of the spherical domain. By identifying  $r_0$  with  $r_A + r_B$  one sees that the  
14 continuum and stochastic approaches agree to within the choice of a reference concentration in (6). When there are  
15  $N$  non-interacting walkers in the shell, the spatially averaged mean first passage time remains unchanged. A proof of  
16 this is given in Appendix A.

To estimate the magnitude of  $k_e$  and hence the time scale in a typical solvent, we use the Stokes-Einstein relation

$$D = \frac{k_B T}{6\pi\mu r'}$$

to estimate the diffusion coefficient, where  $k_B$  is Boltzmann's constant,  $T$  is the absolute temperature,  $\mu$  is the solvent  
viscosity, and  $r'$  is the hydrodynamic radius of the molecule. Assuming a solvent viscosity  $\mu = 9$  poise, one finds that  
 $k_B T/\mu = 4.5 \cdot 10^{-12} \text{ cm}^3/\text{sec}$ , and if one assumes both molecules have the same encounter radius and a hydrodynamic  
radius equal to that radius, then

$$k_e = \frac{8}{3} \frac{k_B T}{\mu} = 1.2 \cdot 10^{-11} \text{ cm}^3/\text{sec}$$

or in molar units

$$k_e = 7.2 \cdot 10^9 \text{ M}^{-1} \text{ sec}^{-1}.$$

17 Based on this estimate, there appear to be few biochemical reactions that are severely diffusion-limited *in vitro*, since  
18 most of the available second-order rate constants lie in the range of  $10^4 - 10^7 (\text{M sec})^{-1}$  (*cf.* Table 1). It may be that  
19 *in vivo* measurements will show more diffusion influence on reaction rates.

**Table 1.** On- and off-rates for second-order reactions

Receptor	Ligand	$k_f$ ( $M^{-1} sec^{-1}$ )	$k_r$ ( $sec^{-1}$ )	Ref
Insulin	Insulin	$1.6 \times 10^5$	$3.3 \times 10^{-3}$	[27]
EGF	EGF	$3 \times 10^6$	$2 \times 10^{-3}$	[27]
Fibronectin	Fibronectin	$1.17 \times 10^4$	$1 \times 10^{-2}$	[27]
Surface bound protein	BMP	$1.67 \times 10^7$	$3.33 \times 10^{-2}$	[49]
Type I/II BMP receptor	BMP	$4 \times 10^5$	$6.67 \times 10^{-2}$	[49]
Vkg	Dpp	$3.92 \times 10^3$	$2.90 \times 10^{-3}$	[50]
Dcg1	Dpp	$3.20 \times 10^3$	$2.07 \times 10^{-3}$	[50]
Human Collagen IV	BMP4	$2.75 \times 10^4$	$2.50 \times 10^{-3}$	[50]
Cv-2	BMP	$1.67 \times 10^6$	$3.33 \times 10^{-3}$	[38]
Receptor	BMP	$4.0 \times 10^5$	$4.0 \times 10^{-4}$	[38]
Receptor	BMP bound Cv-2	$3.3 \times 10^7$	$3.3 \times 10^{-2}$	[38]
Cv-2	BMP bound Receptor	$3.3 \times 10^5$	$3.3 \times 10^{-2}$	[38]

When reactions are strictly diffusion-limited, rather than just diffusion-influenced, a precise description involves tracking individual particles as they diffuse in space, and algorithms for this have been developed [2,14]. If one describes the motion of the Brownian particles A and B with stochastic differential equations, then in certain regimes the positions evolve according to

$$dx_i = \sqrt{2D_i}dW_i \quad i = A, B,$$

where  $W_i$  are independent standard Wiener processes. This is a more fundamental approach that also indicates a shortcoming in the previous analysis, because the previous analysis assumes that the motion of the two particles is perfectly correlated, which is not true if each executes an independent Brownian motion.

Several different attempts to correctly treat diffusion-limited reactions via a compartment-based master equation have been made. Fange *et al.*, [16] first derive a discretized description using spherical shells around a single particle, and then lift this description to a compartmental model using modified diffusion kinetic coefficients that reflect the discretization. However, in deriving the modified rates the authors use a model in which the diffusing particle is confined to a spherical shell around the reaction site, and thus implicitly assume that the particles will react with probability one. In 3D this is not strictly true, and in general it is an approximation that may be difficult to justify. Others have addressed the relationship between the continuum description (4) and the master equation for a diffusion-limited reaction [21]. It is shown that one cannot expect that the limit of the master equation is (4), since bimolecular reactions disappear in this limit. In [22] a pseudo-potential to capture the singular behavior is used, while the reaction rates in a compartmentalized model are modified so that the encounter probability of the molecules does not depend on the compartment size [15]. A more fundamental approach is to derive the evolution equation for the pair distribution function, as was done in a continuum description [45], but the discretization problem was not addressed there.

Numerous authors have addressed the issue of how to choose a suitable compartment size  $h$  when reactions are not diffusion-limited. Most criteria are based on the premise that the compartment size should be small enough that all mobile species in a compartment can traverse the compartment on the time scale of the fastest reaction in a compartment, since only then a compartment can be considered spatially uniform. For bimolecular reactions this implies that all pairs of reactants have equal probability of reacting. Thus most criteria hinge on the relation between the diffusion time scale for a chosen compartment size and a characteristic reaction time. As noted earlier, reaction rates scale as  $V^p \sim h^{pd}$  where  $p = 1, 0, -1$ , for zero-, first- and second-order reactions, respectively, and  $d$  is the space

dimension. Thus a characteristic reaction time scales with the compartment size as follows.

$$\tau_r \sim \begin{cases} \frac{c^*}{k_0 h^d} & : \text{ zero-order} \\ \frac{1}{k_1} & : \text{ first-order} \\ \frac{h^d}{k_2 c^*} & : \text{ second-order} \end{cases}$$

1 Here  $k_j$  is a concentration-based reference kinetic coefficient, usually the deterministic value, and  $c^*$  is a reference  
2 concentration.

3 A characteristic diffusion time scale is given by  $\tau_d = h^2 / \min_i D_i$  and thus the following conditions on  $h$

4 
$$h^{2+d} < \frac{\min_i D_i}{k_0 (c^*)^{-1}} : \text{ zero-order}$$

5 
$$h^2 < \frac{\min_i D_i}{k_1} : \text{ first-order}$$

6 
$$h^{2-d} < \frac{\min_i D_i}{k_2 c^*} : \text{ second-order}$$

7 will ensure that  $\tau_d \leq \tau_r$ . Since no molecular interaction are involved in zero- and first-order reactions, one might  
8 conclude that the first two estimates are irrelevant. However, they arise naturally in the criterion for convergence to  
9 spatial uniformity derived later, and they indicate the crossover in  $h$  from a regime in which diffusion is fast compared  
10 to reaction to one in which the reverse holds. Understanding these regimes plays a role in stochastic simulations using  
11 Gillespie's method, because if the compartment is too small the diffusion steps dominate the computations. A more  
12 detailed analysis using the known solutions for the mean and variance in systems of first-order reactions [17], which  
13 is in progress, will lead to a more precise characterization of this.

14 Stundzia and Lumsden [44] describe diffusion as a jump process, but rather than using a constant jump rate, they  
15 use the inverse of the mean first passage time from a compartment to its neighbors. Baras and Mansour [6] suggest  
16 that the compartment size should be smaller than a two-particle correlation length, but larger than a mean free path  
17  $\lambda$ , defined as the average distance traveled by a particle between two reactive collisions. Since the correlation length  
18 is smaller than the mean free path in dilute solutions, they assert that the compartment size should be chosen with  
19 the order of mean free path *i.e.* ,

20 
$$h \approx \mathcal{O}(\lambda).$$

21 Since this only applies for gases we do not consider it further here. Bernstein [7] applies Gillespie's algorithm to  
22 simulate diffusion with spatially-inhomogeneous coefficients in non-uniform Cartesian grids using a finite volume  
23 approximation with either Neumann or Dirichlet boundary conditions. To apply Gillespie's algorithm to reaction-  
24 diffusion networks the slowest diffusion time should be much less than the fastest reaction time, *i.e.* ,  $\tau_d \ll \tau_r$ .

25 Isaacson and Peskin [23] suggest a lower and upper bound for the compartment size based on three facts: the  
26 compartment size should be much larger than the mean free path  $\lambda$  so that the system can be considered in local  
27 equilibrium in each compartment due to nonreactive collisions; the compartment size should be much smaller than  
28 the length scale of the system size  $L_x$ ; and the time scale for diffusion through each compartment should be faster  
29 than the time scale for the fastest bimolecular reaction. This leads to the condition

30 
$$\max\left(\lambda, \frac{k}{D}\right) \ll h \ll L_x \tag{8}$$

31 but as we noted earlier, a reference concentration is needed to properly define a reaction time scale for bimolecular  
32 reactions.

33 Erban and Chapman [15] suggest simulation algorithms for reaction-diffusion processes in which diffusion is treated  
34 as a jump process. They give a simple example of a bimolecular reaction for which the stationary distributions are

1 explicitly known, and point out the limitations of current SSAs by comparing the stationary distributions. Using a  
 2 modified algorithm, they show that if the compartment size is larger than a critical value, the multi-compartment  
 3 model can reproduce the known stationary distribution correctly. The reactions considered are  $\emptyset \rightarrow A$  and  $A+B \xrightarrow{k} B$   
 4 with diffusing species,  $A$  and  $B$ , and  $k$  is the deterministic reaction rate of the bimolecular reaction. Then the lower  
 5 bound on the compartment size is

$$6 \quad h \geq h_{crit}$$

$$7 \quad h_{crit} \approx \frac{k}{4(D_A + D_B)}.$$

Table 2: Previous criteria for the compartment size

Reference	Criteria
Bernstein	$\tau_D \approx \frac{h^2}{2dD} \ll \tau_c$
Isaacson and Peskin	$\max\left(\lambda, \frac{k}{D}\right) \ll h \ll L_x$
Erban and Chapman	$\frac{k}{4(D_A + D_B)} \leq h$

8  
 9 All of these estimates implicitly assume that all reacting species also diffuse, but this is rarely the case in biological  
 10 problems, since diffusible species may bind to receptors or other essentially immobile proteins. In addition, none of the  
 11 criteria deal with the possibility of reactions on the boundary of the domain. In later sections we develop a criterion  
 12 based on the full chemical network for a reacting system in which some species may be immobilized either in the  
 13 interior of the domain or on the boundary, and we compare the criteria in Table 2 with the criterion that emerges  
 14 from our analysis.

15 Another aspect inadequately addressed in previously-cited work is the effect of compartment size on the magnitude  
 16 of the stochastic fluctuations, as measured by the CV of solutions. As suggested by the example of diffusion alone,  
 17 the noise can be expected to grow as the compartment size decreases, and this raises the question of how to choose  $h$   
 18 so as to ensure that the discretized system is accurate, and yet minimizes some measure of the noise. In the following  
 19 section we analyze linear reaction networks to address this issue, and show that a certain scaling of the coefficient  
 20 of variation stabilizes as  $h$  decreases, which leads to an optimal choice of  $h$  (in the sense that it is taken as large as  
 21 possible). In Section 3 we develop a general criterion for the choice of compartment size that applies to all orders of  
 22 reactions and both diffusing and non-diffusing species.

## 23 2. Linear stochastic reaction-diffusion networks

### 24 2.1. Measures of the fluctuations in a compartmental system

25 We consider a first-order chemical reaction-diffusion network, and from previous work we know that the reactions  
 26 fall into one of four classes: production from a source, degradation, conversion, and catalytic production from a  
 27 source [17]. Production from a source is an input reaction of the form  $\emptyset \rightarrow \mathcal{M}_i$ ; degradation is a reaction  $\mathcal{M}_i \rightarrow \emptyset$ ;  
 28 conversion is a reaction  $\mathcal{M}_j \rightarrow \mathcal{M}_i$ ; and catalytic production from a source is a reaction  $\emptyset \xrightarrow{\mathcal{M}_j} \mathcal{M}_i$ . We assume that  
 29 degradation, conversion, and catalyzed production from a source occur at all spatial positions, but production from  
 30 a source may be spatially nonuniform. Suppose that the domain  $\Omega$  is a rectangular parallelepiped with dimensions  
 31  $L_x \times L_y \times L_z$  of volume  $\mathbf{V} = L_x L_y L_z$ . We divide  $\Omega$  into  $N_c = N_x N_y N_z$  compartments, each of which has volume  
 32  $\mathbf{V}/N_c$ . Define the index  $k = (k_1, k_2, k_3)$  to denote a compartment, where  $k_1, k_2, k_3 = 1, \dots, N_x, N_y, N_z$ , respectively,  
 33 and set  $\boldsymbol{\eta} = (k, i)$  where  $i = 1, \dots, s$  denotes the species. Let  $N_i^k(t) \equiv N^\boldsymbol{\eta}(t)$  be a random variable that represents the  
 34 number of molecules of species  $\mathcal{M}_i$  in the  $k^{th}$  compartment at time  $t$ , and let  $N^k$  denote the vector of  $N_i$ s. Several  
 35 different measures of the noise will be introduced later to understand the dependence of noise on the discretization,

1 all of which involve the means and variances of the components in the network, and thus we first analyze the evolution  
2 of these quantities.

3 The mean  $M(t)$  is an  $sN_c$ -dimensional vector of first moments with elements defined by

$$4 \quad [M(t)]_{\mathbf{i}(\boldsymbol{\eta})} = E[N^{\boldsymbol{\eta}}(t)], \quad (9)$$

5 where the index function  $\mathbf{i}$ , which is defined as  $\mathbf{i}(\boldsymbol{\eta}) = i + ((k_1 - 1)N_y N_z + (k_2 - 1)N_z + (k_3 - 1))s$ , labels the  
6 components. Define  $\boldsymbol{\zeta} = (q, j)$  as the index for the  $j^{\text{th}}$  species in the  $q^{\text{th}}$  compartment. Then the matrix of second  
7 moments  $V(t)$  is an  $sN_c \times sN_c$  matrix with elements

$$8 \quad [V(t)]_{\mathbf{i}(\boldsymbol{\eta}), \mathbf{i}(\boldsymbol{\zeta})} = E[N^{\boldsymbol{\eta}}(t)N^{\boldsymbol{\zeta}}(t)] - E[N^{\boldsymbol{\eta}}(t)]E[N^{\boldsymbol{\zeta}}(t)].$$

9 The covariance matrix,  $Cov(t)$ , is defined as

$$10 \quad [Cov(t)]_{\mathbf{i}(\boldsymbol{\eta}), \mathbf{i}(\boldsymbol{\zeta})} = E[N^{\boldsymbol{\eta}}(t)N^{\boldsymbol{\zeta}}(t)] - E[N^{\boldsymbol{\eta}}(t)]E[N^{\boldsymbol{\zeta}}(t)],$$

11 and therefore can be expressed in terms of  $M(t)$  and  $V(t)$  as

$$12 \quad Cov(t) = V(t) - M(t)M(t)^T + M_d(t),$$

13 where  $M_d(t) = \text{diag}[M(t)]$  is a diagonal matrix whose entries are those of  $M(t)$ .

14 As in [17], let  $\mathcal{K}$  be the  $s \times s$  reaction matrix for conversion or degradation of species within a compartment, let  $\mathcal{D}$  be  
15 the  $s \times s$  diagonal diffusion matrix, let  $K^{cat}$  be the  $s \times s$  matrix wherein the  $(i, j)$ th element is the catalytic production  
16 rate from a source of the  $i^{\text{th}}$  species catalyzed by the  $j^{\text{th}}$  species, and let  $K^S$  be the  $sN_c \times sN_c$  diagonal matrix with  
17 the  $\mathbf{i}(\boldsymbol{\eta})^{\text{th}}$  diagonal element representing an input of the  $i^{\text{th}}$  species from a source into the  $k^{\text{th}}$  compartment<sup>2</sup>. Further,  
18 let  $\Delta$  be the discrete Laplacian for the domain  $\Omega$  with Neumann boundary conditions;  $\Delta$  encodes the topology of the  
19 network, and more general topologies can be treated similarly [35]. Finally, we let  $\mathbf{u}$  and  $I_{N_c}$  be a vector of length  
20  $sN_c$ , all of whose entries are 1, and the unit matrix of dimension  $N_c \times N_c$ , respectively. We let  $\mathbf{k}^s$  be the vector whose  
21 components are the diagonal elements of  $K^S$ , and we define the rank one square matrix  $S = [\mathbf{k}^s | \mathbf{k}^s | \dots | \mathbf{k}^s]$  whose  
22 columns are  $\mathbf{k}^s$ .

23 These definitions lead to a reaction-diffusion matrix defined as  $\Omega \equiv \Delta \otimes \mathcal{D} + I_{N_c} \otimes \mathcal{K}$  [17], and we assume  
24 throughout that  $\Omega$  is semi-simple, that the kinetic system is stable, and that the spectrum  $\sigma(\Omega)$  lies in the closed  
25 left-half plane, denoted  $\overline{LHP}$ . A spectral representation of  $\Omega$  is given in Appendix B. We consider two cases defined  
26 by whether  $\Omega$  has a zero eigenvalue or not (multiple zero eigenvalues can be treated similarly). The presence or  
27 absence of a zero eigenvalue is determined solely by the kinetic matrix, and thus we exclude Turing-type instabilities  
28 that arise from the interaction of reaction and diffusion. If  $\mathcal{G}_c$  is strongly connected, as in the case treated here, the  
29 Laplacian has exactly one zero eigenvalue for Neumann boundary conditions, and  $\sigma(\Omega) \subset LHP$  if the system is either  
30 closed with degradation or open with production from a source and degradation. In the absence of inputs the mean  
31 and variance decay to zero as  $t \rightarrow \infty$ , which is of no interest, and in the presence of inputs they are determined by  
32 the inputs as  $t \rightarrow \infty$ . On the other hand, if there is exactly one zero eigenvalue of  $\Omega$  and no inputs, the steady-state  
33 probability distribution is determined by the eigenvector corresponding to the zero eigenvalue [17]. Other cases may  
34 arise, but are not treated here.

35 The first and second moments  $M(t)$  and  $V(t)$  are solutions of the ordinary differential equations

$$36 \quad \frac{dM(t)}{dt} = \Omega M(t) + \mathbf{k}^s \quad (10)$$

$$37 \quad \frac{dV(t)}{dt} = \Omega V(t) + [\Omega V(t)]^T + C(t) + C(t)^T$$

---

<sup>2</sup> This formulation can be generalized to allow catalyzed production from a source to be space-dependent as well, but we do not include this here.

1 where  $C(t) = W(t) + \mathbf{k}^s M(t)^T$  [17]. Here  $W(t)$  is a block-diagonal matrix with elements defined as

$$2 \quad [W(t)]_{i(\boldsymbol{\eta}),i(\boldsymbol{\zeta})} = \begin{cases} K_{ij}^{cat}[M(t)]_{i(\boldsymbol{\zeta})} & \text{if } k = q \\ 0 & \text{otherwise.} \end{cases}$$

3 Define indices  $l, m, \text{ and } n$  which run over compartments, let  $\mu, \chi, \text{ and } \gamma$  be indices that label species, and define  
4 the indices  $\mathbf{l} = (l, \mu)$ ,  $\mathbf{m} = (m, \chi)$ , and  $\mathbf{n} = (n, \gamma)$ . The eigenvalues of  $\Omega = \Delta \otimes \mathcal{D} + I_{N_c} \otimes \mathcal{K}$  can be computed  
5 once those of  $\Delta$ , denoted by  $\alpha_l$ , are known [35]. They are eigenvalues of the matrix pencil  $\mathcal{K} + \alpha_l \mathcal{D}$  and thus are the  
6 solutions of

$$7 \quad |\mathcal{K} + \alpha_l \mathcal{D} - \lambda_1 I_s| = 0,$$

8 The corresponding eigenvectors of  $\Omega$  are the tensor product of eigenvectors of  $\Delta$  and those of  $\mathcal{K} + \alpha_l \mathcal{D}$ . Let  $\phi_l$  and  
9  $\phi_l^*$  be the eigenvector and adjoint eigenvector of  $\Delta$  for the eigenvalue  $\alpha_l$ , and let  $\varphi_1$  and  $\varphi_1^*$  be the corresponding  
10 eigenvector and adjoint eigenvector of  $\mathcal{K} + \alpha_l \mathcal{D}$  for  $\lambda_1$ . Then  $\phi_l \otimes \varphi_1$  is the eigenvector of  $\Omega$  corresponding to the  
11 eigenvalue  $\lambda_1$ , and the projection  $P_1$  associated with  $\lambda_1$  is defined as

$$12 \quad P_1 = (\phi_l * \overline{\phi_l^*}) \otimes (\varphi_1 * \overline{\varphi_1^*}),$$

13 where  $\overline{\phi_l^*}$  and  $\overline{\varphi_1^*}$  are complex conjugates of  $\phi_l^*$  and  $\varphi_1^*$ , respectively. Here  $*$  is the dyad product defined operationally  
14 as  $(u * v)w = \langle v, w \rangle u$ . Given these, we can compute the solution for the first two moments in terms of  $\lambda_1$  and  $P_1$  in  
15 the semisimple case.

Later we focus primarily on the steady-state level of fluctuations, and therefore we define

$$M_\infty \equiv \lim_{t \rightarrow \infty} M(t) \quad \text{and} \quad V_\infty \equiv \lim_{t \rightarrow \infty} V(t).$$

16 First, consider the case in which  $\sigma(\Omega) \subset LHP$ ; then one finds that

$$17 \quad M_\infty = - \sum_{\mathbf{l}} \frac{P_{\mathbf{l}}}{\lambda_{\mathbf{l}}} \mathbf{k}^s, \tag{11}$$

$$18 \quad V_\infty = \sum_{\mathbf{m}} \sum_{\mathbf{l}} \left[ \frac{1}{\lambda_{\mathbf{l}}(\lambda_{\mathbf{l}} + \lambda_{\mathbf{m}})} P_{\mathbf{m}} K^S (P_{\mathbf{l}} S)^T + \frac{1}{\lambda_{\mathbf{m}}(\lambda_{\mathbf{l}} + \lambda_{\mathbf{m}})} P_{\mathbf{m}} S K^S P_{\mathbf{l}}^T \right.$$

$$19 \quad \left. + \sum_{\mathbf{n}} \frac{P_{\mathbf{m}} \{ \text{diag}[P_{\mathbf{n}} S] (K^{cat})^T + K^{cat} \text{diag}[P_{\mathbf{n}} S] \} P_{\mathbf{l}}^T}{\lambda_{\mathbf{n}}(\lambda_{\mathbf{l}} + \lambda_{\mathbf{m}})} \right],$$

20 where the sums range over the eigenvalues of  $\mathcal{K} + \alpha_l \mathcal{D}$ .

21 To assess the fluctuations in the network we can use one of several forms of a coefficient of variation, defined  
22 as the standard deviation divided by the mean. The first measure results from defining the noise component- and  
23 compartment-wise, which may be appropriate when assessing the effect of fluctuations in a morphogen used to define  
24 the boundary between two tissue types. In this case one computes the standard deviation of the number of molecules  
25 for the  $i^{th}$  species in the  $k^{th}$  compartment divided by its mean, *viz.*,

$$26 \quad (CV(t))^\boldsymbol{\eta} \equiv \frac{\sqrt{E[N^\boldsymbol{\eta}(t)^2] - E[N^\boldsymbol{\eta}(t)]^2}}{E[N^\boldsymbol{\eta}(t)]} \tag{12}$$

27 where  $\boldsymbol{\eta} = (k, i)$ . This reflects the fluctuations of each species in each compartment, and thereby leads to a total of  
28  $sN_c$  measures, but for our purpose a single global measure that averages over all species and compartments is more  
29 appropriate. There are many ways to do this – one could for instance use the average of the component measures in  
30 (12) or the maximum of these. However, we use a measure based on a normalized covariance matrix [47], which is  
31 defined as

$$32 \quad [\Sigma_0(t)]_{i(\boldsymbol{\eta}),i(\boldsymbol{\zeta})} = \frac{E[N^\boldsymbol{\eta}(t)N^\boldsymbol{\zeta}(t)] - E[N^\boldsymbol{\eta}(t)]E[N^\boldsymbol{\zeta}(t)]}{E[N^\boldsymbol{\eta}(t)]E[N^\boldsymbol{\zeta}(t)]}$$

$$\equiv (M_d(t)^{-1} Cov(t) M_d(t)^{-1})_{i(\boldsymbol{\eta}), i(\boldsymbol{\zeta})}.$$

We then define a generalized coefficient of variation as the square root of the maximum eigenvalue of  $\Sigma_0(t)$ , and denote it

$$CV^*(t) \equiv \sqrt{\lambda_{max}(\Sigma_0(t))}. \quad (13)$$

Since the covariance matrix is symmetric and at least positive semi-definite, this is well-defined.

We define other steady-state variables as

$$Cov_\infty \equiv \lim_{t \rightarrow \infty} Cov(t) \quad M_{d,\infty} \equiv \lim_{t \rightarrow \infty} M_d(t) \quad (CV_\infty)^\eta \equiv \lim_{t \rightarrow \infty} (CV(t))^\eta \quad \Sigma_{0,\infty} \equiv \lim_{t \rightarrow \infty} \Sigma_0(t)$$

and then the steady-state  $CV^*$  is given by

$$CV^* = \sqrt{\lambda_{max}(\Sigma_{0,\infty})} = \sqrt{\lambda_{max}(M_{d,\infty}^{-1}(V_\infty - M_\infty M_\infty^T + M_{d,\infty})M_{d,\infty}^{-1})}.$$

In the following proposition, we express  $CV^*$  in terms of  $M_\infty$  alone when the eigenvalues of  $\Omega$  are in the *LHP* and there are no catalyzed inputs to the system. When there is exactly one zero eigenvalue of  $\Omega$  and no inputs, we express  $CV^*$  in terms of  $M_{d,\infty}$ ,  $M_d(0)$ , and  $P_s$ , where  $P_s$  is the projection corresponding to the zero eigenvalue.

**Proposition 1.** *Suppose that the eigenvalues of  $\Omega$  are in the LHP; then*

1. *If there are no non-catalyzed inputs ( $K^S = 0$ ), then  $M_{d,\infty} = Cov_\infty = 0$*

2. *If  $K^S \neq 0$  then*

$$Cov_\infty = M_{d,\infty} + \sum_{\mathbf{l}, \mathbf{m}, \mathbf{n}} \frac{P_{\mathbf{m}} \left\{ \text{diag}[P_{\mathbf{n}} S] (K^{\text{cat}})^T + K^{\text{cat}} \text{diag}[P_{\mathbf{n}} S] \right\} P_{\mathbf{l}}^T}{\lambda_{\mathbf{n}} (\lambda_{\mathbf{l}} + \lambda_{\mathbf{m}})}.$$

and

$$\Sigma_{0,\infty} = M_{d,\infty}^{-1} + M_{d,\infty}^{-1} \left\{ \sum_{\mathbf{l}, \mathbf{m}, \mathbf{n}} \frac{P_{\mathbf{m}} \left\{ \text{diag}[P_{\mathbf{n}} S] (K^{\text{cat}})^T + K^{\text{cat}} \text{diag}[P_{\mathbf{n}} S] \right\} P_{\mathbf{l}}^T}{\lambda_{\mathbf{n}} (\lambda_{\mathbf{l}} + \lambda_{\mathbf{m}})} \right\} M_{d,\infty}^{-1}$$

If  $K^{\text{cat}} = 0$ , the second term vanishes and

$$CV^* = \sqrt{\frac{1}{\min_{\boldsymbol{\eta}} [M_\infty]_{i(\boldsymbol{\eta})}}}. \quad (14)$$

3. *Suppose that  $\sigma(\Omega) \subset \overline{\text{LHP}}$  and there is exactly one zero eigenvalue of  $\Omega$  and no inputs; then*

$$Cov_\infty = M_{d,\infty} - P_s M_d(0) P_s^T$$

where  $P_s$  is the projection corresponding to the zero eigenvalue and

$$\Sigma_{0,\infty} = M_{d,\infty}^{-1} - M_{d,\infty}^{-1} P_s M_d(0) P_s^T M_{d,\infty}^{-1}. \quad (15)$$

*Proof.* The proof is given in Appendix C.  $\square$

It is clear from the foregoing that  $CV^*$  is an increasing function of the number of compartments in the system, since the mean number of molecules of each species in each compartment decreases. However, as the following example illustrates, and as will be proven later, a scaled version of  $CV^*$  stabilizes as the compartment number increases. Define  $V_c = \mathbf{V}/N_c$  and the scaled variables

$$\overline{M}_\infty \equiv \frac{M_\infty}{V_c}, \quad \overline{CV}^* \equiv \sqrt{V_c} CV^*, \quad (\overline{CV}_\infty)^\eta \equiv \sqrt{V_c} (CV_\infty)^\eta, \quad (16)$$

where  $\boldsymbol{\eta} = (k, i)$  and  $(X)^{(k,i)}$  represents the component of  $X$  corresponding to the  $i^{\text{th}}$  species in the  $k^{\text{th}}$  compartment. Eq. (14) shows that in the absence of catalyzed inputs, the least abundant species, evaluated over all compartments,

1 determines  $CV^*$ , and one sees that convergence of  $\overline{M}_\infty$  implies convergence of  $\overline{CV}^*$  as  $N_c \rightarrow \infty$ . Convergence of the  
2 former as  $N_c \rightarrow \infty$  is shown in Theorem 1 below.

3 Denote by  $\mathcal{K}'$  the reaction matrix for conversion or degradation of species in a compartment for systems with  
4 non-catalytic production of diffusing species, and rearrange the species order so that  $\mathcal{K}'$  can be partitioned into block  
5 matrices as

$$6 \quad \mathcal{K}' = \begin{bmatrix} \mathcal{R} & \mathcal{S} \\ \mathcal{T} & \mathcal{W} \end{bmatrix}.$$

7 Here  $\mathcal{R}$  ( $\mathcal{W}$ ) is the reaction matrix for conversion between diffusing (non-diffusing) species or degradation of diffusing  
8 (non-diffusing) species, and  $\mathcal{S}$  ( $\mathcal{T}$ ) is the reaction matrix for conversion of non-diffusing (diffusing) species to diffusing  
9 (non-diffusing) species. We denote by  $X(t)$  and  $Y(t)$  the mean vectors for diffusing and non-diffusing species numbers,  
10 with each element defined as in (9), and write the governing equations for these means as follows:

$$11 \quad \begin{aligned} \frac{dX(t)}{dt} &= (\Delta \otimes \mathcal{D} + I_{N_c} \otimes \mathcal{R}) X(t) + (I_{N_c} \otimes \mathcal{S}) Y(t) + \mathbf{k}^s, \\ \frac{dY(t)}{dt} &= (I_{N_c} \otimes \mathcal{T}) X(t) + (I_{N_c} \otimes \mathcal{W}) Y(t). \end{aligned} \quad (17)$$

12 Let  $X_\infty$  and  $Y_\infty$  be the steady-state solutions of (17). Assuming that  $\sigma(\mathcal{W}) \subset LHP$ , the steady-state mean vector  
13 for non-diffusing species can be expressed in terms of  $X_\infty$ , where  $X_\infty$  is the solution of

$$14 \quad (\Delta \otimes \mathcal{D} + I_{N_c} \otimes \mathcal{K}) X_\infty + \mathbf{k}^s = 0, \quad (18)$$

15 where  $\mathcal{K} \equiv \mathcal{R} - \mathcal{S}\mathcal{W}^{-1}\mathcal{T}$ . To show convergence of  $X_\infty/(N_A V_c)$  to the solution of the continuum deterministic  
16 reaction-diffusion system, we consider a 1-dimensional domain  $[0, L_x]$  for simplicity, and we let  $\tilde{\alpha}_l$  be the solution of  
17 the following scalar problem:

$$18 \quad \begin{aligned} \frac{d^2 \tilde{q}(\mathbf{x})}{d\mathbf{x}^2} &= \tilde{\alpha} \tilde{q}(\mathbf{x}), & \mathbf{x} \in [0, L_x], \\ \tilde{q}'(\mathbf{x}) &= 0, & \mathbf{x} = 0, L_x. \end{aligned} \quad (19)$$

19 **Theorem 1.** *Let  $\tilde{\mathcal{D}}$  be the diffusion matrix for the continuum problem, and suppose that*

$$20 \quad \sigma(\Omega), \sigma(\mathcal{W}), \sigma(\mathcal{K} + \alpha_l \mathcal{D}), \sigma(\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}) \subset LHP,$$

21 *and that  $\mathcal{K} + \alpha_l \mathcal{D}$  and  $\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}$  are semi-simple. Assume that  $K^{cat} = 0$  and assume that only diffusing species are  
22 produced from a source and production occurs only in the left-most compartment. Then,*

- 23 •  $[\overline{M}_\infty]_{\mathbf{i}(\eta)}$  converges to the steady-state solution of the corresponding continuum deterministic reaction-diffusion  
24 system as  $N_c \rightarrow \infty$ ,
- 25 •  $\overline{CV}^*$  converges to the limit of  $\sqrt{\frac{1}{\min_\eta [\overline{M}_\infty]_{\mathbf{i}(\eta)}}}$  as  $N_c \rightarrow \infty$ .

26 *Proof.* The proof is given in Appendix D.  $\square$

27 For systems with no inputs, it follows from (15) that if we show that each element of  $V_c \left( M_{d,\infty}^{-1} P_s M_d(0) P_s^T M_{d,\infty}^{-1} \right)$   
28 converges to zero as  $N_c \rightarrow \infty$ , convergence of  $\overline{M}_\infty$  will imply convergence of  $\overline{CV}^*$  as  $N_c \rightarrow \infty$ , as stated in the  
29 following theorem, again under the assumption of a one-dimensional domain.

30 **Theorem 2.** *Suppose that there are no inputs, that  $\sigma(\Omega) \subset \overline{LHP}$ , and there is exactly one zero eigenvalue of  $\Omega$ .  
31 Then,*

- 32 •  $[\overline{M}_\infty]_{\mathbf{i}(\eta)}$  converges to the steady-state solution of the corresponding continuum deterministic reaction-diffusion  
33 system as  $N_c \rightarrow \infty$ ,
- 34 •  $\overline{CV}^*$  converges to the limit of  $\sqrt{\frac{1}{\min_\eta [\overline{M}_\infty]_{\mathbf{i}(\eta)}}}$  as  $N_c \rightarrow \infty$ .



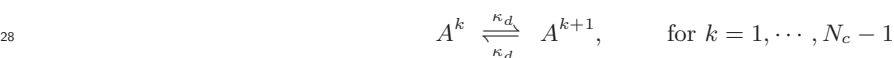
1 *Proof.* The proof is given in Appendix E.  $\square$

2 In essence these results show that the concentration-based CV for the discretized problem stabilizes.

### 3 2.2. Example

4 The example in this section is motivated by the classical French flag paradigm of pattern formation in a one-  
5 dimensional system discussed earlier. In that context, a morphogen is produced at one end of a 1D domain, diffuses  
6 into the domain, and binds to receptors and is perhaps degraded by a first-order process [51]. The problem is to  
7 reliably partition the domain into 3 equal-size sub-domains (corresponding to blue, white and red, beginning at  
8 the staff) by setting thresholds that determine the extent of each domain. A system that exemplifies this arises in  
9 anterior-posterior pattern formation in the fruit fly *Drosophila melanogaster*. The *Drosophila* embryo is approximately  
10 ellipsoidal, and is surrounded by a thin fluid layer, called the perivitelline (PV) space, bounded by the outer membrane.  
11 The coordinate frame of the embryo is first established by gradients of inherited maternal factors in the anterior-  
12 posterior (AP) direction and by gradients of factors in the PV space in the dorsal-ventral direction. The first level of AP  
13 patterning is mediated by the morphogen Bicoid, which is a transcription factor that is transcribed from maternally-  
14 inherited mRNA localized at the anterior end of the embryo. Because production is localized at the anterior end, the  
15 concentration of the Bicoid protein forms a monotone distribution with the high point at the anterior end.

16 We consider a rectangular solid continuum whose major axis lies along the x axis, and we let  $L_x = 275 \mu m$ ,  
17  $L_y = 5 \mu m$ , and  $L_z = 0.5 \mu m$  represent the lengths of each side of the system, motivated by a slice of the PV space  
18 in *Drosophila* [48]. Since  $L_x \gg L_y, L_z$ , we set  $N_y = N_z = 1$ , and determine the appropriate discretization defined  
19 by  $N_x = N_c$  in the x-direction. We suppose that there is one diffusing species,  $A$ , and two non-diffusing species,  $B$   
20 and  $C$ , which represent ligand, receptor with ligand bound, and downstream signal, respectively. Assuming that the  
21 number of receptors is large, as it is in many biological systems, receptor-ligand binding is described as a first order  
22 reaction. The complete set of reactions is as follows.



29 Here superscript  $k$  denotes the species in the  $k^{th}$  compartment.

30 All reactions are either production from a source, degradation, or conversion, in the terminology of Gadgil *et al.*  
31 [17]; there is no catalytic production from a source. The reaction (20) simply describes a linearized ligand-binding to  
32 receptors. The reaction (21) describes production of species  $A$  from a source located at the leftmost compartment,  
33 and thus  $A$  has an input only to the first compartment. The coefficients  $\kappa_m$ , for  $m = 5, 6, 7, 8, 9$ , are stochastic  
34 reaction rate constants and  $\kappa_d$  is the diffusion rate constant for species  $A$ . These are derived from the corresponding  
35 deterministic rate constants estimated in [49], and both the deterministic and the stochastic parameters are given in  
36 Table 3.

Table 3: Deterministic and stochastic parameters in Example 2.2

Deterministic		Stochastic	
$\kappa_5$	$1 nM^{-1} min^{-1}$	$\kappa_5$	$k_5 / (\mathcal{N}_A V_c) \approx 2.42 \times 10^{-3} N_c min^{-1}$
$\kappa_{-5}$	$2 min^{-1}$	$\kappa_{-5}$	$2 min^{-1}$
$\kappa_6$	$1 min^{-1}$	$\kappa_6$	$1 min^{-1}$
$\kappa_7$	$0.03 min^{-1}$	$\kappa_7$	$0.03 min^{-1}$
$\kappa_8$	$250 nM \mu m min^{-1}$	$\kappa_8$	$k_8 (\mathcal{N}_A \mathbf{V}) / L_x \approx 376.38 min^{-1}$
$\kappa_9$	$0.03 min^{-1}$	$\kappa_9$	$0.03 min^{-1}$

$c_R$	$320 \text{ nM}$	$R$	$c_R(\mathcal{N}_A V_c) \approx 132484/N_c$
$D$	$4380 \mu\text{m}^2 \text{ min}^{-1}$	$\kappa_d$	$D/(L_x/N_c)^2 \approx 5.79 \times 10^{-2} N_c^2 \text{ min}^{-1}$
$L_x$	$275 \mu\text{m}$		
$L_y$	$5 \mu\text{m}$		
$L_z$	$0.5 \mu\text{m}$		
$\mathbf{V}$	$687.5 \mu\text{m}^3$		

<sup>2</sup> In the kinetic scheme  $\kappa_5$  is a stochastic rate constant for ligand-receptor binding, and thus depends on the volume,  
<sup>1</sup>  
<sup>3</sup> but since we assume that the receptor density is large compared to the signal, the product of the binding constant and  
<sup>4</sup> the receptor density is a pseudo-first-order rate constant, and hence independent of the volume. All other reactions  
<sup>5</sup> except for production of species  $A$  are first order, and the deterministic and stochastic rate constants are independent  
<sup>6</sup> of the volume and hence  $N_c$ . To standardize the input as we change  $N_c$ , we must hold the total flux of  $A$  constant.  
<sup>7</sup> An estimate of an input on a volumetric basis given in [49] is converted to a flux per unit area by multiplying by the  
<sup>8</sup> length of the longest edge,  $L_x = 275 \mu\text{m}$ ; this yields  $k_8$ . The corresponding stochastic rate,  $\kappa_8$ , is given as follows:

$$\kappa_8 = \frac{k_8(\mathcal{N}_A \mathbf{V})}{L_x}.$$

<sup>10</sup> Therefore,  $\kappa_8$  does not depend on  $N_c$ . The stochastic diffusion rate constant  $\kappa_d$  is computed by dividing the continuum  
<sup>11</sup> deterministic diffusion rate constant by  $(L_x/N_c)^2$ , via discretization of the Laplacian, and thus scales as  $N_c^2$ .

<sup>12</sup> Let  $X(t)$  be the first moment vector for the diffusing species  $A$  and  $Y(t)$  be the first moment vector for the non-  
<sup>13</sup> diffusing species  $B$  and  $C$ .  $[X(t)]_{i(k,1)}$  denotes the mean number of molecules of species  $A$  in the  $k^{\text{th}}$  compartment,  
<sup>14</sup> and  $[Y(t)]_{i(q,1)}$  and  $[Y(t)]_{i(q,2)}$  represent the mean numbers of species  $B$  and  $C$  in the  $q^{\text{th}}$  compartment, respectively.  
<sup>15</sup> The evolution of the first moments is governed by

$$\frac{dX(t)}{dt} = (\Delta \otimes \mathcal{D} + I_{N_c} \otimes \mathcal{R}) X(t) + (I_{N_c} \otimes \mathcal{S}) Y(t) + \mathbf{k}^s \quad (22)$$

$$\frac{dY(t)}{dt} = (I_{N_c} \otimes \mathcal{T}) X(t) + (I_{N_c} \otimes \mathcal{W}) Y(t). \quad (23)$$

<sup>18</sup> The diffusion matrix, reaction matrices, and a matrix for production rate from a source are given as

$$\mathcal{D} = \kappa_d, \quad \mathcal{R} = -\kappa_5 R - \kappa_6, \quad \mathcal{S} = \begin{bmatrix} \kappa_{-5} & 0 \end{bmatrix}, \quad \mathcal{T} = \begin{bmatrix} \kappa_5 R \\ 0 \end{bmatrix},$$

$$\mathcal{W} = \begin{bmatrix} -\kappa_{-5} - \kappa_7 & 0 \\ \kappa_7 & -\kappa_9 \end{bmatrix}, \quad \mathbf{k}^s = [\kappa_8, 0, \dots, 0]^T.$$

<sup>21</sup> In this example, the eigenvalues of all principal submatrices of the reaction matrices  $\mathcal{R}$  and  $\mathcal{W}$  are in the *LHP*.

<sup>22</sup> Let  $X_\infty$  be the steady-state first moment vector for the number of molecules of species  $A$  and let  $Y_\infty$  be the  
<sup>23</sup> steady-state first moment vector for the numbers of molecules of species  $B$  and  $C$ . Using the fact that species  $B$  and  
<sup>24</sup>  $C$  do not diffuse, we compute  $Y_\infty$  in terms of  $X_\infty$ .

$$Y_\infty = - (I_{N_c} \otimes (\mathcal{W}^{-1} \mathcal{T})) X_\infty.$$

<sup>26</sup> Then we find that

$$[Y_\infty]_{i(k,1)} = \frac{\kappa_5 R}{\kappa_{-5} + \kappa_7} [X_\infty]_{i(k,1)} \quad (24)$$

$$[Y_\infty]_{i(k,2)} = \frac{\kappa_5 \kappa_7 R}{\kappa_9 (\kappa_{-5} + \kappa_7)} [X_\infty]_{i(k,1)} \quad (25)$$

<sup>29</sup> Converting the steady-state first moment of  $B$  into that of  $A$ , the effective reaction rate of species  $A$  at steady-state  
<sup>30</sup> becomes

$$\mathcal{K} \equiv \mathcal{R} - \mathcal{S} \mathcal{W}^{-1} \mathcal{T}$$

$$= -\frac{\kappa_5 \kappa_7 R}{\kappa_{-5} + \kappa_7} - \kappa_6.$$

Note that  $\mathcal{K}$  is a scalar since there is exactly one diffusing species,  $A$ . Then,  $X_\infty$  satisfies

$$(\Delta \otimes \mathcal{D} + I_{N_c} \otimes \mathcal{K}) X_\infty + \mathbf{k}^s = 0. \quad (26)$$

Define  $\overline{X}_\infty \equiv X_\infty / V_c$ . To calculate the steady-state first moment for species  $A$ , we first compute the eigenvalues and corresponding projections of the reaction-diffusion matrix  $\Omega = \Delta \otimes \mathcal{D} + I_{N_c} \otimes \mathcal{K}$ . As before, let  $\alpha_l$  denote an eigenvalue of  $\Delta$ , let  $\lambda_l$  be an eigenvalue of  $\Omega$ , and let  $P_l$  be the corresponding projection of  $\Omega$ . Then according to (11), the scaled steady-state first moment of  $A$  is

$$\overline{X}_\infty = -\frac{1}{V_c} \sum_l \frac{P_l}{\lambda_l} \mathbf{k}^s. \quad (27)$$

In Figure 3 we show the relation between  $\alpha_l$  and  $\lambda_l$  for  $N_c = 10, 20, 30, 40$ . The range of  $\alpha_l$  is fixed at  $[-4, 0]$  by the structure of the network and the boundary conditions, but the range of  $\lambda_l$  increases as  $N_c$  is increased because higher spatial frequencies are captured with increasing  $N_c$ . Since we impose Neumann data on the boundary in the discrete problem, the smallest eigenvalue in magnitude is  $\lambda_{N_c,1} \approx -5.73$  corresponding to  $\alpha_{N_c} = 0$ . The dominant term in (27) corresponds to  $\lambda_{N_c,1}$ , and is independent of  $N_c$ . Using (24) and (25), we compute  $\max_k (\overline{CV}_\infty)^{k,i}$  for  $i = A, B, C$

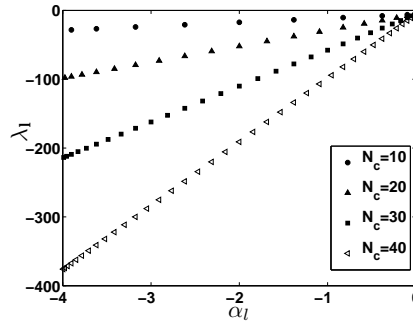


Fig. 3. Eigenvalues of  $\Delta$  and  $\Omega$  when  $N_c = 10, 20, 30, 40$

in terms of  $\min_k [\overline{X}_\infty]_{i(k,1)}$ .

$$\max_k (\overline{CV}_\infty)^{k,A} = \sqrt{\frac{1}{\min_k [\overline{X}_\infty]_{i(k,1)}}} \quad (28)$$

$$\max_k (\overline{CV}_\infty)^{k,B} = \sqrt{\frac{\kappa_{-5} + \kappa_7}{\kappa_5 R}} \sqrt{\frac{1}{\min_k [\overline{X}_\infty]_{i(k,1)}}} \quad (29)$$

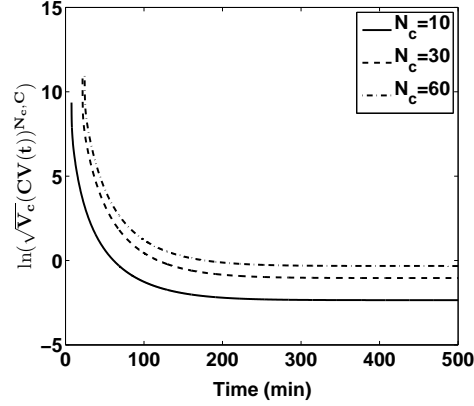
$$\max_k (\overline{CV}_\infty)^{k,C} = \sqrt{\frac{(\kappa_{-5} + \kappa_7) \kappa_9}{\kappa_5 \kappa_7 R}} \sqrt{\frac{1}{\min_k [\overline{X}_\infty]_{i(k,1)}}} \quad (30)$$

Also, we have

$$\overline{CV}^* = \sqrt{\frac{1}{\min \left( \min_k [\overline{X}_\infty]_{i(k,1)}, \frac{\kappa_5 R}{\kappa_{-5} + \kappa_7} \min_k [\overline{X}_\infty]_{i(k,1)}, \frac{\kappa_5 \kappa_7 R}{\kappa_9 (\kappa_{-5} + \kappa_7)} \min_k [\overline{X}_\infty]_{i(k,1)} \right)}}.$$

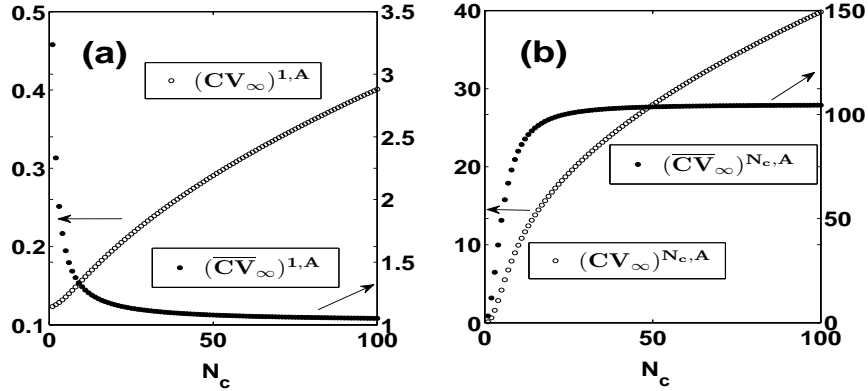
In Appendix D we prove that  $\overline{X}_\infty / \mathcal{N}_A$  converges to the concentration satisfying the corresponding continuum model in which the source is located at  $x = 0$ . Due to the source location at  $x = 0$ , the steady-state concentration is monotone decreasing in  $x$ . One can show that the same is true in the discrete problem, and the proof is left to the reader.

1 Since we later only use the steady-state measure for the noise to determine the number of compartments, one has  
 2 to check that the steady-state is reached rapidly. In Figure 4 we show the transient evolution of  $\sqrt{V_c}(CV(t))^{N_c, C}$  in  
 3 the last compartment for  $N_c = 10, 30, 60$ . In this example species  $C$  serves as the downstream signal, and as we see in  
 4 Figure 4, the noise in the downstream signal stabilizes very quickly, which indicates that a compartment size based  
 on the steady-state noise level is appropriate here. Of course it may not always be.



**Fig. 4.** The evolution of  $\ln(\sqrt{V_c}(CV(t))^{N_c, C})$  in time for  $N_c = 10, 30, 60$ , when production is restricted to the first compartment and initial values are zero.

5  
 6 In Figures 5 and 6, we display the steady state noise for the kinetic scheme in Example 2.2 for the unscaled and  
 7 scaled CVs as a function of  $N_c$ . In Figure 5, we show  $(CV_\infty)^{k, A}$  for  $k = 1, N_c$  as we vary  $N_c$ . The values of  $(CV_\infty)^{k, B}$   
 8 and  $(CV_\infty)^{k, C}$  are simply scaled versions of  $(CV_\infty)^{k, A}$  and are not shown.  $[X_\infty]_{i(k, 1)}$  has its maximum and minimum  
 9 values in the first and the last compartments, respectively, by virtue of the monotonicity of the profile. Therefore, using  
 10 Proposition 1,  $(CV_\infty)^{k, i}$  has its maximum and minimum values in the last and the first compartments, respectively.  
 It is apparent in the figure that as  $N_c$  increases,  $(CV_\infty)^{k, i}$  gets larger due to the smaller number of molecules in each



**Fig. 5.** The CV of species A in the 1-dimensional model.

11  
 12 compartment, but  $(\overline{CV}_\infty)^{k, i}$  stabilizes due to convergence of  $\overline{X}_\infty$  to the corresponding concentration multiplied by  
 13  $\mathcal{N}_A$  as  $N_c \rightarrow \infty$ . This indicates that the contribution of diffusion to this scaled measure is dominated by the kinetic  
 14 contribution for large enough  $N_c$ . This explanation could be made rigorous by analyzing the asymptotic behavior of  
 15 the covariance as  $h \rightarrow 0$ , but we will not pursue this here.

1 In Figure 6 we illustrate the effect of the source location on the global measures  $CV^*$  and  $\overline{CV}^*$ . In (a) the source  
 2 for species  $A$  is in the first compartment as previously, while in (b) the input is divided amongst all compartments,  
 3 scaled so that the total input is fixed and independent of  $N_c$ . In (a)  $\overline{CV}^*$  stabilizes at about  $N_c = 30$ , while in (b)  
 4  $\overline{CV}^*$  is constant in  $N_c$ , whereas  $CV^*$  is monotone increasing in  $N_c$  in both cases. The stabilization of  $\overline{CV}^*$  with  $N_c$   
 5 suggests that the minimum number of compartments needed for an accurate representation of the noise is defined  
 6 by the smallest  $N_c$  at which  $\overline{CV}^*$  reaches a chosen percentage of the asymptotic value. As we show in the following  
 section, this agrees remarkably well with a criterion based on convergence of solutions to a uniform state. Although it

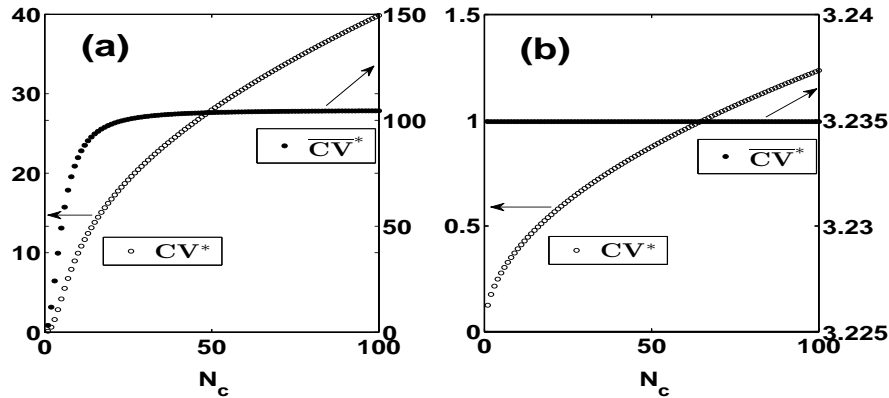


Fig. 6. A comparison of the noise for (a) spatially-nonuniform and (b) spatially-uniform inputs.

7 is not shown in (a), an increase of the diffusion constant decreases the value of  $N_c$  at which  $\overline{CV}^*$  stabilizes, as is to be  
 8 expected. The results in (b) suggest that distributing the input is an effective way of reducing the noise, but of course  
 9 this cannot be done if  $A$  is a morphogen that is used to determine cell types in a developing tissue. A comparison of  
 10  $(\overline{CV}_\infty)^{N_c, A}$  in Figure 5(b) and  $\overline{CV}^*$  in Figure 6(a) shows that these two values are equal, since  $[X_\infty]_{i(N_c, 1)}$  is the  
 11 smallest among all first moments of species  $A$  in all compartments, which is what is used for computing  $\overline{CV}^*$ .

12 In Section 1.4 we discussed different criteria to determine the compartment size, and here we apply Bernstein's  
 13 criterion to the foregoing example. Denote  $A_{tot}$ ,  $B_{tot}$ , and  $C_{tot}$  as the total numbers of molecules of species  $A$ ,  $B$ ,  
 14 and  $C$  at steady state, respectively. To compute the reaction timescale we have to compute the propensity of each  
 15 reaction, for which we use the steady-state first moments for species  $A$ ,  $B$ , and  $C$ , and this yields  $A_{tot} = 65.69$  and  
 16  $B_{tot} = C_{tot} = 10356$ . Since the propensity of conversion from  $A$  to  $B$  is the largest, we use this to compute the  
 17 timescale of the process,  $\tau_c$ . Thus

$$18 \quad \tau_c \approx \frac{1}{320 \text{ min}^{-1} \times \frac{A_{tot}}{N_c}}$$

$$19 \quad = \frac{1}{76.44 \times h_x} \text{ min.}$$

20 The diffusion timescale computed using the diffusion coefficient of species  $A$  is

$$21 \quad \frac{h_x^2}{2D} = \frac{h_x \mu\text{m}^2}{2 \times 4380 \mu\text{m}^2 \text{ min}^{-1}} = \frac{h_x^2}{8760} \text{ min.}$$

22 If we demand that the diffusion timescale be much smaller than the reaction timescale, we obtain

$$23 \quad h_x \ll \left( \frac{8760}{76.44} \right)^{1/3} \mu\text{m}$$

$$24 \quad \approx 4.86 \mu\text{m}.$$

25 In Section 3 we derive an upper bound for the compartment size for general systems, and as we show in Section  
 26 4, this leads to an estimate  $h_x < 8.25 \mu\text{m}$  (which corresponds to  $N_c > 33$ ). This result confirms the fact that it  
 27

1 is not necessary to discretize in the  $y$  or  $z$ -direction, since those dimensions are less than the maximum allowable  
 2 compartment size.

### 3 3. An upper bound for the compartment size

4 As was discussed earlier, current criteria for choosing an appropriate cell or compartment size for discretizing a  
 5 reaction-diffusion system are based on first estimating the smallest reaction time-scale and then choosing a compart-  
 6 ment size to ensure that the diffusion time scale is less than this. Since this is based on the objective of ensuring that  
 7 all molecules within a cell are accessible to each other, an alternate approach is to ask what conditions guarantee that  
 8 the solution of the coupled reaction-diffusion system converges to spatially-uniform solutions, whether stationary or  
 9 time-dependent. This obviously requires that diffusion dominates reaction in an appropriate sense, and that sense  
 10 was first developed in [31,3]. There it was shown that every species approaches a spatially-uniform solution at an  
 11 exponential rate in time when the diffusion constant for each species is sufficiently large relative to a measure of the  
 12 sensitivity of the kinetic network. More precisely, if  $c$  is the vector of species concentrations and  $\bar{c}$  is the spatial average  
 13 of  $c$ , then under the assumption that all species diffuse, the authors prove that  $\|c(\mathbf{x}, t) - \bar{c}(t)\|_{L_2} \rightarrow 0$  exponentially in  
 14  $t$  in any bounded domain  $\Omega$  in 1-, 2-, or 3-dimensional space if  $|\alpha_1 \delta| > \hat{r}$ . Here  $\delta$  is the smallest diffusion coefficient,  
 15  $\alpha_1$  is the largest non-zero eigenvalue of the Laplacian with homogeneous Neumann conditions on  $\partial\Omega$ , and  $\hat{r}$  is the  
 16 maximum Euclidean norm, taken over an appropriate set, of the Jacobian of the reaction terms. That result was gen-  
 17 eralized in [10]. Here we extend this result to allow non-diffusing species and reactions at the boundary, as described  
 18 below, and use the result to compute a maximal compartment size for a discretized reaction-diffusion system. We do  
 19 this in two steps – first we treat homogeneous Neumann boundary data to analyze the effect of non-diffusing species,  
 20 and then we summarize the results for other boundary conditions.

#### 21 3.1. Homogeneous Neumann conditions

22 Let  $\Omega \subset R^3$  be a domain with a smooth boundary  $\partial\Omega$ . Denote generic spatial locations as  $\mathbf{x} = (x, y, z)$  and  
 23  $\boldsymbol{\xi} = (\xi_x, \xi_y, \xi_z)$ , where either can lie in the interior of or on the boundary of  $\Omega$ . Let  $u(\mathbf{x}, t) \in R^m$  and  $v(\mathbf{x}, t) \in R^n$   
 24 denote the concentrations of diffusing and non-diffusing species, respectively, that react in  $\Omega$  at time  $t$ , and let  
 25  $w(\mathbf{x}, t) \in R^p$  denote the concentrations of non-diffusing species that do not affect the evolution of any species in  $u$   
 26 and  $v$ . We write the governing equations and the boundary conditions for all species as follows.

$$\begin{aligned}
 27 \quad & \frac{\partial u(\mathbf{x}, t)}{\partial t} = \mathcal{D}\Delta u(\mathbf{x}, t) + \mathcal{R}(u(\mathbf{x}, t), v(\mathbf{x}, t)), & \mathbf{x} \in \Omega \\
 28 \quad & \frac{\partial v(\mathbf{x}, t)}{\partial t} = \mathcal{S}(u(\mathbf{x}, t), v(\mathbf{x}, t)), & \mathbf{x} \in \Omega \\
 29 \quad & \frac{\partial w(\mathbf{x}, t)}{\partial t} = \mathcal{T}(u(\mathbf{x}, t), v(\mathbf{x}, t), w(\mathbf{x}, t)), & \mathbf{x} \in \Omega \\
 30 \quad & \frac{\partial u(\mathbf{x}, t)}{\partial n} = 0, & \mathbf{x} \in \partial\Omega \\
 31 \quad & u(\mathbf{x}, 0) = u_0(\mathbf{x}), & \mathbf{x} \in \Omega \\
 32 \quad & v(\mathbf{x}, 0) = v_0(\mathbf{x}), & \mathbf{x} \in \Omega \\
 33 \quad & w(\mathbf{x}, 0) = w_0(\mathbf{x}), & \mathbf{x} \in \Omega
 \end{aligned}$$

34 Here differentiation with respect to  $n$  is along the outward unit normal vector to  $\Omega$ , and  $\mathcal{D}$  is a diagonal matrix of  
 35 diffusion coefficients. Note that the non-diffusing species in  $w(\mathbf{x}, t)$ , which do not affect any species in  $u$  or  $v$ , can  
 36 be ignored in the following, since they do not affect the convergence to a uniform solution of the diffusing species.  
 37 Hereafter we exclude the governing equation for  $w(\mathbf{x}, t)$  from the system equations, and this leads to the following

1 simplified set of equations that we use for the analysis<sup>3</sup>.

$$\begin{aligned}
\frac{\partial u(\mathbf{x}, t)}{\partial t} &= \mathcal{D}\Delta u(\mathbf{x}, t) + \mathcal{R}(u(\mathbf{x}, t), v(\mathbf{x}, t)), & \mathbf{x} \in \Omega \\
\frac{\partial v(\mathbf{x}, t)}{\partial t} &= \mathcal{S}(u(\mathbf{x}, t), v(\mathbf{x}, t)), & \mathbf{x} \in \Omega \\
\frac{\partial u(\mathbf{x}, t)}{\partial n} &= 0, & \mathbf{x} \in \partial\Omega \\
u(\mathbf{x}, 0) &= u_0(\mathbf{x}), & \mathbf{x} \in \Omega \\
v(\mathbf{x}, 0) &= v_0(\mathbf{x}), & \mathbf{x} \in \Omega
\end{aligned} \tag{31}$$

3 We assume that  $\mathcal{R}$  and  $\mathcal{S}$  are  $C^1$  in the non-negative cone of  $(u, v)$ -space, and in general both are nonlinear functions  
4 of  $(u, v)$ .

5 Define the vector  $c(\mathbf{x}, t) \equiv [u(\mathbf{x}, t)^T, v(\mathbf{x}, t)^T]^T$  of all species concentrations and let  $c_0(\mathbf{x})$  be the initial distri-  
6 bution. For a given  $c_0(\mathbf{x})$ , define  $C_0 = \{c_0(\mathbf{x}) | \mathbf{x} \in \bar{\Omega}\}$ . In the following theorem we assume that the concentration  
7  $c(\mathbf{x}, t)$ , which is the image of  $c_0(\mathbf{x})$  under the mapping defined by the integral representation given in (43) and (44)  
8 is contained in a closed, bounded, convex set  $C_\infty \supseteq C_0$ , for all  $t \in [0, \infty)$ . We define the spatial average of the  
9 concentrations for diffusing species as

$$10 \quad \bar{u}(t) \equiv \frac{1}{|\Omega|} \int_{\Omega} u(\mathbf{x}, t) d\mathbf{x},$$

11 and we further assume that for each  $t > 0$ , there exists a spatially-uniform solution  $\bar{v}(t)$  satisfying

$$\begin{aligned}
\frac{d\bar{v}(t)}{dt} &= \mathcal{S}(\bar{u}(t), \bar{v}(t)), \\
\bar{v}(0) &= \frac{1}{|\Omega|} \int_{\Omega} v_0(\mathbf{x}) d\mathbf{x}.
\end{aligned} \tag{32}$$

13 Given  $\bar{u}(t)$ , the existence and uniqueness of  $\bar{v}$  is guaranteed by the smoothness assumptions, but  $\bar{v}(t)$  may not be the  
14 spatial average of  $v(\mathbf{x}, t)$  when  $\mathcal{S}$  is a nonlinear function of either  $u$  or  $v$ .

15 We define the  $L_2$  norm

$$16 \quad \|f(\mathbf{x}, t)\|_{L_2}^2 \equiv \langle f(\mathbf{x}, t), f(\mathbf{x}, t) \rangle_{L_2} \equiv \int_{\Omega} \|f(\mathbf{x}, t)\|_E^2 d\mathbf{x}$$

17 where  $\|\cdot\|_E$  is the Euclidean matrix norm, and we define  $\bar{c}(t) \equiv [\bar{u}(t)^T, \bar{v}(t)^T]^T$ . Then we say that  $\|c(\mathbf{x}, t) - \bar{c}(t)\|_{L_2} \rightarrow 0$   
18 if  $\|u(\mathbf{x}, t) - \bar{u}(t)\|_{L_2} \rightarrow 0$  and  $\|v(\mathbf{x}, t) - \bar{v}(t)\|_{L_2} \rightarrow 0$ .

19 The Jacobians of a function  $f : R^m \times R^n \rightarrow R^m \times R^n$  with respect to  $u$  and  $v$  are denoted

$$20 \quad D_u f(u, v) \equiv \frac{\partial f(u, v)}{\partial u}, \quad D_v f(u, v) \equiv \frac{\partial f(u, v)}{\partial v}.$$

21 We define measures of kinetic sensitivity via the Jacobians of the reaction terms as follows

$$\begin{aligned}
\hat{r}_u &\equiv \sup_{c \in C_\infty} \|D_u \mathcal{R}(c)\|_E, & \hat{r}_v &\equiv \sup_{c \in C_\infty} \|D_v \mathcal{R}(c)\|_E, \\
\hat{s}_u &\equiv \sup_{c \in C_\infty} \|D_u \mathcal{S}(c)\|_E, & \hat{s}_v &\equiv \inf_{c \in C_\infty} |\sigma(W_c D_v \mathcal{S}(c))|.
\end{aligned} \tag{33}$$

23 The matrix  $W_c$  is defined in the following.

24 For a Hermitian operator  $H$ , let  $\lambda_m(H)$  and  $\lambda_M(H)$  be the smallest and the largest eigenvalues of  $H$ , which are  
25 defined as

$$26 \quad \lambda_M(H) = \sup_x \{ \langle x, Hx \rangle | \|x\| = 1 \} \quad \text{and} \quad \lambda_m(H) = \inf_x \{ \langle x, Hx \rangle | \|x\| = 1 \}, \tag{34}$$

---

<sup>3</sup> In fact, as we show in an example later, inclusion of such species can lead to an inappropriate estimate of the compartment size, and for this reason we include it in the above.

1 respectively, and define  $\|H\| \equiv \max\{\lambda_M(H), -\lambda_m(H)\}$ . An operator  $H$  is said to be uniformly positive if  $\lambda_m(H) > 0$   
 2 [12], which we denote as  $H \gg 0$ . If  $H \gg 0$ , the norm defined as  $\|x\|_H^2 = \langle x, Hx \rangle$  satisfies

$$3 \quad \lambda_m(H)\|x\|^2 \leq \|x\|_H^2 \leq \lambda_M(H)\|x\|^2 = \|H\|\|x\|^2. \quad (35)$$

4 In case all species are diffusible, the proof of the convergence result in [31] does not require that the spectrum of  
 5 the Jacobian of the kinetics lies in the *LHP*; only that a norm of the Jacobian can be dominated by the diffusion  
 6 terms. However this fails when there are non-diffusing species, and this raises several technical difficulties, even when  
 7 the spectrum of the Jacobian of the kinetics lies in the *LHP* pointwise in time. These are overcome in part by use of  
 8 a time-dependent metric, but this raises additional difficulties. We assume that for all  $c \in C_\infty$ ,

$$9 \quad \sigma(D_v \mathcal{S}(c)) \subset \text{LHP}, \quad (36)$$

10 where  $\sigma(A)$  denotes the spectrum of  $A$ . A generalized Lyapunov theorem (Theorem 5.1 in [12]) states that if  $A$  is  
 11 a real bounded linear operator on a real Hilbert space, and if  $\sigma(A) \subset \text{LHP}$ , then there exists a uniformly positive  
 12 operator  $W_A$  such that  $W_A A \ll 0$ . Thus in view of (36), for each  $t > 0$  there is a uniformly positive operator  $W_c$  such  
 13 that

$$14 \quad W_c D_v \mathcal{S}(c) \ll 0. \quad (37)$$

15 Since  $c$  is time-dependent  $W$  is also, and one cannot apply the standard result which states that for a linear system  
 16 with a real constant matrix  $A$ , the zero solution of

$$17 \quad \frac{dz}{dt} = Az, \quad (38)$$

is exponentially stable if and only if the Lyapunov equation

$$A^T W + W A = -Q$$

has a positive definite solution  $W$  for any positive definite matrix  $Q$  [18]. When  $A$  is time-dependent the spectral  
 condition no longer suffices, but it is known that the fundamental solution  $U(t, s)$  of the nonautonomous version of  
 (38) is exponentially stable if and only if the Lyapunov equation

$$\frac{dW}{dt} = -(A^T(t)W(t) + W(t)A(t) + Q)$$

18 has a solution for any positive definite matrix  $Q$  [37]. Here we do not assume exponential stability, but later we  
 19 require a bound on the derivative with respect to time of  $W_c$ , and one can see that this is equivalent to requiring  
 20 that the Jacobian  $D_v \mathcal{S}(c)$  does not vary too rapidly. We define

$$21 \quad \begin{aligned} \lambda_m &\equiv \inf_{c \in C_\infty} \lambda_m(W_c), \\ \lambda_M &\equiv \sup_{c \in C_\infty} \lambda_M(W_c), \\ w &\equiv \frac{1}{2} \sup_{U \in \hat{C}_\infty} \left\| \frac{\partial W_c}{\partial t} \right\|_E. \end{aligned} \quad (39)$$

22 In the definition of  $w$  the supremum is taken over all solutions  $U = (u(\mathbf{x}, t), v(\mathbf{x}, t), \bar{v}(t))$  where  $(u(\mathbf{x}, t), v(\mathbf{x}, t))$  satisfy  
 23 (31), the third component satisfies (32), and  $(\mathbf{x}, \tau) \in (\Omega, [0, \infty))$ .

24 The following theorem gives conditions for exponential convergence in time of  $c(\mathbf{x}, t)$  to a spatially-uniform  
 25 solution under homogeneous Neumann boundary conditions. In essence the condition requires that the smallest non-  
 26 zero diffusion coefficient should be large enough compared to some function of the kinetic sensitivities,  $\hat{r}_u, \hat{r}_v, \hat{s}_u$ , and  
 27  $\hat{s}_v$ , and of the constants,  $\lambda_m, \lambda_M$ , and  $w$ , defined by the positive operator  $W_c$ .



1 **Theorem 3.**  $\|c(\mathbf{x}, t) - \bar{c}(t)\|_{L_2} \rightarrow 0$  exponentially in  $t$  if

$$\begin{aligned} & (i) \quad \sigma(D_v \mathcal{S}(c)) \subset LHP \text{ for all } c \in C_\infty, \\ & (ii) \quad \check{s}_v > w, \\ & (iii) \quad |\alpha_1 \delta| > \hat{r}_u + \frac{\hat{r}_v \hat{s}_u}{\check{s}_v - w} \cdot \frac{\lambda_M^2}{\lambda_m}, \end{aligned} \tag{40}$$

3 where  $\delta = \min_i D_{ii}$  and  $\alpha_1$  is the largest non-zero eigenvalue of the scalar problem

$$\begin{aligned} \Delta \phi(\mathbf{x}) &= \alpha \phi(\mathbf{x}), \quad \mathbf{x} \in \Omega, \\ \frac{\partial \phi}{\partial n} &= 0, \quad \mathbf{x} \in \partial \Omega. \end{aligned} \tag{41}$$

7 *Proof.* Define the Green's function  $G(\mathbf{x}, \boldsymbol{\xi}, t)$  as the solution of

$$\begin{aligned} \frac{\partial G(\mathbf{x}, \boldsymbol{\xi}, t)}{\partial t} &= \mathcal{D} \Delta G(\mathbf{x}, \boldsymbol{\xi}, t), \quad \mathbf{x} \in \Omega, \\ \frac{\partial G(\mathbf{x}, \boldsymbol{\xi}, t)}{\partial n} &= 0, \quad \mathbf{x} \in \partial \Omega, \\ G(\mathbf{x}, \boldsymbol{\xi}, 0) &= \delta(\mathbf{x}, \boldsymbol{\xi}). \end{aligned} \tag{42}$$

9  $G$  is a diagonal matrix and  $G_{ii}$  represents the  $i^{th}$  diagonal element. Then for  $\mathbf{x} \in \Omega$ ,  $u$  and  $v$  satisfy

$$u(\mathbf{x}, t) = \int_{\Omega} G(\mathbf{x}, \boldsymbol{\xi}, t) u_0(\boldsymbol{\xi}) d\boldsymbol{\xi} + \int_0^t \int_{\Omega} G(\mathbf{x}, \boldsymbol{\xi}, t - \tau) \mathcal{R}(u(\boldsymbol{\xi}, \tau), v(\boldsymbol{\xi}, \tau)) d\boldsymbol{\xi} d\tau, \tag{43}$$

$$v(\mathbf{x}, t) = v_0(\mathbf{x}) + \int_0^t \mathcal{S}(u(\mathbf{x}, \tau), v(\mathbf{x}, \tau)) d\tau. \tag{44}$$

12 The  $i^{th}$  diagonal element of  $G$  satisfies

$$\int_{\Omega} G_{ii}(\mathbf{x}, \boldsymbol{\xi}, t - \tau) d\boldsymbol{\xi} = 1,$$

14 and by defining

$$G_{ii}^0(\mathbf{x}, \boldsymbol{\xi}, t) \equiv G_{ii}(\mathbf{x}, \boldsymbol{\xi}, t) - \frac{1}{|\Omega|}, \tag{45}$$

16 we have

$$\int_{\Omega} G_{ii}^0(\mathbf{x}, \boldsymbol{\xi}, t) d\boldsymbol{\xi} = 0. \tag{46}$$

18 Therefore

$$\begin{aligned} u(\mathbf{x}, t) &= \int_{\Omega} \left( \frac{1}{|\Omega|} + G^0(\mathbf{x}, \boldsymbol{\xi}, t) \right) u_0(\boldsymbol{\xi}) d\boldsymbol{\xi} \\ &\quad + \int_0^t \int_{\Omega} \left( \frac{1}{|\Omega|} + G^0(\mathbf{x}, \boldsymbol{\xi}, t - \tau) \right) \mathcal{R}(u(\boldsymbol{\xi}, \tau), v(\boldsymbol{\xi}, \tau)) d\boldsymbol{\xi} d\tau, \end{aligned} \tag{47}$$

20 and it follows that

$$\bar{u}(t) = \frac{1}{|\Omega|} \int_{\Omega} u_0(\boldsymbol{\xi}) d\boldsymbol{\xi} + \frac{1}{|\Omega|} \int_0^t \int_{\Omega} \mathcal{R}(u(\boldsymbol{\xi}, \tau), v(\boldsymbol{\xi}, \tau)) d\boldsymbol{\xi} d\tau \tag{48}$$

22 and

$$0 = \int_{\Omega} G^0(\mathbf{x}, \boldsymbol{\xi}, t) \bar{u}(0) d\boldsymbol{\xi} + \int_0^t \int_{\Omega} G^0(\mathbf{x}, \boldsymbol{\xi}, t - \tau) \mathcal{R}(\bar{u}(\tau), \bar{v}(\tau)) d\boldsymbol{\xi} d\tau. \tag{49}$$

1 We define

$$2 \quad \Psi(\mathbf{x}, t) \equiv u(\mathbf{x}, t) - \bar{u}(t), \quad \Phi(\mathbf{x}, t) \equiv v(\mathbf{x}, t) - \bar{v}(t),$$

3 and then find that

$$4 \quad \Psi(\mathbf{x}, t) = \int_{\Omega} G^0(\mathbf{x}, \boldsymbol{\xi}, t) \Psi(\boldsymbol{\xi}, 0) d\boldsymbol{\xi} + \int_0^t \int_{\Omega} G^0(\mathbf{x}, \boldsymbol{\xi}, t - \tau) R(\boldsymbol{\xi}, \tau) d\boldsymbol{\xi} d\tau \quad (50)$$

5 where

$$6 \quad R(\boldsymbol{\xi}, \tau) \equiv \mathcal{R}(u(\boldsymbol{\xi}, \tau), v(\boldsymbol{\xi}, \tau)) - \mathcal{R}(\bar{u}(\tau), \bar{v}(\tau)).$$

7 Consider the diffusion problem

$$8 \quad \begin{aligned} \frac{\partial \psi(\mathbf{x}, t)}{\partial t} &= \mathcal{D}_{ii} \Delta \psi(\mathbf{x}, t), & \mathbf{x} \in \Omega, \\ \frac{\partial \psi(\mathbf{x}, t)}{\partial n} &= 0, & \mathbf{x} \in \partial \Omega. \end{aligned} \quad (51)$$

9 The solution of (51) can be written

$$10 \quad \psi(\mathbf{x}, t) = \sum_l a_l \phi_l(\mathbf{x}) e^{\alpha_l \mathcal{D}_{ii} t}$$

11 where the eigenvalues  $\alpha_l$  are non-positive and the  $\phi_l(\mathbf{x})$ 's are the corresponding orthonormal eigenfunctions. For those  
12 having zero mean over  $\Omega$ ,  $\alpha_l < 0$  and  $\psi(\mathbf{x}, t)$  satisfies

$$13 \quad \begin{aligned} \frac{d}{dt} \|\psi\|_{L_2}^2 &= \frac{d}{dt} \langle \psi, \psi \rangle_{L_2} = 2 \left\langle \psi, \frac{\partial \psi}{\partial t} \right\rangle_{L_2} \\ 14 \quad &= 2 \langle \psi, \mathcal{D}_{ii} \Delta \psi \rangle_{L_2} \leq 2\alpha_1 D_{ii} \|\psi\|_{L_2}^2 \end{aligned}$$

15 where  $\alpha_1$  is the largest non-zero eigenvalue. It follows that  $\psi(\mathbf{x}, t)$  satisfies

$$16 \quad \|\psi(\mathbf{x}, t)\|_{L_2} \leq e^{-|\alpha_1| \mathcal{D}_{ii} t} \|\psi(\mathbf{x}, 0)\|_{L_2}. \quad (52)$$

The Green's function defined at (42) has an eigenfunction expansion in terms of the  $\phi_l$ 's, and from this it follows that the first term in (50) satisfies the inequality

$$\left\| \int_{\Omega} G^0(\mathbf{x}, \boldsymbol{\xi}, t) \Psi(\boldsymbol{\xi}, 0) d\boldsymbol{\xi} \right\|_{L_2} \leq e^{-|\alpha_1| \delta t} \|\Psi(\mathbf{x}, 0)\|_{L_2}.$$

17 Using this, and the inequality

$$18 \quad \|\mathcal{R}(u(\boldsymbol{\xi}, \tau), v(\boldsymbol{\xi}, \tau)) - \mathcal{R}(\bar{u}(\tau), \bar{v}(\tau))\|_{L_2} \leq \hat{r}_u \|\Psi(\mathbf{x}, \tau)\|_{L_2} + \hat{r}_v \|\Phi(\mathbf{x}, \tau)\|_{L_2}, \quad (53)$$

19 it follows from (50) that

$$20 \quad \begin{aligned} e^{|\alpha_1| \delta t} \|\Psi(\mathbf{x}, t)\|_{L_2} &\leq \|\Psi(\mathbf{x}, 0)\|_{L_2} \\ 21 \quad &+ \hat{r}_u \int_0^t e^{|\alpha_1| \delta \tau} \|\Psi(\mathbf{x}, \tau)\|_{L_2} d\tau + \hat{r}_v \int_0^t e^{|\alpha_1| \delta \tau} \|\Phi(\mathbf{x}, \tau)\|_{L_2} d\tau. \end{aligned} \quad (54)$$

22 From (32), we have

$$23 \quad \bar{v}(t) = \bar{v}(0) + \int_0^t \mathcal{S}(\bar{u}(\tau), \bar{v}(\tau)) d\tau \quad (55)$$

1 and therefore the difference  $\Phi(\mathbf{x}, t) \equiv v(\mathbf{x}, t) - \bar{v}(t)$  satisfies

$$2 \quad \Phi(\mathbf{x}, t) = \Phi(\mathbf{x}, 0) + \int_0^t \{S(u(\mathbf{x}, \tau), v(\mathbf{x}, \tau)) - S(\bar{u}(\tau), \bar{v}(\tau))\} d\tau. \quad (56)$$

3 Since  $C_\infty$  is convex, for each  $\mathbf{x} \in \Omega$  and  $\tau \in [0, t < \infty)$ , there exist  $c_1(\mathbf{x}, \tau)$  and  $c_2(\mathbf{x}, \tau)$  such that

$$4 \quad S(u(\mathbf{x}, \tau), v(\mathbf{x}, \tau)) - S(\bar{u}(\tau), \bar{v}(\tau)) = D_u S(c_1(\mathbf{x}, \tau)) (u(\mathbf{x}, \tau) - \bar{u}(\tau)) + D_v S(c_2(\mathbf{x}, \tau)) (v(\mathbf{x}, \tau) - \bar{v}(\tau)).$$

5 As a result, (56) can be rewritten as

$$6 \quad \Phi(\mathbf{x}, t) = \Phi(\mathbf{x}, 0) + \int_0^t \{D_u S(c_1(\mathbf{x}, \tau)) \Psi(\mathbf{x}, \tau) + D_v S(c_2(\mathbf{x}, \tau)) \Phi(\mathbf{x}, \tau)\} d\tau. \quad (57)$$

7 In view of (36) and (37), for each  $c \in C_\infty$  we can define a weighted Euclidean norm by  $W_c$  as

$$8 \quad \|A\|_{E, W_c}^2 \equiv \langle A, W_c A \rangle_E.$$

9 Since  $W_c$  is a uniformly positive operator for each  $c \in C_\infty$ ,  $\lambda_m > 0$  in (39). Using (35) and (39), for each  $\mathbf{x} \in \Omega$  and  
10  $t \in [0, \infty)$ , we obtain the upper and lower bounds

$$11 \quad \lambda_m \|\Phi(\mathbf{x}, t)\|_E^2 \leq \|\Phi(\mathbf{x}, t)\|_{E, W_{c_2(\mathbf{x}, t)}}^2 \leq \lambda_M \|\Phi(\mathbf{x}, t)\|_E^2. \quad (58)$$

12 Define a weighted  $L_2$  norm as

$$13 \quad \|f(\mathbf{x}, t)\|_{L_2, W_f}^2 \equiv \int_\Omega \|f(\mathbf{x}, t)\|_{E, W_{f(\mathbf{x}, t)}}^2 d\mathbf{x}.$$

14 Then,

$$15 \quad \lambda_m \|\Phi(\mathbf{x}, t)\|_{L_2}^2 \leq \|\Phi(\mathbf{x}, t)\|_{L_2, W_{c_2}}^2 \leq \lambda_M \|\Phi(\mathbf{x}, t)\|_{L_2}^2, \quad (59)$$

16 and differentiating this we obtain

$$17 \quad \begin{aligned} \frac{d}{dt} \|\Phi(\mathbf{x}, t)\|_{E, W_{c_2(\mathbf{x}, t)}}^2 &= \langle \Phi(\mathbf{x}, t), D_t W_{c_2(\mathbf{x}, t)} \Phi(\mathbf{x}, t) \rangle_E + \langle \Phi(\mathbf{x}, t), W_{c_2(\mathbf{x}, t)} D_t \Phi(\mathbf{x}, t) \rangle_E \\ &+ \langle D_t \Phi(\mathbf{x}, t), W_{c_2(\mathbf{x}, t)} \Phi(\mathbf{x}, t) \rangle_E. \end{aligned} \quad (60)$$

18 From (57) and (60) it follows that

$$19 \quad \begin{aligned} \frac{d}{dt} \|\Phi(\mathbf{x}, t)\|_{E, W_{c_2(\mathbf{x}, t)}}^2 &= \langle \Phi(\mathbf{x}, t), D_t W_{c_2(\mathbf{x}, t)} \Phi(\mathbf{x}, t) \rangle_E \\ 20 \quad &+ \langle \Phi(\mathbf{x}, t), W_{c_2(\mathbf{x}, t)} D_u S(c_1(\mathbf{x}, \tau)) \Psi(\mathbf{x}, t) \rangle_E + \langle \Phi(\mathbf{x}, t), W_{c_2(\mathbf{x}, t)} D_v S(c_2(\mathbf{x}, \tau)) \Phi(\mathbf{x}, t) \rangle_E \\ 21 \quad &+ \langle D_u S(c_1(\mathbf{x}, \tau)) \Psi(\mathbf{x}, t), W_{c_2(\mathbf{x}, t)} \Phi(\mathbf{x}, t) \rangle_E + \langle D_v S(c_2(\mathbf{x}, \tau)) \Phi(\mathbf{x}, t), W_{c_2(\mathbf{x}, t)} \Phi(\mathbf{x}, t) \rangle_E. \end{aligned}$$

22 After applying the Cauchy-Schwarz inequality and using the fact that  $\sigma(W_c D_v S(c)) \subset LHP$  for all  $c \in C_\infty$ , we  
23 obtain

$$24 \quad \begin{aligned} \frac{d}{dt} \|\Phi(\mathbf{x}, t)\|_{E, W_{c_2(\mathbf{x}, t)}}^2 &\leq 2w \|\Phi(\mathbf{x}, t)\|_E^2 + 2\hat{s}_u \lambda_M \|\Phi(\mathbf{x}, t)\|_E \|\Psi(\mathbf{x}, t)\|_E \\ 25 \quad &+ \left\langle \Phi(\mathbf{x}, t), \left( W_{c_2(\mathbf{x}, t)} D_v S(c_2(\mathbf{x}, \tau)) + [D_v S(c_2(\mathbf{x}, \tau))]^T W_{c_2(\mathbf{x}, t)} \right) \Phi(\mathbf{x}, t) \right\rangle_E \\ 26 \quad &\leq 2\hat{s}_u \lambda_M \|\Phi(\mathbf{x}, t)\|_E \|\Psi(\mathbf{x}, t)\|_E - (2\check{s}_v - 2w) \|\Phi(\mathbf{x}, t)\|_E^2 \end{aligned} \quad (61)$$

1 where  $[D_v \mathcal{S}(c_2(\mathbf{x}, \tau))]^T$  is the transpose of  $D_v \mathcal{S}(c_2(\mathbf{x}, \tau))$  for fixed  $\mathbf{x} \in \Omega$  and  $\tau \in [0, \infty)$ . From the second condition  
 2 in (40), we assume that  $\check{s}_v - w > 0$ . Then using (58) and dividing by  $2\|\Phi(\mathbf{x}, t)\|_{E, W_{c_2}(\mathbf{x}, t)}$ , (61) becomes

$$3 \quad \frac{d}{dt} \|\Phi(\mathbf{x}, t)\|_{E, W_{c_2}(\mathbf{x}, t)} \leq \frac{\hat{s}_u \lambda_M}{\sqrt{\lambda_m}} \|\Psi(\mathbf{x}, t)\|_E - \frac{\check{s}_v - w}{\lambda_M} \|\Phi(\mathbf{x}, t)\|_{E, W_{c_2}(\mathbf{x}, t)}. \quad (62)$$

4 From this one obtains

$$5 \quad e^{|\alpha_1 \delta| t} \|\Phi(\mathbf{x}, t)\|_{E, W_{c_2}(\mathbf{x}, t)} \leq \|\Phi(\mathbf{x}, 0)\|_{E, W_{c_2}(\mathbf{x}, 0)} + \frac{\hat{s}_u \lambda_M}{\sqrt{\lambda_m}} \int_0^t e^{|\alpha_1 \delta| \tau} \|\Psi(\mathbf{x}, \tau)\|_E d\tau \\ 6 \quad + \left( |\alpha_1 \delta| - \frac{\check{s}_v - w}{\lambda_M} \right) \int_0^t e^{|\alpha_1 \delta| \tau} \|\Phi(\mathbf{x}, \tau)\|_{E, W_{c_2}(\mathbf{x}, \tau)} d\tau \quad (63)$$

7 which the reader can verify by integration by parts. Integrating (63) over  $\mathbf{x} \in \Omega$ , we get

$$8 \quad e^{|\alpha_1 \delta| t} \|\Phi(\mathbf{x}, t)\|_{L_2, W_{c_2}} \leq \|\Phi(\mathbf{x}, 0)\|_{L_2, W_{c_2}} + \frac{\hat{s}_u \lambda_M}{\sqrt{\lambda_m}} \int_0^t e^{|\alpha_1 \delta| \tau} \|\Psi(\mathbf{x}, \tau)\|_{L_2} d\tau \\ 9 \quad + \left( |\alpha_1 \delta| - \frac{\check{s}_v - w}{\lambda_M} \right) \int_0^t e^{|\alpha_1 \delta| \tau} \|\Phi(\mathbf{x}, \tau)\|_{L_2, W_{c_2}} d\tau. \quad (64)$$

10 We now have the estimates on the diffusing and non-diffusing species needed, and we combine these as follows.

11 Define

$$12 \quad g(t) \equiv \left[ e^{|\alpha_1 \delta| t} \|\Psi(\mathbf{x}, t)\|_{L_2}, \quad e^{|\alpha_1 \delta| t} \|\Phi(\mathbf{x}, t)\|_{L_2, W_{c_2}} \right]^T;$$

13 then from (54) and (64) we have

$$14 \quad g(t) \leq g(0) + \int_0^t \begin{bmatrix} \hat{r}_u & \frac{\hat{r}_v}{\sqrt{\lambda_m}} \\ \frac{\hat{s}_u \lambda_M}{\sqrt{\lambda_m}} & |\alpha_1 \delta| - \frac{\check{s}_v - w}{\lambda_M} \end{bmatrix} g(\tau) d\tau,$$

where the inequality is defined componentwise. Since the off-diagonal elements of the matrix  $K$  in the integral are positive, the map defined by the right-hand side preserves the order in the positive cone of  $R^2$ [40], and it follows that

$$g(t) \leq e^{Kt} g(0).$$

15 Therefore

$$16 \quad \begin{bmatrix} \|\Psi(\mathbf{x}, t)\|_{L_2} \\ \|\Phi(\mathbf{x}, t)\|_{L_2, W_{c_2}} \end{bmatrix} \leq e^{At} \begin{bmatrix} \|\Psi(\mathbf{x}, 0)\|_{L_2} \\ \|\Phi(\mathbf{x}, 0)\|_{L_2, W_{c_2}} \end{bmatrix}$$

17 where

$$18 \quad A = \begin{bmatrix} -|\alpha_1 \delta| + \hat{r}_u & \frac{\hat{r}_v}{\sqrt{\lambda_m}} \\ \frac{\hat{s}_u \lambda_M}{\sqrt{\lambda_m}} & -\frac{\check{s}_v - w}{\lambda_M} \end{bmatrix}.$$

19 Consequently, if all eigenvalues of  $A$  have negative real part, then it follows, after using (59), that  $\|c(\mathbf{x}, t) - \bar{c}(t)\|_{L_2} \rightarrow 0$   
 20 exponentially in  $t$ .

21 We can compute the characteristic equation explicitly and find the factored form

$$22 \quad (\lambda + |\alpha_1 \delta| - \hat{r}_u) \left( \lambda + \frac{\check{s}_v - w}{\lambda_M} \right) - \frac{\hat{r}_v}{\sqrt{\lambda_m}} \frac{\hat{s}_u \lambda_M}{\sqrt{\lambda_m}} = 0.$$

23 Therefore the  $\lambda$ 's are negative if

$$24 \quad (|\alpha_1 \delta| - \hat{r}_u) + \frac{\check{s}_v - w}{\lambda_M} > 0 \\ (|\alpha_1 \delta| - \hat{r}_u) \frac{\check{s}_v - w}{\lambda_M} - \frac{\hat{r}_v \hat{s}_u \lambda_M}{\lambda_m} > 0. \quad (65)$$

1 Since we assume that  $\hat{s}_v > w$ , these conditions are satisfied if

$$2 \quad |\alpha_1 \delta| > \hat{r}_u + \frac{\hat{r}_v \hat{s}_u}{\hat{s}_v - w} \cdot \frac{\lambda_M^2}{\lambda_m}, \quad (66)$$

3 which then guarantees exponential in  $t$  convergence to the uniform solution. This proves the theorem.  $\square$

4 *Remark 1.* If all species diffuse, the criterion in (66) reduces to

$$5 \quad |\alpha_1 \delta| > \hat{r}_u,$$

6 since  $\mathcal{S}(u(\mathbf{x}, t), v(\mathbf{x}, t)) = 0$ . This is the condition given in [3].

*Remark 2.* If  $\Omega = [0, h_x] \times [0, h_y] \times [0, h_z]$ ,

$$\alpha_i \mathcal{D} = - \left( \frac{l_1 \pi}{h_x} \right)^2 \mathcal{D}_x - \left( \frac{l_2 \pi}{h_y} \right)^2 \mathcal{D}_y - \left( \frac{l_3 \pi}{h_z} \right)^2 \mathcal{D}_z$$

7 for  $l_1, l_2, l_3 = 0, 1, \dots$  where  $\mathcal{D}_x$ ,  $\mathcal{D}_y$ , and  $\mathcal{D}_z$  are diffusion matrices of diffusing species in  $x$ -,  $y$ -, and  $z$ - directions. If  
8 diffusion is isotropic, we have

$$9 \quad |\alpha_1 \delta| = \min \left\{ \left( \frac{\pi}{h_x} \right)^2, \left( \frac{\pi}{h_y} \right)^2, \left( \frac{\pi}{h_z} \right)^2 \right\} \min_i \mathcal{D}_{ii}.$$

10 If one adopts the criterion that the largest computational cell size for a stochastic simulation of a reaction-diffusion  
11 system should be small enough that the cell is uniform on time scales relevant to the coupled deterministic reaction  
12 network, one can apply the criterion in (66) to predict the minimum number of computational cells. We do this in  
13 detail for Example 2.2, as later shown in Example 4.1. There we set  $W_c = 1$  since  $D_v \mathcal{S}$  is a scalar, and therefore  
14  $\lambda_m = \lambda_M = 1$  and  $w = 0$  in (66).

### 15 3.2. Other types of boundary conditions

16 The convergence result in Theorem 3 can be extended in several ways. For example, when there are no non-diffusing  
17 species, exponential convergence to a uniform state under homogeneous Robin boundary conditions on  $\partial\Omega$  was proven  
18 in [10]. That result can be extended to allow homogeneous Robin conditions on  $\partial\Omega_1 \subset \partial\Omega$  and homogeneous Neumann  
19 data on  $\partial\Omega/\partial\Omega_1$ , since it is easy to show that the principal eigenvalue for the spectral problem with mixed data is  
20 strictly less than zero.

21 When there are non-diffusing species, the Neumann conditions in (31) are replaced by

$$22 \quad \begin{aligned} \mathcal{D} \frac{\partial u(\mathbf{x}, t)}{\partial n} &= -\mathcal{B}u(\mathbf{x}, t), & \mathbf{x} \in \partial\Omega_1, \\ \frac{\partial u(\mathbf{x}, t)}{\partial n} &= 0, & \mathbf{x} \in \partial\Omega/\partial\Omega_1. \end{aligned} \quad (67)$$

23 We assume that  $\mathcal{B}$  is a diagonal matrix with positive diagonal elements, and thus the reactions represent degradation  
24 or sequestration on the boundary, or transfer through the boundary. In any case, the flux of diffusing species to  $\partial\Omega_1$   
25 balances reactions of diffusing species on  $\partial\Omega_1$ .

We could allow a slightly more general boundary condition by replacing the term  $-\mathcal{B}u(\mathbf{x}, t)$  in (67) with  $-\mathcal{B}(u(\mathbf{x}, t) - u^s)$ , where  $u^s$  is a solution of

$$\mathcal{R}(u^s, v^s) = 0 \quad \text{and} \quad \mathcal{S}(u^s, v^s) = 0,$$

26 but by translating the steady state, we can assume that  $(u^s, v^s) = (0, 0)$ , which we do hereafter.

27 We then have the following.

**Theorem 4.** Consider the system (31) with boundary conditions replaced by (67). Further, suppose that the system admits the uniform steady state  $(u^s, v^s) = (0, 0)$ . Let  $\alpha_{i1}$  be the principal eigenvalue of the scalar problem

$$\begin{aligned} \Delta\phi(\mathbf{x}) &= \alpha_i\phi(\mathbf{x}), & \mathbf{x} \in \Omega, \\ \frac{\partial\phi}{\partial n} &= \frac{\mathcal{B}_{ii}}{\mathcal{D}_{ii}}\phi, & \mathbf{x} \in \partial\Omega_1, \\ \frac{\partial\phi}{\partial n} &= 0, & \mathbf{x} \in \partial\Omega/\partial\Omega_1, \end{aligned} \quad (68)$$

and suppose that

$$\begin{aligned} (i) \quad & \sigma(D_v\mathcal{S}(c)) \subset LHP \text{ for all } c \in C_\infty, \\ (ii) \quad & \check{s}_v > w, \\ (iii) \quad & \delta_1 > \hat{r}_u + \frac{\hat{r}_v \hat{s}_u}{\check{s}_v - w} \cdot \frac{\lambda_M^2}{\lambda_m}, \end{aligned}$$

where  $\delta_1 \equiv \min_i(|\alpha_{i1}|/\mathcal{D}_{ii})$ . Then  $\|c(\mathbf{x}, t)\|_{L_2} \rightarrow 0$  exponentially in  $t$ .

*Proof.* The proof follows along the same lines as that of Theorem 3. The equations are cast into integral form using a Green's function for the Robin problem. From the analog of (51) one finds that since the principal eigenvalue  $\alpha_{i1} < 0$  [41],  $\psi(\mathbf{x}, t)$  satisfies the inequality

$$\|\psi(\mathbf{x}, t)\|_{L_2} \leq e^{-|\alpha_{i1}|\mathcal{D}_{ii}t} \|\psi(\mathbf{x}, 0)\|_{L_2}, \quad (69)$$

where  $\alpha_{i1}$  is the largest eigenvalue of the scalar problem analogous to (51). This leads to the estimates

$$\begin{aligned} e^{\delta_1 t} \|u(\mathbf{x}, t)\|_{L_2} &\leq \|u_0(\mathbf{x})\|_{L_2} + \hat{r}_u \int_0^t e^{\delta_1 \tau} \|u(\mathbf{x}, \tau)\|_{L_2} d\tau \\ &\quad + \frac{\hat{r}_v}{\sqrt{\lambda_m}} \int_0^t e^{\delta_1 \tau} \|v(\mathbf{x}, \tau)\|_{L_2, W_{c_4}} d\tau, \end{aligned} \quad (70)$$

$$\begin{aligned} e^{\delta_1 t} \|v(\mathbf{x}, t)\|_{L_2, W_{c_4}} &\leq \|v_0(\mathbf{x})\|_{L_2, W_{c_4}} + \frac{\hat{s}_u \lambda_M}{\sqrt{\lambda_m}} \int_0^t e^{\delta_1 \tau} \|u(\mathbf{x}, \tau)\|_{L_2} d\tau \\ &\quad + \left( \delta_1 - \frac{\check{s}_v - w}{\lambda_M} \right) \int_0^t e^{\delta_1 \tau} \|v(\mathbf{x}, \tau)\|_{L_2, W_{c_4}} d\tau, \end{aligned} \quad (71)$$

and from these it follows that  $\|c(\mathbf{x}, t)\|_{L_2} \rightarrow 0$  exponentially in  $t$  under the hypotheses in the statement of the theorem. Here  $W_{c_4}$  in (70) and (71) is analogous to  $W_{c_2}$  in (64). This is used to estimate  $\mathcal{S}(u(\mathbf{x}, \tau), v(\mathbf{x}, \tau))$ , rather than  $\mathcal{S}(u(\mathbf{x}, \tau), v(\mathbf{x}, \tau)) - \mathcal{S}(\bar{u}(\tau), \bar{v}(\tau))$  as in Theorem 3. The functions  $c_3$  and  $c_4$  are defined as follows: for each  $\mathbf{x} \in \Omega$  and  $\tau$  where  $0 \leq \tau \leq t < \infty$ , there exist  $c_3(\mathbf{x}, \tau)$  in  $(0, u(\mathbf{x}, \tau))$  and  $c_4(\mathbf{x}, \tau)$  in  $(0, v(\mathbf{x}, \tau))$  satisfying

$$\mathcal{S}(u(\mathbf{x}, \tau), v(\mathbf{x}, \tau)) = D_u \mathcal{S}(c_3(\mathbf{x}, \tau))u(\mathbf{x}, \tau) + D_v \mathcal{S}(c_4(\mathbf{x}, \tau))v(\mathbf{x}, \tau).$$

The details of the proof are left to the reader.  $\square$

Other types of boundary conditions can be treated under suitable constraints on the boundary values. For example, if non-homogeneous Dirichlet conditions of the form  $u(\mathbf{x}, t) = u^B$  for  $\mathbf{x} \in \partial\Omega$  are imposed, then a uniform steady state exists only if there is a constant vector  $v^s$  such that

$$\mathcal{R}(u^B, v^s) = 0 \quad \text{and} \quad \mathcal{S}(u^B, v^s) = 0.$$

In other words, the boundary values must coincide with the  $u$ -component of a constant, spatially-uniform steady state of the governing reaction-diffusion equations. In this case, as in the above, the proof of convergence to a uniform solution follows that from Robin data, since the principal eigenvalue of the associated spectral problem is negative [41].

## 4. Applications

In this section we apply the results in Section 2 and 3 to Example 2.2 and to a simple nonlinear reaction-diffusion system with the bimolecular reaction  $2A \rightleftharpoons C$ . We compute  $\overline{CV}^*$  as a function of  $N_c$  and determine at what number of compartments,  $\overline{CV}^*$  stabilizes, so as to determine the number of compartments in the stochastic model. We also compute the upper bound for the computational cell size by applying Theorem 3 to obtain the minimal number of compartments, and we compare this number to the one determined by  $\overline{CV}^*$ . Erban and Chapman [15] also analyzed a system with bimolecular reaction,  $2A \rightarrow C$ , but their reaction is irreversible and there is production of  $A$  from a source in their model. They consider both the compartment-based model using a master equation and the molecular-based model using stochastic differential equations. We consider a multi-compartment model to illustrate another aspect of how the criterion of Theorem 3 can be used.

### 4.1. Example 2.2 revisited

Consider the reactions involving species  $A$ ,  $B$ , and  $C$  in Example 2.2 with corresponding deterministic rate constants given in Table 3. Species  $A$  can diffuse, species  $B$  and  $C$  do not, and species  $C$  does not affect species  $A$  and  $B$ . Let  $u(x, t)$ ,  $v(x, t)$ , and  $w(x, t)$  be the concentrations of species  $A$ ,  $B$ , and  $C$ , respectively. Since  $C$  does not affect the dynamics of  $A$  or  $B$ , we first consider the system with only  $A$  and  $B$ . The domain is  $\Omega = [0, h_x]$  and  $\partial\Omega = \{0, h_x\}$ .  $u(x, t)$  and  $v(x, t)$  satisfy (31) with

$$\begin{aligned} \mathcal{R}(u(x, t), v(x, t)) &= \begin{bmatrix} -(k_5 c_R + k_6) & k_{-5} \\ k_5 c_R & -(k_{-5} + k_7) \end{bmatrix} \begin{bmatrix} u(x, t) \\ v(x, t) \end{bmatrix}, \\ \mathcal{S}(u(x, t), v(x, t)) &= \begin{bmatrix} k_5 c_R & -(k_{-5} + k_7) \end{bmatrix} \begin{bmatrix} u(x, t) \\ v(x, t) \end{bmatrix}, \\ \mathcal{D} &= D. \end{aligned}$$

Since  $D_v \mathcal{S}$  is a scalar, we may set  $W_c = 1$ . Then, the variables defined by Jacobian of the reaction terms are found to be

$$\hat{r}_u = k_5 c_R + k_6, \quad \hat{r}_v = k_{-5}, \quad \hat{s}_u = k_5 c_R, \quad \hat{s}_v = k_{-5} + k_7.$$

After computing the convergence criterion given in Theorem 3, we obtain the following upper bound for the computational cell size that guarantees convergence to a spatially-uniform solution in the deterministic calculation.

$$h_x < \sqrt{\frac{D\pi^2}{\left(\frac{k_5 c_R (2k_{-5} + k_7)}{k_{-5} + k_7} + k_6\right)}} \approx 8.25 \mu m$$

We then use  $h_x$  as the compartment size for the corresponding stochastic model, apply the fact that  $h_x = L_x/N_c$ , and replace the deterministic reaction rate constants by stochastic reaction rate constants. As a result we obtain the following lower bound for the number of compartments,

$$N_c > \sqrt{\frac{\left(\frac{\kappa_5 R (2\kappa_{-5} + \kappa_7)}{\kappa_{-5} + \kappa_7} + \kappa_6\right) L_x^2}{D\pi^2}} \approx 33. \quad (72)$$

Comparing this lower bound for  $N_c$  to the number at which the  $\overline{CV}^*$  in Figure 6(a) stabilizes, we conclude that (72) gives a good estimate for the maximum allowable compartment size. This shows that Theorem 3, applied appropriately, provides a rational basis for computing the minimum number of compartments. However, some thought is required, as the following illustrates.

1 Suppose that we had not recognized that  $C$  does not affect the upstream reactions. Now  $w$  becomes  $v_2$  and the  
2 reaction matrices are

$$3 \quad \mathcal{R}(u(x,t), v(x,t)) = \begin{bmatrix} -(k_5 c_R + k_6) & k_{-5} & 0 \end{bmatrix} \begin{bmatrix} u(x,t) \\ v_1(x,t) \\ v_2(x,t) \end{bmatrix},$$

$$4 \quad \mathcal{S}(u(x,t), v(x,t)) = \begin{bmatrix} k_5 c_R & -(k_{-5} + k_7) & 0 \\ 0 & k_7 & -k_9 \end{bmatrix} \begin{bmatrix} u(x,t) \\ v_1(x,t) \\ v_2(x,t) \end{bmatrix},$$

$$5 \quad \mathcal{D} = D.$$

6 Since  $\sigma\left(\frac{(D_v \mathcal{S}(c) + [D_v \mathcal{S}(c)]^T)}{2}\right) \subset LHP$  for all  $c \in C_\infty$ , we can take  $W_c$  to be the identity matrix. The sensitivities  
7 defined by the Jacobians of the reaction terms are found to be

$$8 \quad \hat{r}_u = k_5 c_R + k_6, \quad \hat{r}_v = k_{-5},$$

$$9 \quad \hat{s}_u = k_5 c_R, \quad \hat{s}_v = \frac{(k_{-5} + k_7 + k_9) - \sqrt{(k_{-5} + k_7 - k_9)^2 + k_7^2}}{2}.$$

10 Using  $\alpha_l = -\left(\frac{l\pi}{h_x}\right)^2$  and applying Theorem 3, we find that

$$11 \quad h_x < \sqrt{\frac{D\pi^2}{\left(\frac{k_5 c_R (k_{-5} + \hat{s}_v)}{\hat{s}_v} + k_6\right)}} \approx 1.41 \mu m.$$

12 Treating  $[0, h_x]$  as one compartment, we obtain a lower bound for the number of compartments,

$$13 \quad N_c > \sqrt{\frac{\left(\frac{\kappa_5 R (\kappa_{-5} + \hat{s}_v)}{\hat{s}_v} + \kappa_6\right) L_x^2}{D\pi^2}} \approx 195. \quad (73)$$

14 Comparing (72) and (73), we see that the condition in Theorem 3 over estimates a lower bound for the number of  
15 compartments if we include the non-diffusing species which do not affect the diffusing species. The difference between  
16 (72) and (73) arises from the fact that  $\kappa_{-5} + \kappa_7 \approx 2.03 \text{ min}^{-1}$  in (72), whereas  $\hat{s}_v \approx 0.03 \text{ min}^{-1}$  appears in (73).  
17 The difference reflects the fact that the relaxation time of the non-diffusible species (more generally, a measure of the  
18 sensitivity of the non-diffusible species) enters into the calculation of the overall relaxation time, and this is much  
19 longer when  $C$  is included. Clearly if downstream uncoupled components relax faster than upstream components  
20 their inclusion will not have this effect.

#### 21 4.2. Why network properties must be considered

22 Condition (i) in Theorem 3 is very important, since without it diffusive instabilities (such as the Turing instability)  
23 can arise [31, 32]. In fact, if a self-activating species is also immobile, instabilities of arbitrarily-short wavelengths can  
24 arise [34], which precludes establishing a minimum non-zero compartment size. To illustrate some of the effects of  
25 violating this condition, we investigate a simple reaction-diffusion network which does not satisfy this condition.

26 *Example 1.* Consider a model for the glycolytic reactions, which involves positive feedback and leads to a generic  
27 back-activation oscillator [3]. The kinetic steps are



29 and assuming that the enzyme is far from saturation, these can be described via polynomials. To illustrate some  
30 of the problems as simply as possible, we consider a two-compartment model, since the deterministic case for that  
31 system has been analyzed in detail [3]. We assume that  $Y$  does not diffuse and therefore the governing equations are

$$32 \quad \frac{du_1(t)}{dt} = \frac{D_x}{(L_x/2)^2} (u_2(t) - u_1(t)) + k_0 - k_1 u_1(t) v_1(t)^2 - k_2 u_1(t),$$



**Table 4.** Deterministic and stochastic parameters in Example 1

Deterministic		Stochastic	
$k_0$	$5.25 \text{ nM min}^{-1}$	$\kappa_0$	$k_0 \times (\mathcal{N}_A V_c) \approx 158.22 \text{ min}^{-1}$
$k_1$	$1 \text{ nM}^{-2} \text{ min}^{-1}$	$\kappa_1$	$k_1 / (\mathcal{N}_A V_c)^2 \approx 0.0011 \text{ min}^{-1}$
$k_2$	$0.005 \text{ min}^{-1}$	$\kappa_2$	$k_2 = 0.005 \text{ min}^{-1}$
$k_3$	$5 \text{ min}^{-1}$	$\kappa_3$	$k_3 = 5 \text{ min}^{-1}$
$D_x$	$1000 \mu\text{m}^2 \text{ min}^{-1}$	$\kappa_{d,x}$	$D_x / (L_x/2)^2 = 0.4 \text{ min}^{-1}$
$u_1(0)$	$2 \text{ nM}$	$N^{(1,1)}(0)$	$2 \text{ nM} \times \frac{6.022 \times 10^{-1} \mu\text{m}^{-3}}{1 \text{ nM}} \times 50 \mu\text{m}^3 \approx 60$
$u_2(0)$	$2 \text{ nM}$	$N^{(2,1)}(0)$	$2 \text{ nM} \times \frac{6.022 \times 10^{-1} \mu\text{m}^{-3}}{1 \text{ nM}} \times 50 \mu\text{m}^3 \approx 60$
$v_1(0)$	$7 \text{ nM}$	$N^{(1,2)}(0)$	$7 \text{ nM} \times \frac{6.022 \times 10^{-1} \mu\text{m}^{-3}}{1 \text{ nM}} \times 50 \mu\text{m}^3 \approx 210$
$v_2(0)$	$0 \text{ nM}$	$N^{(2,2)}(0)$	$0 \text{ nM} \times \frac{6.022 \times 10^{-1} \mu\text{m}^{-3}}{1 \text{ nM}} \times 50 \mu\text{m}^3 \approx 0$
$L_x$	$100 \mu\text{m}$		
$L_y$	$1 \mu\text{m}$		
$L_z$	$1 \mu\text{m}$		
$\mathbf{V}$	$100 \mu\text{m}^3$		

$$\frac{dv_1(t)}{dt} = k_1 u_1(t) v_1(t)^2 + k_2 u_1(t) - k_3 v_1(t),$$

$$\frac{du_2(t)}{dt} = \frac{D_x}{(L_x/2)^2} (u_1(t) - u_2(t)) + k_0 - k_1 u_2(t) v_2(t)^2 - k_2 u_2(t),$$

$$\frac{dv_2(t)}{dt} = k_1 u_2(t) v_2(t)^2 + k_2 u_2(t) - k_3 v_2(t),$$

where  $u_k(t)$  and  $v_k(t)$  are concentrations of species  $X$  and  $Y$  in the  $k^{\text{th}}$  compartment at time  $t$  for  $k = 1, 2$ . The term  $k_1 u_k(t) v_k(t)^2$  comes from the positive feedback step in which  $Y$  activates its production. The uniform steady state is

$$u^* = u_1^* = u_2^* = \frac{\frac{k_0}{k_3}}{\frac{k_2}{k_3} + \left(\frac{k_0}{k_3} \sqrt{\frac{k_1}{k_3}}\right)^2},$$

$$v^* = v_1^* = v_2^* = \frac{k_0}{k_3}.$$

The diffusion matrix and the Jacobian of the kinetic terms are

$$\mathcal{D} = \begin{bmatrix} \frac{D_x}{(L_x/2)^2} & 0 \\ 0 & 0 \end{bmatrix},$$

$$\mathcal{K}(u_k(t), v_k(t)) = \begin{bmatrix} -(k_2 + k_1 v_k(t)^2) & -2k_1 u_k(t) v_k(t) \\ k_2 + k_1 v_k(t)^2 & 2k_1 u_k(t) v_k(t) - k_3 \end{bmatrix}.$$

The stability of the uniform steady state to spatially uniform disturbance is determined by the eigenvalues of  $\mathcal{K}(u^*, v^*)$  and the stability of the uniform steady state to nonuniform disturbance is determined by the eigenvalues of  $\mathcal{K}(u^*, v^*) - 2\mathcal{D}$  [3].

We choose parameter values so that the uniform steady state is unstable, in which case a well-mixed system evolves to a unique periodic solution. This solution is also a solution of the coupled systems, since both cells are identical, and therefore a stochastic simulation should be expected to exhibit periodic behavior if averaged over many realizations. Using the stochastic and deterministic parameters given in Table 4, we compare the time evolution of  $X$  and  $Y$  in the two compartment model. Let  $N^{(k,1)}(t)$  and  $N^{(k,2)}(t)$  represent the numbers of molecules of  $X$  and  $Y$  in the  $k^{\text{th}}$  compartment at time  $t$ , respectively.

Note that in Table 4, we multiply  $k_0$  by the volume of each compartment ( $V_c$ ) to obtain the corresponding stochastic rate constant, since the corresponding reaction is production from a source occurring in both compartments. Similarly, we divide  $k_1$  by square of the volume of each compartment to obtain the stochastic rate constant, since the corresponding reaction,  $X + 2Y \rightarrow 3Y$ , is trimolecular. Diffusion coefficients in both the deterministic and the

1 stochastic two-compartment model are scaled by the square of each compartment size due to the discretization of  
 2 the Laplacian. For the parameter values chosen, the second diagonal element of  $\mathcal{K}(u^*, v^*)$ , corresponding to the  
 3 non-diffusing species  $Y$ , is positive, which reflects the self-activation of that species. Thus a deterministic spatially-  
 4 continuous system with these kinetics does not satisfy condition (i) in Theorem 3, and thus we cannot apply the  
 5 criterion to determine an appropriate discretization. Furthermore, we have

$$\begin{aligned} 6 \quad & Tr(\mathcal{K}(u^*, v^*)) > 0, \\ 7 \quad & det(\mathcal{K}(u^*, v^*)) > 0, \\ 8 \quad & Tr(\mathcal{K}(u^*, v^*) - 2\mathcal{D}) > 0, \\ 9 \quad & det(\mathcal{K}(u^*, v^*) - 2\mathcal{D}) > 0, \end{aligned}$$

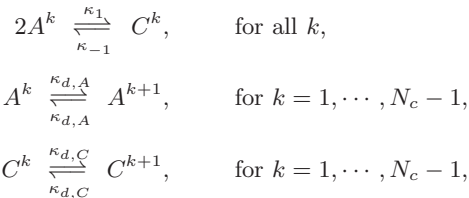
10 and therefore the eigenvalues of  $\mathcal{K}(u^*, v^*)$  and  $\mathcal{K}(u^*, v^*) - 2\mathcal{D}$  are positive and the uniform steady state is unstable  
 11 to both uniform and nonuniform disturbances.

12 In Figure 7 we compare the stochastic and deterministic simulations of the two-compartment model. In (a) and  
 13 (b), we show the uniform periodic solution of the deterministic two-compartment model for initial values given in  
 14 Table 4. One sees that when  $u_k(t)$  approaches zero,  $v_k(t)$  approaches to its maximum. In (c)–(f) and (g)–(j) we give  
 15 two realizations of stochastic simulations of the two-compartment model. Clearly there is no hint of periodicity in the  
 16 stochastic simulations, despite running for a long time compared to that needed for relaxation to the uniform periodic  
 17 solution in the deterministic case, and the compartments are certainly not synchronized. As in (a) and (b), when  $X$   
 18 peaks,  $Y$  in the same compartment has a local minimum value. However, the local minimum and maximum values  
 19 of the numbers of molecules for the same species vary, and the burst times of the same species in two compartments  
 20 are not synchronized. One reason for the discrepancy between the deterministic and stochastic solutions is that there  
 21 is also a stable non-uniform periodic solution in the deterministic system, and though the initial data lies in the  
 22 domain of attraction of the uniform solution for the deterministic problem, the non-uniform solution appears to exert  
 23 a significant influence on the stochastic evolution. A detailed analysis of the deterministic case is done on Ashkenazi  
 24 and Othmer [3], where the reader can see the full complexity of the solution set for this simple system.

25 The complexity of the solution set is solely the result of the self-activation in the local dynamics – without that the  
 26 system would not show oscillations and the condition (i) in Theorem 3 would be satisfied. Thus a full understanding  
 27 of the local dynamics is needed to determine whether a sufficiently small compartment size will produce meaningful  
 28 results in a stochastic simulation.

### 29 4.3. The reaction $2A \rightleftharpoons C$ in a distributed setting

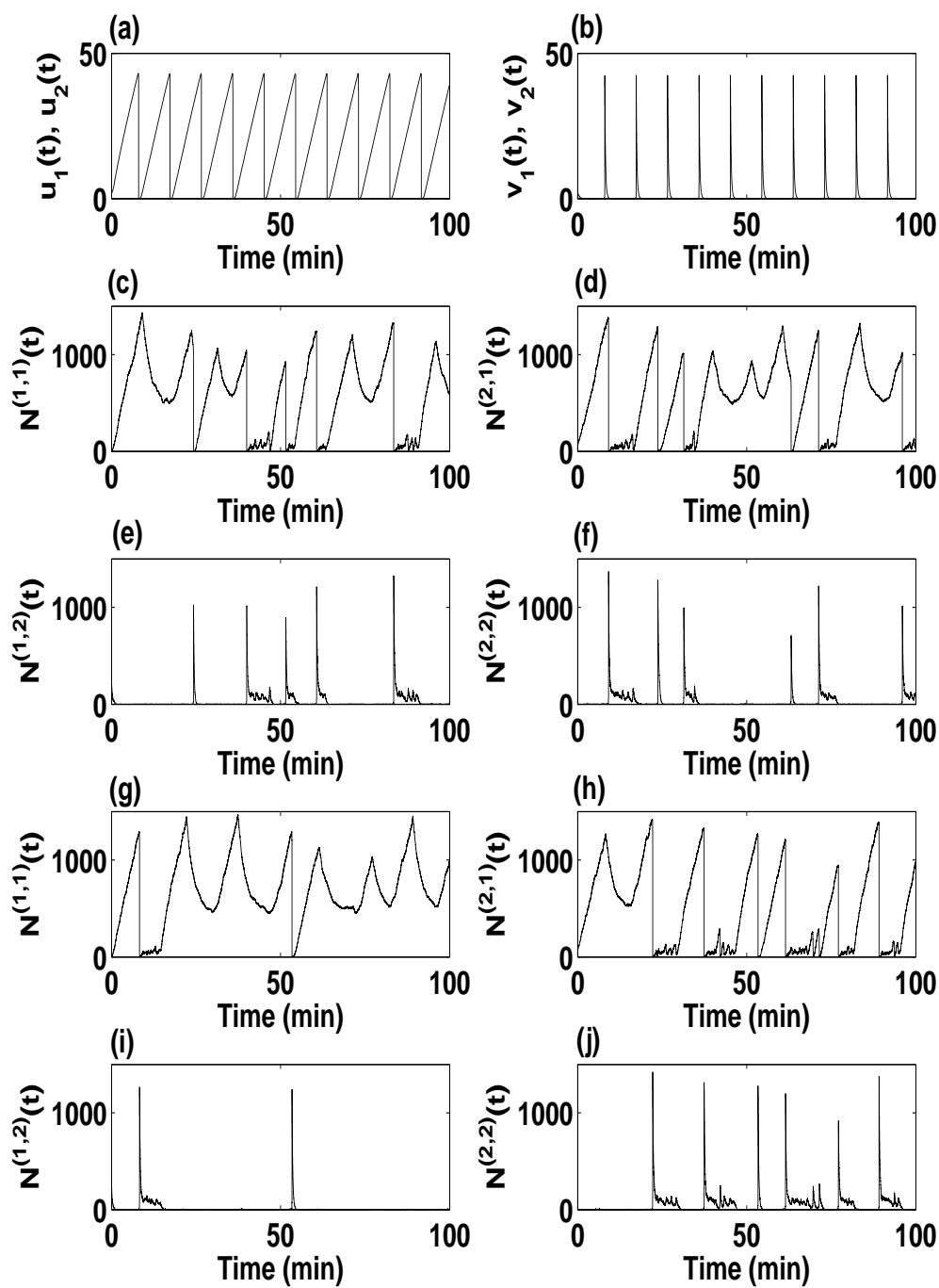
30 Consider a multi-compartment system of  $N_c$  compartments. Let  $A^k$  and  $C^k$  denote species  $A$  and  $C$  in the  $k^{th}$   
 31 compartment for  $k = 1, \dots, N_c$ . The system has the following reactions



35 wherein the parameter values are as given in Table 5.

36 In Table 5,  $\kappa_1$  is divided by the volume of the compartment ( $V_c$ ) to obtain the stochastic rate constant  $\kappa_1$ .  
 37 As before, we assume that  $L_x \gg L_y, L_z$ , and discretize  $L_x$  into  $N_c$  compartments. Since this system is closed, the  
 38 components are strongly connected, the deficiency is zero, and the steady-state probability density function has a  
 39 product form [1]. (Zero-deficiency of the network means that  $\nu$  does not annihilate any elements in the range of  $\mathcal{E}$   
 40 [33].)

41 Let  $N^{(k,1)}(t)$  and  $N^{(k,2)}(t)$  represent the numbers of molecules of species  $A$  and  $C$  in the  $k^{th}$  compartment at time  
 42  $t$ , respectively. Then,  $\sum_{k=1}^{N_c} (N^{(k,1)}(t) + 2N^{(k,2)}(t))$  is conserved and equal to its initial value  $N_0$ . The steady-state



**Fig. 7.** Deterministic and stochastic simulation for the two-compartment model.  $u_k(t)$  and  $v_k(t)$  are concentrations of species  $X$  and  $Y$  in the  $k^{\text{th}}$  compartment at time  $t$ .  $N^{(k,1)}(t)$  and  $N^{(k,2)}(t)$  are the numbers of molecules of species  $X$  and  $Y$  in the  $k^{\text{th}}$  compartment at time  $t$ .

**Table 5.** Deterministic and stochastic parameters in Example 4.3

Deterministic		Stochastic	
$k_1$	$0.05 \text{ nM}^{-1} \text{ min}^{-1}$	$\kappa_1$	$k_1/(\mathcal{N}_A V_c) \approx 1.21 \times 10^{-4} N_c \text{ min}^{-1}$
$k_{-1}$	$0.05 \text{ min}^{-1}$	$\kappa_{-1}$	$0.05 \text{ min}^{-1}$
$D_A = D_C$	$73 \mu\text{m}^2 \text{ sec}^{-1}$	$\kappa_{d,A} = \kappa_{d,C}$	$D_A/(L_x/N_c)^2 \approx 9.65 \times 10^{-4} N_c^2 \text{ sec}^{-1}$
	$0.73 \mu\text{m}^2 \text{ sec}^{-1}$		$D_A/(L_x/N_c)^2 \approx 9.65 \times 10^{-6} N_c^2 \text{ sec}^{-1}$
$L_x$	$275 \mu\text{m}$		
$L_y$	$5 \mu\text{m}$		
$L_z$	$0.5 \mu\text{m}$		
$\mathbf{V}$	$687.5 \mu\text{m}^3$		

1 probability density function can be expressed as [1]

$$2 \quad P_\infty(n^{(1,1)}, n^{(1,2)}, \dots, n^{(k,1)}, n^{(k,2)}, \dots, n^{(N_c,1)}, n^{(N_c,2)}) = M \prod_{k=1}^{N_c} \frac{(w_A^k)^{n^{(k,1)}}}{(n^{(k,1)})!} \frac{(w_C^k)^{n^{(k,2)}}}{(n^{(k,2)})!} \quad (74)$$

3 where  $w_A^k$  and  $w_C^k$  for  $k = 1, \dots, N_c$  are the components of the steady-state solution of the following deterministic  
4 system.

$$5 \quad \frac{dw_A^k(t)}{dt} = -2\kappa_1 w_A^k(t)^2 + 2\kappa_{-1} w_C^k(t) + \kappa_{d,A} \left[ (w_A^{k-1}(t) - w_A^k(t)) I_{\{k \neq 1\}} + (w_A^{k+1}(t) - w_A^k(t)) I_{\{k \neq N_c\}} \right] \quad (75)$$

$$6 \quad \frac{dw_C^k(t)}{dt} = \kappa_1 w_A^k(t)^2 - \kappa_{-1} w_C^k(t) + \kappa_{d,C} \left[ (w_C^{k-1}(t) - w_C^k(t)) I_{\{k \neq 1\}} + (w_C^{k+1}(t) - w_C^k(t)) I_{\{k \neq N_c\}} \right] \quad (76)$$

Here  $I$  is the indicator function,  $w_A^k$  and  $w_C^k$  satisfy the conservation law,  $\sum_{k=1}^{N_c} (w_A^k + 2w_C^k) = 1$ , and in (74),  $M$  is a  
normalizing constant that depends on  $N_c$ . The steady-state solution of (75) and (76) is spatially uniform and given  
by

$$w_A^k = Kd, \quad w_C^k = \frac{K}{2}d^2,$$

where

$$K = \frac{\kappa_{-1}}{2\kappa_1}, \quad d = \frac{-1 + \sqrt{1 + \frac{4}{KN_c}}}{2}.$$

7 Since  $d$  and  $K$  do not depend on the diffusion coefficients of species  $A$  and  $C$ , the steady-state probability density  
8 function does not depend on them neither. (Compare the earlier example of diffusion only in Section 1.4, which led  
9 to a multinomial distribution independent of the diffusion constant.) Using  $P_\infty(\cdot)$  in (74) and parameters given in  
10 Table 5, the computed  $CV^*$  and  $\overline{CV}^*$  for  $N_0 = 6$  is given in Table 6. For  $N_c \geq 2$ ,  $\overline{CV}^*$  is approximately constant.  
11 Thus the results suggest that the number of compartments should be at least 2 in a multi-compartment stochastic  
model.

**Table 6.**  $CV^*$  and  $\overline{CV}^*$  when  $N_0 = 6$ 

$N_c$	$CV^*$	$\overline{CV}^*$
1	3.7164	97.4456
2	5.3659	99.4871
3	6.5719	99.4871
4	7.5886	99.4871
5	8.4843	99.4871
6	9.2941	99.4871
7	10.0387	99.4871
8	10.7319	99.4871
9	11.3829	99.4871
10	11.9986	99.4871

1 Next, we apply Theorem 3 to estimate an upper bound for each compartment size in the stochastic model. Let  
 2  $u_1(\mathbf{x}, t)$  and  $u_2(\mathbf{x}, t)$  be concentrations of species  $A$  and  $C$  on  $\mathbf{x}$  at time  $t$ , and denote  $u(\mathbf{x}, t) \equiv [u_1(\mathbf{x}, t), u_2(\mathbf{x}, t)]^T$ .  
 3 The reaction and diffusion matrices are given as

$$4 \quad \mathcal{R}(u(\mathbf{x}, t)) = \begin{bmatrix} -2k_1 u_1(\mathbf{x}, t)^2 + 2k_{-1} u_2(\mathbf{x}, t) \\ k_1 u_1(\mathbf{x}, t)^2 - k_{-1} u_2(\mathbf{x}, t) \end{bmatrix},$$

$$5 \quad \mathcal{D} = \begin{bmatrix} D_A & 0 \\ 0 & D_C \end{bmatrix}.$$

6 Since we assume that both species  $A$  and  $C$  diffuse, (40) is reduced to

$$7 \quad |\alpha_1 \delta| > \hat{r}_u \tag{77}$$

8 where  $\alpha_1 = -(\pi/h_x)^2$  and  $\delta = D_A$ . The Jacobian matrix for the kinetic terms is

$$9 \quad D_u \mathcal{R}(u(\mathbf{x}, t)) = \begin{bmatrix} -4k_1 u_1(\mathbf{x}, t) & 2k_{-1} \\ 2k_1 u_1(\mathbf{x}, t) & -k_{-1} \end{bmatrix}.$$

10 Using  $\|D_u \mathcal{R}(u(\mathbf{x}, t))\|_E = \sqrt{20(k_1 u_1(\mathbf{x}, t))^2 + 5(k_{-1})^2}$ , we have

$$11 \quad \hat{r}_u = \max_{u_1(\mathbf{x}, t) \in C_\infty} \sqrt{20(k_1 u_1(\mathbf{x}, t))^2 + 5(k_{-1})^2}.$$

12 For fixed volume  $V_c$ ,  $\hat{r}_u$  has a maximum value when all molecules of species  $A$  are located in one compartment.  
 13 Therefore, we approximate the maximum value of  $u_1(\mathbf{x}, t)$  by  $N_0/(\mathcal{N}_A V_c) = N_0/(\mathcal{N}_A h_x L_y L_z)$  where  $N_0 = 6$  is the  
 14 conserved number of molecules. Using the upper bound of  $\hat{r}_u$ , we find the following is a sufficient condition to satisfy  
 15 (77):

$$16 \quad \frac{D_A \pi^2}{h_x^2} > \sqrt{20 \left( \frac{k_1 N_0}{\mathcal{N}_A h_x L_y L_z} \right)^2 + 5(k_{-1})^2}.$$

17 It follows that  $h_x$  must be chosen so that

$$18 \quad 5(k_{-1})^2 h_x^4 + 20 \left( \frac{k_1 N_0}{\mathcal{N}_A L_y L_z} \right)^2 h_x^2 - D_A^2 \pi^4 < 0,$$

19 which leads to the following condition on  $h_x$ .

$$20 \quad h_x < 621.79 \mu m \quad \text{when } D_A = D_C = 73 \mu m^2 \text{ sec}^{-1}$$

$$21 \quad h_x < 61.93 \mu m \quad \text{when } D_A = D_C = 0.73 \mu m^2 \text{ sec}^{-1}$$

22 Said otherwise, a single compartment suffices if  $L_x < 622 \mu m$ , which compares well with the result in Table  
 23 6, which depends on  $L_x$  through the rate constants, and are computed for  $L_x = 275 \mu m$ . However when  $D_A =$   
 24  $D_C = 0.73 \mu m^2 \text{ sec}^{-1}$ , the upper bound for the compartment size gives  $N_c > 4$  which overestimates the number of  
 25 compartments needed according to Table 6. This is not surprising, since the theoretical estimate is based on the  
 26 relaxation rate to the uniform solution, and for the second case, in which diffusion is 100-fold slower, the approach to  
 27 the uniform solution takes much longer and at least four compartments are needed to capture the temporal evolution.  
 28 Once again, care is needed in applying the criterion.

29 Finally, we compare our result to the one using Bernstein's criterion for the compartment size introduced in  
 30 Section 1.4. Consider the case when  $D_A = D_C = 73 \mu m^2 \text{ sec}^{-1}$ . To compute the reaction timescale, we use the  
 31 reaction with the largest propensity, since Bernstein sets the reaction time as the reciprocal of the propensity. Denote

1  $A_{tot}$  and  $C_{tot}$  as the total numbers of molecules of species  $A$  and  $C$ , respectively. Using  $A_{tot} + 2C_{tot} = 6$ , we compute

$$2 \quad \kappa_1 \frac{A_{tot}}{N_c} \left( \frac{A_{tot}}{N_c} - 1 \right) \approx (1.21 \times 10^{-4} N_c) \frac{6}{N_c} \left( \frac{6}{N_c} - 1 \right) \approx 0.0044 \left( \frac{1}{N_c} - \frac{1}{6} \right),$$

$$3 \quad \kappa_{-1} \frac{C_{tot}}{N_c} \approx 0.05 \frac{3}{N_c} = 0.15 \left( \frac{1}{N_c} \right).$$

4 Therefore, the propensity of the first-order reaction,  $C \xrightarrow{\kappa_1} 2A$ , is the largest. Based on the conservation law, we assume  
5 that the total number of molecules of species  $C$  is 3. The reaction time-scale is given as

$$6 \quad \tau_C \approx \frac{1}{0.05 \text{ min}^{-1} \times \frac{3}{N_c}}$$

$$7 \quad = \frac{1833.33}{h_x} \text{ min.}$$

8 The diffusion time-scale is computed using the diffusion coefficient of species  $A$ .

$$9 \quad \frac{h_x^2}{2D_A} \approx \frac{h_x^2 \mu\text{m}^2}{2 \times 4380 \mu\text{m}^2 \text{ min}^{-1}} = \frac{h_x^2}{8760} \text{ min}$$

10 Since the diffusion time-scale should be much faster than the reaction time-scale, we have

$$11 \quad h_x \ll (8760 \times 1833.33)^{1/3} \mu\text{m}$$

$$12 \quad \approx 252.30 \mu\text{m}.$$

13 Similarly, when  $D_A = D_C = 0.73 \mu\text{m}^2 \text{ sec}^{-1}$ , Bernstein's criterion yields  $h_x \ll 54.36 \mu\text{m}$ . Therefore, in both cases the  
14 upper bound for the compartment size using our criterion is larger than the one obtained from Bernstein's criterion.

## 15 5. Discussion

16 A number of criteria have been suggested for determining the appropriate compartment size for a stochastic treatment  
17 of a reaction-diffusion system, but none incorporates measures of the noise in the process, nor do they account for the  
18 properties of the integrated reaction network. We incorporate those factors here and develop a criterion for choosing  
19 the compartment size.

20 We first define a suitably scaled, generalized coefficient of variation as an appropriate measure for the noise level of  
21 the system, and use this scaled measure,  $\overline{CV}^*$ , to determine a lower bound for the number of compartments based on  
22 simulations of a particular network. Using results from [17], we compute the asymptotic mean  $M_\infty$  and variance  $V_\infty$ ,  
23 and we express  $\overline{CV}^*$  in terms of  $M_\infty$  for specific cases of a linear reaction-diffusion network. We show the convergence  
24 of each component in  $\overline{M}_\infty$  multiplied by some constant to the corresponding concentration in the continuum model  
25 for an open linear network with production from a source, and for a closed linear network with no inputs and no  
26 degradation (in case there is exactly one zero eigenvalue in the reaction matrix). We then show computationally that  
27 the scaled measure converges for a sufficiently large number of compartments in a linear network, which suggests the  
28 minimum value for the number of compartments.

29 However, it is not easy in general to compute the minimum value of the number of compartments using  $\overline{CV}^*$   
30 analytically, and therefore we developed an alternate criterion based on convergence of solutions of the deterministic  
31 reaction-diffusion system to a spatially-uniform solution. In previous work, conditions for this convergence were  
32 derived for closed systems in which all species diffuse [31,3], and we extend this here to allow both non-diffusing  
33 species and degradation of diffusing species on the boundary of the domain. Of course the system size plays a role  
34 in the convergence, and this leads to an estimate of the minimal compartment size by requiring that the dynamics  
35 converge to a spatially-uniform solution for that compartment size. The exponential convergence condition involved  
36 is applicable to the nonlinear reaction-diffusion networks for which it is known that solutions are bounded in  $L_2$ ,  
37 which include the majority of biologically-realistic systems. We apply the results to a simple dimerization reaction  
38 that illustrates some of the issues that must be considered in general.

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<sup>3</sup> paper.

## 1 APPENDICES

2 **A. Proof of the mean first passage time for  $N$  walkers**

3 Consider the first passage time for annihilation beginning on a spherical surface of radius  $r \in (a, b)$ . Define  $P_i(R, t|r, s)$   
 4 as the probability that the  $i^{\text{th}}$  particle starts on a spherical surface of radius  $r$  at time  $s$  and ends on a spherical  
 5 surface of radius  $R$  at time  $t$ . Define  $\tau_i(r) \equiv \tau_i$  as the first passage time for annihilation of the  $i^{\text{th}}$  particle when it  
 6 begins on the spherical surface of radius  $r$  and  $\tau(r_1, r_2, \dots, r_N) \equiv \tau$  as the first passage time for annihilation of the  
 7  $N$  particles, where for  $i = 1, 2, \dots, N$ , the  $i^{\text{th}}$  particle begins on a spherical surface of radius  $r_i$ . Define a cumulative  
 8 distribution function for  $\tau_i$  as

$$\begin{aligned} 9 \quad F_i(r, t) &\equiv P(\tau_i \leq t), \\ 10 \quad &= \int_0^t -\frac{d}{ds} \int_a^b P_i(R, s|r, 0) dR ds = 1 - \int_a^b P_i(R, t|r, 0) dR, \end{aligned}$$

11 and define the operator  $\Delta_r$  as

$$\begin{aligned} 12 \quad \Delta_r f(r, t) &= \frac{1}{r^2} \frac{\partial}{\partial r} \left( r^2 \frac{\partial f(r, t)}{\partial r} \right), \\ 13 \quad \Delta_r g(r) &= \frac{1}{r^2} \frac{d}{dr} \left( r^2 \frac{dg(r)}{dr} \right). \end{aligned}$$

14 Then, for  $i = 1, \dots, N$ ,  $F_i(r, t)$  is a solution of

$$\begin{aligned} &\frac{\partial}{\partial t} F_i(r, t) = D \Delta_r F_i(r, t), \\ 15 \quad &F_i(r, t)|_{\{r=a\}} = 1, \\ &\frac{\partial}{\partial r} F_i(r, t)|_{\{r=b\}} = 0. \end{aligned} \tag{78}$$

16 Let  $F(r_1, r_2, \dots, r_N, t)$  be the cumulative distribution function of  $\tau$  when  $N$  particles are initially located on spherical  
 17 surface of radius  $r_i \in (a, b)$  for  $i = 1, \dots, N$ . Then

$$\begin{aligned} 18 \quad P(\tau > t) &= 1 - F(r_1, r_2, \dots, r_N, t) \\ 19 \quad &= \prod_{i=1}^N (1 - F_i(r_i, t)). \end{aligned}$$

20 The mean first passage time of  $N$  particles is defined as  $\lambda \equiv E[\tau]$ , and is computed in terms of the cumulative  
 21 distribution function of the first passage time of each particle.

$$\begin{aligned} 22 \quad \lambda(r_1, r_2, \dots, r_N) &= \int_0^\infty P(\tau > s) ds \\ 23 \quad &= \int_0^\infty \prod_{i=1}^N (1 - F_i(r_i, s)) ds \end{aligned}$$

24 Using (78),  $\lambda$  satisfies

$$\begin{aligned} 25 \quad D \Delta_{r_i} \lambda(r_1, r_2, \dots, r_N) &= - \int_0^\infty (D \Delta_{r_i} F_i(r_i, s)) \prod_{j \neq i} (1 - F_j(r_j, s)) ds \\ 26 \quad &= - \int_0^\infty \frac{\partial F_i(r_i, s)}{\partial s} \prod_{j \neq i} (1 - F_j(r_j, s)) ds. \end{aligned} \tag{79}$$

27 Summing (79) over  $i = 1, \dots, N$ , we find that

$$28 \quad \sum_{i=1}^N D \Delta_{r_i} \lambda(r_1, r_2, \dots, r_N) = \int_0^\infty \frac{\partial \prod_{i=1}^N (1 - F_i(r_i, s))}{\partial s} ds$$



$$\begin{aligned}
&= \lim_{t \rightarrow \infty} \prod_{i=1}^N (1 - F_i(r_i, t)) - \prod_{i=1}^N (1 - F_i(r_i, 0)) \\
&= -1.
\end{aligned}$$

$\lambda$  also satisfies the boundary conditions

$$\begin{aligned}
&\lambda(r_1, r_2, \dots, r_N)|_{\{r_i=a\}} = 0, \\
&\frac{\partial}{\partial r_i} \lambda(r_1, r_2, \dots, r_N)|_{\{r_i=b\}} = 0,
\end{aligned}$$

for each  $i = 1, \dots, N$ . Substituting  $r_0 = a$ ,  $r_1 = b$ , and  $\tau = \tau_i$  in (7) in Section 1.4, for each  $i = 1, \dots, N$ ,  $E[\tau_i(r)]$  satisfies

$$\begin{aligned}
D\Delta_r E[\tau_i(r)] &= -1, \\
E[\tau_i(a)] &= 0, \\
E[\tau'_i(b)] &= 0.
\end{aligned}$$

Thus,  $\lambda$  is given as

$$\lambda(r_1, r_2, \dots, r_N) = \frac{1}{N} \sum_{i=1}^N E[\tau_i(r_i)] \prod_{j \neq i} (1 - \delta(r_j - a)).$$

## B. Spectral representation of the reaction-diffusion matrix $\Omega$

In this section, we give the explicit form of the eigenvalues and eigenvectors of the reaction-diffusion matrix  $\Omega$ , which are used in Section 2.1. Consider the first-order chemical reaction-diffusion network in the 3-dimensional parallelepiped with dimension  $L_x \times L_y \times L_z$ .  $\Delta_n$  is a finite difference approximation of the 1-dimensional Laplacian with homogeneous Neumann conditions. Then, modulo terms in  $h_x^{-2}$ , the entries of  $\Delta_n$  are

$$(\Delta_n)_{ij} = \begin{cases} -1 & \text{when } i = j = 1 \text{ or } i = j = n, \\ -2 & \text{when } i = j, i \neq 1, \text{ and } i \neq n, \\ 1 & \text{when } |i - j| = 1, \\ 0 & \text{otherwise.} \end{cases}$$

Let  $\mathcal{D}_x$ ,  $\mathcal{D}_y$ , and  $\mathcal{D}_z$  be diagonal matrices whose diagonal elements are the scaled diffusion coefficients for diffusion along the  $x$ ,  $y$ , and  $z$  directions, respectively. The reaction-diffusion matrix  $\Omega$  is expressed as

$$\Omega = \overline{\Delta}_z \otimes \mathcal{D}_z + \overline{\Delta}_y \otimes \mathcal{D}_y + \overline{\Delta}_x \otimes \mathcal{D}_x + (I_{N_z} \otimes I_{N_y} \otimes I_{N_x}) \otimes \mathcal{K}$$

where

$$\begin{aligned}
\overline{\Delta}_x &\equiv I_{N_z} \otimes I_{N_y} \otimes \Delta_{N_x}, \\
\overline{\Delta}_y &\equiv I_{N_z} \otimes \Delta_{N_y} \otimes I_{N_x}, \\
\overline{\Delta}_z &\equiv \Delta_{N_z} \otimes I_{N_y} \otimes I_{N_x}.
\end{aligned}$$

Define an index for compartments as  $l = (l_1, l_2, l_3)$ . Let  $\mathbf{1} = (l, \mu)$  for  $l_1, l_2, l_3 = 1, \dots, N_x, N_y, N_z$ , respectively, and for  $\mu = 1, \dots, s$ . Denote by  $\alpha_{l_1}$ ,  $\alpha_{l_2}$  and  $\alpha_{l_3}$  the eigenvalues of  $\Delta_{N_x}$ ,  $\Delta_{N_y}$ , and  $\Delta_{N_z}$  given as

$$\begin{aligned}
\alpha_{l_1} &= -4 \sin^2 \left( \frac{\pi l_1}{2N_x} \right) (1 - \delta_{l_1 N_x}), \\
\alpha_{l_2} &= -4 \sin^2 \left( \frac{\pi l_2}{2N_y} \right) (1 - \delta_{l_2 N_y}),
\end{aligned}$$

$$\alpha_{l_3} = -4 \sin^2 \left( \frac{\pi l_3}{2N_z} \right) (1 - \delta_{l_3 N_z}),$$

where  $\delta_{ij}$  is the Kronecker delta. Denote  $\phi_{l_1}$ ,  $\phi_{l_2}$ , and  $\phi_{l_3}$  as the corresponding eigenvectors of  $\Delta_{N_x}$ ,  $\Delta_{N_y}$ , and  $\Delta_{N_z}$  with elements given as

$$\begin{aligned} (\phi_{l_1})_{l'_1} &= \sqrt{\frac{2}{N_x}} \cos \left( \frac{\pi l_1 (l'_1 - 1/2)}{N_x} \right) (1 - \delta_{l_1 N_x}) + \sqrt{\frac{1}{N_x}} \delta_{l_1 N_x}, \\ (\phi_{l_2})_{l'_2} &= \sqrt{\frac{2}{N_y}} \cos \left( \frac{\pi l_2 (l'_2 - 1/2)}{N_y} \right) (1 - \delta_{l_2 N_y}) + \sqrt{\frac{1}{N_y}} \delta_{l_2 N_y}, \\ (\phi_{l_3})_{l'_3} &= \sqrt{\frac{2}{N_z}} \cos \left( \frac{\pi l_3 (l'_3 - 1/2)}{N_z} \right) (1 - \delta_{l_3 N_z}) + \sqrt{\frac{1}{N_z}} \delta_{l_3 N_z}, \end{aligned} \quad (80)$$

for  $l'_1, l'_2, l'_3 = 1, \dots, N_x, N_y, N_z$ , respectively. Then, the eigenvalue of  $\bar{\Delta}_x + \bar{\Delta}_y + \bar{\Delta}_z$  is expressed as  $\alpha_l \equiv \alpha_{l_1} + \alpha_{l_2} + \alpha_{l_3}$  and the corresponding eigenvector and adjoint eigenvector are given as  $\phi_l \equiv \phi_{l_1} \otimes \phi_{l_2} \otimes \phi_{l_3}$  and  $\phi_l^*$ .

Define the complete diffusion matrix as  $\alpha_l \mathcal{D} \equiv \alpha_{l_1} \mathcal{D}_x + \alpha_{l_2} \mathcal{D}_y + \alpha_{l_3} \mathcal{D}_z$ , and then the eigenvalue  $\lambda_1$  of  $\Omega$  is a solution of

$$|\mathcal{K} + \alpha_l \mathcal{D} - \lambda_1 I_{n_s}| = 0.$$

Let  $\varphi_1$  and  $\varphi_1^*$  be the eigenvector and the adjoint eigenvector of  $\mathcal{K} + \alpha_l \mathcal{D}$  corresponding to the eigenvalue  $\lambda_1$ . Then,  $\phi_l \otimes \varphi_1$  is an eigenvector of  $\Omega$  corresponding to  $\lambda_1$ , and the projection  $P_1$  associated with  $\lambda_1$  is

$$P_1 = (\phi_l * \bar{\phi}_l^*) \otimes (\varphi_1 * \bar{\varphi}_1^*) \quad (81)$$

where  $*$  is the dyad product and  $\bar{\phi}_l^*$  is the complex conjugate of  $\phi_l^*$ .

In Example 2.2,  $N_y = N_z = 1$  and we set  $N_x = N_c$ . The eigenvalues of  $\Delta$  are

$$\alpha_l = \begin{cases} -4 \sin^2 \left( \frac{\pi l}{2N_c} \right), & l \neq N_c, \\ 0, & l = N_c, \end{cases} \quad (82)$$

and the corresponding eigenvectors are

$$\phi_l = \begin{cases} \sqrt{\frac{2}{N_c}} \left( \cos \left( \frac{\pi l}{2N_c} \right), \dots, \cos \left( \frac{(2l'-1)\pi l}{2N_c} \right), \dots, \cos \left( \frac{(2N_c-1)\pi l}{2N_c} \right) \right)^T, & l \neq N_c, \\ \sqrt{\frac{1}{N_c}} (1, 1, \dots, 1)^T, & l = N_c. \end{cases} \quad (83)$$

In Example 2.2,

$$\mathcal{K} + \alpha_l \mathcal{D} = -\frac{\kappa_5 \kappa_7 R}{\kappa_{-5} + \kappa_7} - \kappa_6 + \alpha_l \frac{D}{(L_x/N_c)^2},$$

and the eigenvalue of  $\mathcal{K} + \alpha_l \mathcal{D}$  is

$$\lambda_1 = -\frac{\kappa_5 \kappa_7 R}{\kappa_{-5} + \kappa_7} - \kappa_6 + \alpha_l \frac{D}{(L_x/N_c)^2}.$$

The corresponding eigenvector and adjoint eigenvector of  $\mathcal{K} + \alpha_l \mathcal{D}$  are  $\varphi_1 = 1, \varphi_1^* = 1$ . In Example 2.2, we only need the first column of  $P_1$  to compute the first and the second moments, since we assume that production from a source only occurs in the first compartment. Thus we have

$$P_1 \mathbf{k}^s = \kappa_8 P_1^{(1)}$$

where  $P_1^{(1)}$  represents the first column of  $P_1$ . From (81), we obtain the first column of the projection as

$$P_1^{(1)} = (\phi_l * \bar{\phi}_l^*)^{(1)} \otimes (\varphi_1 * \bar{\varphi}_1^*)^{(1)}$$

$$= (\phi_l * \overline{\phi_l^*})^{(1)}.$$

Using  $\phi_l = \overline{\phi_l^*}$ , we compute

$$P_1^{(1)} = \begin{cases} \frac{2}{N_c} \cos\left(\frac{\pi l}{2N_c}\right) \left( \cos\left(\frac{\pi l}{2N_c}\right), \dots, \cos\left(\frac{(2l'-1)\pi l}{2N_c}\right), \dots, \cos\left(\frac{(2N_c-1)\pi l}{2N_c}\right) \right)^T, & l \neq N_c, \\ \frac{1}{N_c} (1, 1, \dots, 1)^T, & l = N_c. \end{cases}$$

### C. Proof of Proposition 1

In this section, we prove Proposition 1 in Section 2.1. Using (10) in Section 2.1, we have

$$\begin{aligned} \frac{d(M(t)M(t)^T)}{dt} &= \frac{dM(t)}{dt}M(t)^T + M(t)\frac{dM(t)^T}{dt} \\ &= (\Omega M(t) + \mathbf{k}^s)M(t)^T + M(t)(\Omega M(t) + \mathbf{k}^s)^T \\ &= \Omega M(t)M(t)^T + [\Omega M(t)M(t)^T]^T + C(t) + C(t)^T - W(t) - W(t)^T. \end{aligned} \quad (84)$$

where  $C(t) = W(t) + \mathbf{k}^s M(t)^T$  and  $W(t)$  is a block-diagonal matrix with elements defined as

$$[W(t)]_{i(\eta), i(\zeta)} = \begin{cases} K_{ij}^{cat} [M(t)]_{i(\zeta)}, & \text{if } k = q, \\ 0, & \text{otherwise.} \end{cases}$$

Recall that  $M_d(t) = \text{diag}[M(t)]$ . Using (84), (10), and

$$\text{Cov}(t) = V(t) - M(t)M(t)^T + M_d(t),$$

we have

$$\begin{aligned} \frac{d\text{Cov}(t)}{dt} &= \frac{dV(t)}{dt} - \frac{d(M(t)M(t)^T)}{dt} + \frac{dM_d(t)}{dt} \\ &= \left\{ \Omega V(t) + [\Omega V(t)]^T + C(t) + C(t)^T \right\} \\ &\quad - \left\{ \Omega M(t)M(t)^T + [\Omega M(t)M(t)^T]^T + C(t) + C(t)^T - W(t) - W(t)^T \right\} \\ &\quad + \frac{dM_d(t)}{dt} \\ &= \Omega \text{Cov}(t) + [\Omega \text{Cov}(t)]^T - \Omega M_d(t) - (\Omega M_d(t))^T + W(t) + W(t)^T \\ &\quad + \frac{dM_d(t)}{dt}. \end{aligned} \quad (85)$$

From this we obtain

$$\frac{d(\text{Cov}(t) - M_d(t))}{dt} = \Omega(\text{Cov}(t) - M_d(t)) + [\Omega(\text{Cov}(t) - M_d(t))]^T + W(t) + W(t)^T. \quad (86)$$

Define  $\text{col}(A)$  as a vector by concatenating all columns of  $A$  in order, *i.e.*  $\text{col}(A) = [(A^{(1)})^T, (A^{(2)})^T, \dots, (A^{(m)})^T]^T$  where  $A^{(l)}$  is the  $l^{\text{th}}$  column of  $A$ . Define  $v(t) = \text{col}(\text{Cov}(t) - M_d(t))$ ,  $\mathcal{V} = \Omega \otimes I_{sN_c} + I_{sN_c} \otimes \Omega$ , and  $\gamma(t) = \text{col}(W(t) + W(t)^T)$ . We change matrices in (86) to column vectors, and obtain [17],

$$\frac{dv(t)}{dt} = \mathcal{V}v(t) + \gamma(t). \quad (87)$$

The solution of (87) is

$$v(t) = e^{\mathcal{V}t}v(0) + \int_0^t e^{\mathcal{V}(t-\tau)}\gamma(\tau) d\tau$$

$$\begin{aligned}
&= \sum_{\mathbf{l}, \mathbf{m}} e^{(\lambda_1 + \lambda_{\mathbf{m}})t} P_{\mathbf{l}} \otimes P_{\mathbf{m}} v(0) \\
&+ \sum_{\mathbf{l}, \mathbf{m}} \int_0^t e^{(\lambda_1 + \lambda_{\mathbf{m}})(t-\tau)} \left( P_{\mathbf{l}} K^{cat} \otimes P_{\mathbf{m}} + P_{\mathbf{l}} \otimes P_{\mathbf{m}} K^{cat} \right) col(M_d(\tau)) d\tau.
\end{aligned} \tag{88}$$

Reverting to the matrix form in (88), we find that

$$\begin{aligned}
(Cov(t) - M_d(t)) &= \sum_{\mathbf{l}, \mathbf{m}} e^{(\lambda_1 + \lambda_{\mathbf{m}})t} P_{\mathbf{m}} (Cov(0) - M_d(0)) P_{\mathbf{l}}^T \\
&+ \sum_{\mathbf{l}, \mathbf{m}} \int_0^t e^{(\lambda_1 + \lambda_{\mathbf{m}})(t-\tau)} \left( P_{\mathbf{m}} M_d(\tau) (P_{\mathbf{l}} K^{cat})^T + P_{\mathbf{m}} K^{cat} M_d(\tau) P_{\mathbf{l}}^T \right) d\tau.
\end{aligned} \tag{89}$$

Next we compute  $M_d(t)$ . Following [17], the evolution of the mean matrix is expressed as

$$M(t) = \sum_{\mathbf{n}} e^{\lambda_{\mathbf{n}} t} P_{\mathbf{n}} M(0) - \sum_{\mathbf{n}} \frac{1 - e^{\lambda_{\mathbf{n}} t}}{\lambda_{\mathbf{n}}} P_{\mathbf{n}} \mathbf{k}^s. \tag{90}$$

Define  $L(t)$  and  $S$  as  $sN_c \times sN_c$  matrices satisfying

$$\begin{aligned}
L(t) &= \left[ M(t) | M(t) | \cdots | M(t) \right]^T, \\
S &= \left[ \mathbf{k}^s | \mathbf{k}^s | \cdots | \mathbf{k}^s \right].
\end{aligned}$$

Define  $diag[b_1, b_2, \dots, b_n]$  as an  $n \times n$  diagonal matrix with the  $i^{th}$  diagonal element equal to  $b_i$ . Define diagonalization of matrices and of vectors as

$$\begin{aligned}
diag[A] &\equiv diag[a_{11}, a_{22}, \dots, a_{nn}], \\
diag[X] &\equiv diag[x_1, x_2, \dots, x_n],
\end{aligned}$$

where  $A$  is an  $n \times n$  matrix with each element  $a_{ij}$  and  $X$  is an  $n$ -dimensional vector with each element  $x_i$ . Diagonalizing both sides in (90), we have

$$M_d(t) = \sum_{\mathbf{n}} e^{\lambda_{\mathbf{n}} t} diag[P_{\mathbf{n}} L(0)^T] - \sum_{\mathbf{n}} \frac{1 - e^{\lambda_{\mathbf{n}} t}}{\lambda_{\mathbf{n}}} diag[P_{\mathbf{n}} S]. \tag{91}$$

In (89), we first calculate  $\int_0^t e^{(\lambda_1 + \lambda_{\mathbf{m}})(t-\tau)} P_{\mathbf{m}} M_d(\tau) (P_{\mathbf{l}} K^{cat})^T d\tau$  using (91), and find that

$$\begin{aligned}
&\int_0^t e^{(\lambda_1 + \lambda_{\mathbf{m}})(t-\tau)} P_{\mathbf{m}} M_d(\tau) (P_{\mathbf{l}} K^{cat})^T d\tau \\
&= \int_0^t e^{(\lambda_1 + \lambda_{\mathbf{m}})(t-\tau)} P_{\mathbf{m}} \left\{ \sum_{\mathbf{n}} e^{\lambda_{\mathbf{n}} \tau} diag[P_{\mathbf{n}} L(0)^T] - \sum_{\mathbf{n}} \frac{1 - e^{\lambda_{\mathbf{n}} \tau}}{\lambda_{\mathbf{n}}} diag[P_{\mathbf{n}} S] \right\} (P_{\mathbf{l}} K^{cat})^T d\tau \\
&= \sum_{\lambda_{\mathbf{n}} \neq \lambda_1 + \lambda_{\mathbf{m}}} \frac{e^{\lambda_{\mathbf{n}} t} - e^{(\lambda_1 + \lambda_{\mathbf{m}})t}}{-\lambda_{\mathbf{n}} + \lambda_1 + \lambda_{\mathbf{m}}} P_{\mathbf{m}} diag[P_{\mathbf{n}} L(0)^T] (P_{\mathbf{l}} K^{cat})^T \\
&+ \sum_{\lambda_{\mathbf{n}} = \lambda_1 + \lambda_{\mathbf{m}}} (t e^{\lambda_{\mathbf{n}} t}) P_{\mathbf{m}} diag[P_{\mathbf{n}} L(0)^T] (P_{\mathbf{l}} K^{cat})^T \\
&- \sum_{\lambda_{\mathbf{n}} \neq \lambda_1 + \lambda_{\mathbf{m}}} \frac{1}{\lambda_{\mathbf{n}}} \left[ \frac{1 - e^{(\lambda_1 + \lambda_{\mathbf{m}})t}}{-\lambda_{\mathbf{n}} + \lambda_1 + \lambda_{\mathbf{m}}} - \frac{e^{\lambda_{\mathbf{n}} t} - e^{(\lambda_1 + \lambda_{\mathbf{m}})t}}{-\lambda_{\mathbf{n}} + \lambda_1 + \lambda_{\mathbf{m}}} \right] P_{\mathbf{m}} diag[P_{\mathbf{n}} S] (P_{\mathbf{l}} K^{cat})^T \\
&- \sum_{\lambda_{\mathbf{n}} = \lambda_1 + \lambda_{\mathbf{m}}} \frac{1}{\lambda_{\mathbf{n}}} \left[ \frac{e^{\lambda_{\mathbf{n}} t} - 1}{\lambda_{\mathbf{n}}} - t e^{\lambda_{\mathbf{n}} t} \right] P_{\mathbf{m}} diag[P_{\mathbf{n}} S] (P_{\mathbf{l}} K^{cat})^T,
\end{aligned} \tag{92}$$

1 and  $\int_0^t e^{(\lambda_1+\lambda_m)(t-\tau)} P_m K^{cat} M_d(\tau) P_1^T d\tau$  as

$$\begin{aligned}
& \int_0^t e^{(\lambda_1+\lambda_m)(t-\tau)} P_m K^{cat} M_d(\tau) P_1^T d\tau \\
&= \sum_{\lambda_n \neq \lambda_1 + \lambda_m} \frac{e^{\lambda_n t} - e^{(\lambda_1+\lambda_m)t}}{-(\lambda_1 + \lambda_m) + \lambda_n} P_m K^{cat} \text{diag}[P_n L(0)^T] P_1^T \\
&+ \sum_{\lambda_n = \lambda_1 + \lambda_m} (te^{\lambda_n t}) P_m K^{cat} \text{diag}[P_n L(0)^T] P_1^T \\
&- \sum_{\lambda_n \neq \lambda_1 + \lambda_m} \frac{1}{\lambda_n} \left[ \frac{1 - e^{(\lambda_1+\lambda_m)t}}{-(\lambda_1 + \lambda_m)} - \frac{e^{\lambda_n t} - e^{(\lambda_1+\lambda_m)t}}{-(\lambda_1 + \lambda_m) + \lambda_n} \right] P_m K^{cat} \text{diag}[P_n S] P_1^T \\
&- \sum_{\lambda_n = \lambda_1 + \lambda_m} \frac{1}{\lambda_n} \left[ \frac{e^{\lambda_n t} - 1}{\lambda_n} - te^{\lambda_n t} \right] P_m K^{cat} \text{diag}[P_n S] P_1^T.
\end{aligned} \tag{93}$$

7 Now we prove the second and the third cases in Proposition 1, using the results of the previous computations.  
8 First, consider the open system with  $\sigma(\Omega) \subset LHP$ . Using (89), (92), and (93) and letting  $t \rightarrow \infty$ , we get

$$\begin{aligned}
Cov_\infty - M_{d,\infty} &= \sum_{l,m} \sum_{\lambda_n \neq \lambda_1 + \lambda_m} \frac{P_m \left\{ \text{diag}[P_n S] (K^{cat})^T + K^{cat} \text{diag}[P_n S] \right\} P_1^T}{\lambda_n (\lambda_1 + \lambda_m)} \\
&+ \sum_{l,m} \sum_{\lambda_n = \lambda_1 + \lambda_m} \frac{P_m \left\{ \text{diag}[P_n S] (K^{cat})^T + K^{cat} \text{diag}[P_n S] \right\} P_1^T}{\lambda_n^2} \\
&= \sum_{l,m,n} \frac{P_m \left\{ \text{diag}[P_n S] (K^{cat})^T + K^{cat} \text{diag}[P_n S] \right\} P_1^T}{\lambda_n (\lambda_1 + \lambda_m)}.
\end{aligned} \tag{94}$$

12 From (94), either if there is no production from a source or if there is no catalytic inputs, which is  $K^S = 0$  ( $S = 0$ )  
13 or  $K^{cat} = 0$ , we have  $Cov_\infty = M_{d,\infty}$ .

14 Second, consider a closed system for which  $\sigma(\Omega)$  is in the closed left-half plane. We only consider the case in  
15 which there is exactly one zero eigenvalue of  $\Omega$  and no inputs. Then (89) becomes

$$(Cov(t) - M_d(t)) = \sum_{l,m} e^{(\lambda_1+\lambda_m)t} P_m (Cov(0) - M_d(0)) P_1^T. \tag{95}$$

17 Letting  $t \rightarrow \infty$  in (95) and using the fact that there is exactly one zero eigenvalue of  $\Omega$  and that the remaining  
18 eigenvalues are negative, we get

$$Cov_\infty - M_{d,\infty} = P_s (Cov(0) - M_d(0)) P_s^T$$

20 where  $P_s$  is the projection corresponding to the zero eigenvalue. Assuming that initial values are deterministic,  
21  $Cov(0) = 0$ . As a result,  $Cov_\infty$  can be expressed in terms of  $M_{d,\infty}$ ,  $M_d(0)$ , and  $P_s$  as follows.

$$Cov_\infty = M_{d,\infty} - P_s M_d(0) P_s^T$$

## 23 D. Proof of Theorem 1

24 To prove the theorem, we first introduce and prove a lemma. The proof of the theorem is given at the end of this  
25 appendix. In the following lemma we show that as  $N_c \rightarrow \infty$ , each component of  $X_\infty / (\mathcal{N}_A V_c)$  converges to the  
26 corresponding value of the steady-state concentration in the continuum deterministic reaction-diffusion system.

27 Let  $u_\infty(\mathbf{x})$  and  $v_\infty(\mathbf{x})$  be the steady-state solution of the corresponding continuum model.  $u_\infty(\mathbf{x})$  and  $v_\infty(\mathbf{x})$   
28 represent concentration vectors for diffusing and non-diffusing species, respectively, which satisfy

$$\begin{aligned}
\tilde{\mathcal{D}} \Delta u_\infty(\mathbf{x}) + \mathcal{R} u_\infty(\mathbf{x}) + \mathcal{S} v_\infty(\mathbf{x}) + \delta(\mathbf{x}, 0) \tilde{k}^s &= 0, \\
\mathcal{T} u_\infty(\mathbf{x}) + \mathcal{W} v_\infty(\mathbf{x}) &= 0.
\end{aligned} \tag{96}$$

1 In (96), each component of  $\tilde{\mathbf{k}}^s$  is defined as

$$2 \quad \tilde{k}_i^s \equiv \mathbf{k}_{i(1,i)}^s \frac{L_x}{\mathcal{N}_A \mathbf{V}},$$

3 where  $\mathbf{V}$  is the volume of the system. Letting  $[\mathbf{k}^s]^{(1)} \equiv [\mathbf{k}_{i(1,i)}^s, \dots, \mathbf{k}_{i(1,m)}^s]^T$ ,  $\tilde{\mathbf{k}}^s$  is written as

$$4 \quad \tilde{\mathbf{k}}^s = \frac{L_x}{\mathcal{N}_A \mathbf{V}} [\mathbf{k}^s]^{(1)}. \quad (97)$$

5 In (96),  $\tilde{\mathcal{D}}$  is a diagonal matrix with diagonal elements the diffusion coefficients in the continuum description, while  
6 diagonal elements of  $\mathcal{D}$  in (18) are the corresponding diffusion coefficients in the discretized system. The two are  
7 related by

$$8 \quad \mathcal{D} = \frac{\tilde{\mathcal{D}}}{(L_x/N_c)^2}.$$

9 **Lemma 1.** Let  $X_\infty$  be the solution of (18) and  $u_\infty(\mathbf{x})$  be the solution of (96). Define  $U_\infty(\mathbf{x})$  as a vector with each  
10 element satisfying

$$11 \quad (U_\infty(\mathbf{x}))_i \equiv \sum_{k=1}^{N_c} \frac{1}{\mathcal{N}_A V_c} [X_\infty]_{i(k,i)} I_{\left\{ \left[ \frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c} \right] \right\}}(\mathbf{x}), \quad i = 1, \dots, m,$$

12 where  $m$  is the number of diffusing species. Then,  $\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_{L_2} = O\left(\frac{\ln N_c}{N_c}\right)$  and converges to 0 as  $N_c \rightarrow \infty$ .

13 Note that for  $\mathbf{x} \in \left[\frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c}\right)$ ,  $(U_\infty(\mathbf{x}))_i$  gives the concentration of the  $i^{\text{th}}$  species in the  $k^{\text{th}}$  compartment.

14 *Proof.* The proof of the Lemma 1 is lengthy, and we split it into 6 steps:

- 15 – **Step 1** Express  $U_\infty(\mathbf{x})$  and  $u_\infty(\mathbf{x})$  in terms of the discrete and continuous Green's functions;
- 16 – **Step 2** Split the error between  $U_\infty(\mathbf{x})$  and  $u_\infty(\mathbf{x})$  into three parts,  $(I)$ ,  $(II)$ , and  $(III)$  and split  $(I)$  into four  
17 parts,  $(i)$ ,  $(ii)$ ,  $(iii)$ , and  $(iv)$ ;
- 18 – **Step 3** To get an upper bound for  $(ii)$ , we prove  $\|Q_1\|_E \leq O(1)$ ;
- 19 – **Step 4** Prove  $\sum_\mu \left| \tilde{\lambda}_1^{-1} \right| \leq O\left(\frac{1}{N_c}\right)$  and finish showing  $\|(ii)\|_E, \|(iii)\|_E \leq O\left(\frac{\ln N_c}{N_c}\right)$ ;
- 20 – **Step 5** Show  $\|(i)\|_E, \|(iv)\|_E \leq o\left(\frac{1}{N_c}\right)$ ;
- 21 – **Step 6** Show  $\|(II)\|_E, \|(III)\|_E \leq O\left(\frac{1}{N_c}\right)$ .

22 **Step 1:**

23 First, we show that  $U_\infty(\mathbf{x})$ , which is defined in terms of  $X_\infty$ , is expressed in the form with a discrete version of  
24 Green's function.  $X_\infty$  is a solution of (18) and is given as

$$25 \quad X_\infty = \sum_1 \frac{1}{\lambda_1} (\phi_l * \overline{\phi_l^*}) \otimes (\varphi_1 * \overline{\varphi_1^*}) \mathbf{k}^s.$$

26 Recall that  $\mathbf{l} = (l, \mu)$  where  $l = (l_1, l_2, l_3)$  is an index for the compartment and  $\mu$  is an index for the species. Also,  
27 recall from Section 2.1 that  $\alpha_l$  and  $\phi_l$  are the eigenvalue and eigenvector of  $\Delta$  and  $\lambda_1$  and  $\varphi_1$  are the eigenvalue and  
28 eigenvector of  $\mathcal{K} + \alpha_l \mathcal{D}$ .  $\phi_l^*$  and  $\varphi_1^*$  are the corresponding adjoint eigenvectors. Denote

$$29 \quad Q_1 \equiv \varphi_1 * \overline{\varphi_1^*}$$

30 and define a spatial variable  $q_l(\mathbf{x})$  as

$$31 \quad q_l(\mathbf{x}) \equiv \sqrt{\frac{N_c}{L_x}} \sum_{l'=1}^{N_c} (\phi_l)_{l'} I_{\left\{ \left[ \frac{(l'-1)L_x}{N_c}, \frac{l'L_x}{N_c} \right] \right\}}(\mathbf{x}), \quad l = 1, 2, \dots, N_c. \quad (98)$$

1 Then, define each element of a spatial vector for the production rates from a source as

$$2 \quad k_i^s(\mathbf{x}) \equiv \sum_{k=1}^{N_c} \frac{\mathbf{k}_i^s(k,i)}{\mathcal{N}_A V_c} I_{\left\{ \left[ \frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c} \right] \right\}}(\mathbf{x}), \quad i = 1, \dots, m,$$

$$3 \quad = \frac{\mathbf{k}_i^s(1,i)}{\mathcal{N}_A V_c} I_{\left\{ \left[ 0, \frac{L_x}{N_c} \right] \right\}}(\mathbf{x}).$$

4 Using  $[\mathbf{k}^s]^{(1)}$ ,  $k^s(\mathbf{x})$  is written as

$$5 \quad k^s(\mathbf{x}) = \frac{1}{\mathcal{N}_A V_c} [\mathbf{k}^s]^{(1)} I_{\left\{ \left[ 0, \frac{L_x}{N_c} \right] \right\}}(\mathbf{x}). \quad (99)$$

6 Using the fact that  $\phi_l = \overline{\phi_l^*}$ , define a matrix for a discrete Green's function as

$$7 \quad G(\mathbf{x}, \boldsymbol{\xi}) \equiv \sum_1 \frac{1}{\lambda_1} q_l(\mathbf{x}) q_l(\boldsymbol{\xi}) Q_1. \quad (100)$$

8 Then, the vector for the scaled species numbers in different locations is expressed as

$$9 \quad U_\infty(\mathbf{x}) = \int_0^{L_x} G(\mathbf{x}, \boldsymbol{\xi}) k^s(\boldsymbol{\xi}) d\boldsymbol{\xi}. \quad (101)$$

10 Next, we show that  $u_\infty(\mathbf{x})$ , which corresponds to  $U_\infty(\mathbf{x})$  in the deterministic reaction-diffusion equations, can also  
 11 be expressed in terms of a Green's function. Using our assumption that  $\sigma(\mathcal{W}) \subset LHP$ , the steady-state concentration  
 12 of non-diffusing species is expressed in terms of the steady-state concentration of diffusing species and (96) is reduced  
 13 to

$$14 \quad \tilde{\mathcal{D}} \Delta u_\infty(\mathbf{x}) + \mathcal{K} u_\infty(\mathbf{x}) + \delta(\mathbf{x}, 0) \tilde{k}^s = 0, \quad \mathbf{x} \in [0, L_x], \quad (102)$$

15 where  $\mathcal{K} = \mathcal{R} - S\mathcal{W}^{-1}\mathcal{T}$ . Consider an eigenvalue problem related to (102).

$$16 \quad \begin{aligned} \tilde{\mathcal{D}} \Delta \mathcal{Y}(\mathbf{x}) + (\mathcal{K} - \lambda I_m) \mathcal{Y}(\mathbf{x}) &= 0, & \mathbf{x} \in [0, L_x], \\ \mathcal{Y}'(\mathbf{x}) &= 0, & \mathbf{x} = 0, L_x \end{aligned} \quad (103)$$

17 Let  $\tilde{\lambda}_1$  and  $\tilde{\varphi}_1$  be a solution of the algebraic eigenvalue problem

$$18 \quad (\mathcal{K} + \tilde{\alpha}_1 \tilde{\mathcal{D}} - \tilde{\lambda}_1 I_m) \tilde{\varphi}_1 = 0,$$

19 and let  $\tilde{\varphi}_1^*$  be the solution in the adjoint algebraic eigenvalue problem. We find that the solution of the scalar problem  
 20 in (19) is

$$21 \quad \tilde{\alpha}_l = - \left( \frac{l\pi}{L_x} \right)^2, \quad l = 0, 1, \dots, \quad (104)$$

22 and

$$23 \quad \tilde{q}_l(\mathbf{x}) = \begin{cases} \sqrt{\frac{1}{L_x}}, & l = 0, \\ \sqrt{\frac{2}{L_x}} \cos\left(\frac{l\pi\mathbf{x}}{L_x}\right), & l \neq 0. \end{cases} \quad (105)$$

24 Then, the eigenfunction of (103) is written as  $\mathcal{Y}_1(\mathbf{x}) \equiv \tilde{q}_l(\mathbf{x}) \tilde{\varphi}_1$ . Using our assumption that  $\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}$  is semi-simple,  
 25 the eigenfunctions are complete and the solution of (103) is given as

$$26 \quad \mathcal{Y}(\mathbf{x}) = \sum_1 \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) \tilde{\varphi}_1.$$

1 Define

$$2 \quad \tilde{Q}_1 \equiv \tilde{\varphi}_1 * \overline{\tilde{\varphi}_1^*}.$$

3 Since we assume that  $\sigma(\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}) \subset LHP$ , we have  $\tilde{\lambda}_1 \neq 0$  for all  $\mathbf{l}$  and the Green's function of the operator  $\tilde{\mathcal{D}}\Delta + \mathcal{K}$   
4 is given as

$$5 \quad \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \equiv \sum_{\mathbf{l}} \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \tilde{Q}_1.$$

6 Then, the steady-state concentration vector which is a solution of (102) is written as

$$7 \quad u_\infty(\mathbf{x}) = \int_0^{L_x} \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \delta(\boldsymbol{\xi}, 0) \tilde{k}^s d\boldsymbol{\xi}. \quad (106)$$

8 **Step 2:**

9 Now, we estimate the error between  $U_\infty(\mathbf{x})$  and  $u_\infty(\mathbf{x})$  and show that  $\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_{L_2}$  is  $O\left(\frac{\ln N_c}{N_c}\right)$ . Define a  
10 projection of the Green's function onto the space with a finite number of frequencies and its remainder as

$$11 \quad \pi_{N_c} \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \equiv \sum_{l < N_c, 1} \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \tilde{Q}_1, \quad (107)$$

$$12 \quad (I_m - \pi_{N_c}) \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \equiv \sum_{l \geq N_c, 1} \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \tilde{Q}_1. \quad (108)$$

13 Subtracting (106) from (101) and breaking  $U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})$  into three parts, we get

$$14 \quad U_\infty(\mathbf{x}) - u_\infty(\mathbf{x}) = \int_0^{L_x} \left( G(\mathbf{x}, \boldsymbol{\xi}) - \pi_{N_c} \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \right) k^s(\boldsymbol{\xi}) d\boldsymbol{\xi} \quad \cdots (I) \quad (109)$$

$$15 \quad + \int_0^{L_x} \pi_{N_c} \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \left( k^s(\boldsymbol{\xi}) - \tilde{k}^s \delta(\boldsymbol{\xi}, 0) \right) d\boldsymbol{\xi} \quad \cdots (II)$$

$$16 \quad - \int_0^{L_x} (I_m - \pi_{N_c}) \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \tilde{k}^s \delta(\boldsymbol{\xi}, 0) d\boldsymbol{\xi} \quad \cdots (III).$$

17 Using (100) and (107), the first part in (109) is computed as

$$18 \quad (I) = \int_0^{L_x} \left( G(\mathbf{x}, \boldsymbol{\xi}) - \pi_{N_c} \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \right) k^s(\boldsymbol{\xi}) d\boldsymbol{\xi}$$

$$19 \quad = \int_0^{L_x} \sum_{1, 0 < l < N_c} \left( \frac{1}{\lambda_1} q_l(\mathbf{x}) q_l(\boldsymbol{\xi}) Q_1 - \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \tilde{Q}_1 \right) k^s(\boldsymbol{\xi}) d\boldsymbol{\xi}. \quad (110)$$

20 In (110), the first term in the parenthesis with  $l = N_c$  is canceled out by the second term in the parenthesis with  
21  $l = 0$ . Using (99), (I) is computed as

$$22 \quad (I) = \int_0^{L_x} \sum_{1, 0 < l < N_c} \left( \frac{1}{\lambda_1} q_l(\mathbf{x}) q_l(\boldsymbol{\xi}) Q_1 - \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \tilde{Q}_1 \right) \frac{1}{N_A V_c} [\mathbf{k}^s]^{(1)} I_{\{[0, \frac{L_x}{N_c}]\}}(\boldsymbol{\xi}) d\boldsymbol{\xi}$$

$$23 \quad = \frac{1}{N_A V} \max_i \mathbf{k}_{i(1,i)}^s \int_0^{\frac{L_x}{N_c}} \sum_{1, 0 < l < N_c} N_c \left( \frac{1}{\lambda_1} q_l(\mathbf{x}) q_l(\boldsymbol{\xi}) Q_1 - \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \tilde{Q}_1 \right) \mathbf{u}_m d\boldsymbol{\xi},$$

24 where  $\mathbf{u}_m$  is an  $m$ -dimensional vector with each element equal to 1. We break (I) into four parts by adding and  
25 subtracting terms as

$$26 \quad ABCD - \tilde{A}\tilde{B}\tilde{C}\tilde{D} = (A - \tilde{A})BCD + \tilde{A}(B - \tilde{B})CD + \tilde{A}\tilde{B}(C - \tilde{C})D + \tilde{A}\tilde{B}\tilde{C}(D - \tilde{D}).$$



1 Then, we get

$$\begin{aligned}
2 \quad (I) &= \frac{1}{\mathcal{N}_A \mathbf{V}} \max_i \mathbf{k}_{i(1,i)}^s \\
3 &\quad \times \left[ \int_0^{\frac{L_x}{N_c}} \sum_{1,0 < l < N_c} N_c \left( \frac{1}{\lambda_1} - \frac{1}{\tilde{\lambda}_1} \right) q_l(\mathbf{x}) q_l(\boldsymbol{\xi}) Q_1 \mathbf{u}_m d\boldsymbol{\xi} \quad \dots (i) \right. \\
4 &\quad + \int_0^{\frac{L_x}{N_c}} \sum_{1,0 < l < N_c} N_c \frac{1}{\tilde{\lambda}_1} (q_l(\mathbf{x}) - \tilde{q}_l(\mathbf{x})) q_l(\boldsymbol{\xi}) Q_1 \mathbf{u}_m d\boldsymbol{\xi} \quad \dots (ii) \\
5 &\quad + \int_0^{\frac{L_x}{N_c}} \sum_{1,0 < l < N_c} N_c \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) (q_l(\boldsymbol{\xi}) - \tilde{q}_l(\boldsymbol{\xi})) Q_1 \mathbf{u}_m d\boldsymbol{\xi} \quad \dots (iii) \\
6 &\quad \left. + \int_0^{\frac{L_x}{N_c}} \sum_{1,0 < l < N_c} N_c \frac{1}{\tilde{\lambda}_1} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) (Q_1 - \tilde{Q}_1) \mathbf{u}_m d\boldsymbol{\xi} \right] \quad \dots (iv).
\end{aligned}$$

7 **Step 3:**

8 We will show that  $\|(ii)\|_E \leq O\left(\frac{\ln N_c}{N_c}\right)$ . Using (98) and (83), for each  $l$  where  $0 < l < N_c$  we get

$$\begin{aligned}
9 \quad |q_l(\mathbf{x})| &= \left| \sqrt{\frac{2}{L_x}} \sum_{l'=1}^{N_c} \cos\left(\frac{(2l'-1)\pi l}{2N_c}\right) I_{\left[\frac{(l'-1)L_x}{N_c}, \frac{l'L_x}{N_c}\right]}(\mathbf{x}) \right| \\
10 &\leq O(1). \tag{111}
\end{aligned}$$

11 Next, using (98) and (105), for each  $l$  where  $0 < l < N_c$  and for  $\mathbf{x} \in \left[\frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c}\right)$ , a difference between  $q_l(\mathbf{x})$   
12 and  $\tilde{q}_l(\mathbf{x})$  is estimated as

$$\begin{aligned}
13 \quad |q_l(\mathbf{x}) - \tilde{q}_l(\mathbf{x})| &= \left| \sqrt{\frac{2}{L_x}} \sum_{l'=1}^{N_c} \cos\left(\frac{(2l'-1)\pi l}{2N_c}\right) I_{\left[\frac{(l'-1)L_x}{N_c}, \frac{l'L_x}{N_c}\right]}(\mathbf{x}) - \sqrt{\frac{2}{L_x}} \cos\left(\frac{l\pi \mathbf{x}}{L_x}\right) \right| \\
14 &= \sqrt{\frac{2}{L_x}} \left| \cos\left(\frac{(2k-1)\pi l}{2N_c}\right) - \cos\left(\frac{l\pi \mathbf{x}}{L_x}\right) \right| \\
15 &= \sqrt{\frac{2}{L_x}} \left| \left(\frac{(2k-1)\pi l}{2N_c} - \frac{l\pi \mathbf{x}}{L_x}\right) \sin b \right| \tag{112}
\end{aligned}$$

$$\begin{aligned}
16 &\leq \sqrt{\frac{2}{L_x}} \cdot \frac{l\pi}{2N_c} \cdot |\sin b| \\
17 &\equiv O\left(\frac{1}{N_c}\right) \times l. \tag{113}
\end{aligned}$$

18 (112) is obtained by the Mean value theorem where  $b$  is between  $\frac{(2k-1)\pi l}{2N_c}$  and  $\frac{l\pi \mathbf{x}}{L_x}$ . Then using (111) and (113),  
19  $\|(ii)\|_E$  is estimated as

$$\begin{aligned}
20 \quad \|(ii)\|_E &= \left\| \int_0^{\frac{L_x}{N_c}} \sum_{1,0 < l < N_c} N_c \frac{1}{\tilde{\lambda}_1} (q_l(\mathbf{x}) - \tilde{q}_l(\mathbf{x})) q_l(\boldsymbol{\xi}) Q_1 \mathbf{u}_m d\boldsymbol{\xi} \right\|_E \\
21 &\leq \left\| \sum_{1,0 < l < N_c} |\tilde{\lambda}_1^{-1}| \int_0^{\frac{L_x}{N_c}} N_c \times O\left(\frac{1}{N_c}\right) \times l \times O(1) d\boldsymbol{\xi} \right\|_E \max_1 \|Q_1 \mathbf{u}_m\|_E \\
22 &\leq O\left(\frac{1}{N_c}\right) \sum_{1,0 < l < N_c} l \cdot |\tilde{\lambda}_1^{-1}| \max_1 \|Q_1\|_E. \tag{114}
\end{aligned}$$

23 Now, we will show that  $\max_1 \|Q_1\|_E \leq O(1)$  by showing  $\max_1 \|\tilde{Q}_1\|_E \leq O(1)$ , because  $\tilde{Q}_1 - Q_1 = o\left(\frac{1}{N_c}\right)$  as we will  
24 show later in (118). To estimate difference between  $Q_1$  and  $\tilde{Q}_1$ , we use a perturbation theory of the linear operator  
25 in a finite-dimensional space (Theorem 5.4, p111 [24]). Define the operator  $A = \mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}$ , and express  $\mathcal{K} + \alpha_l \mathcal{D}$  as a

1 series of perturbed operators from  $A$ . Using (82) and (104), we obtain

$$\begin{aligned}
2 \quad A(\epsilon) &= A - \tilde{\alpha}_l \tilde{\mathcal{D}} + \alpha_l \mathcal{D} \\
3 \quad &= A - \tilde{\alpha}_l \tilde{\mathcal{D}} + \alpha_l \frac{\tilde{\mathcal{D}}}{(L_x/N_c)^2} \\
4 \quad &= A + \left(\frac{l\pi}{L_x}\right)^2 \tilde{\mathcal{D}} - \left(\frac{2N_c \sin\left(\frac{l\pi}{2N_c}\right)}{L_x}\right)^2 \tilde{\mathcal{D}} \\
5 \quad &= A + \left(\frac{l\pi}{L_x}\right)^2 \tilde{\mathcal{D}} - \left(\frac{\sin\left(\frac{l\pi\epsilon}{2}\right)}{\left(\frac{L_x\epsilon}{2}\right)}\right)^2 \tilde{\mathcal{D}}
\end{aligned}$$

6 where  $\epsilon = \frac{1}{N_c}$ . We can show that  $A(\epsilon)$  is differentiable at  $\epsilon = 0$  and  $A'(0) = 0$ . Let  $\lambda$  be the semi-simple eigenvalue  
7 of  $A$  and let  $Q$  be the projection of  $A$  corresponding to  $\lambda$ . Then, using Theorem 5.4 in [24], we get

$$8 \quad \lambda(\epsilon) = \lambda + o(\epsilon), \quad (115)$$

$$9 \quad Q(\epsilon) = Q + o(\epsilon), \quad (116)$$

10 where  $\lambda(\epsilon)$  and  $Q(\epsilon)$  are eigenvalues and projections of  $A(\epsilon)$ . From (115) and (116),  $\frac{\lambda(\epsilon)-\lambda}{\epsilon}$  and  $\left\|\frac{Q(\epsilon)-Q}{\epsilon}\right\|_E$  go to zero  
11 as  $\epsilon \rightarrow 0$ . In our setting,  $\lambda = \tilde{\lambda}_1$  and  $\lambda(\epsilon) = \lambda_1$ . Similarly,  $Q = \tilde{Q}_1$  and  $Q(\epsilon) = Q_1$ . Since we assume that  $A = \mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}$   
12 is semi-simple, the condition that  $\lambda$  is the semi-simple eigenvalue of  $A$  is satisfied. Then, (115) and (116) yields

$$13 \quad \tilde{\lambda}_1 - \lambda_1 = o\left(\frac{1}{N_c}\right), \quad (117)$$

$$14 \quad \tilde{Q}_1 - Q_1 = o\left(\frac{1}{N_c}\right). \quad (118)$$

15 Using (118) and  $\|Q_1 \mathbf{u}_m\|_E \leq \|Q_1\|_E \|\mathbf{u}_m\|_E = \sqrt{m} \|Q_1\|_E$  where  $m$  is the number of diffusing species, we only need  
16 to show that

$$17 \quad \max_1 \|\tilde{Q}_1\|_E = O(1).$$

18 Define the operator  $B = \tilde{\alpha}_l \tilde{\mathcal{D}}$  and consider  $\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}$  as perturbation of  $B$  as follows:

$$19 \quad B(\epsilon) = \left(-\mathcal{K} \left(\frac{L_x\epsilon}{\pi}\right)^2 \tilde{\mathcal{D}}^{-1} + I\right) \tilde{\alpha}_l \tilde{\mathcal{D}}$$

20 where  $\epsilon = \frac{1}{l}$ . As  $\epsilon \rightarrow 0$ ,  $B(\epsilon) \rightarrow B$  and  $B'(0) = 0$ . Therefore, using Theorem 5.4 in [24],

$$21 \quad J(\epsilon) - J = o(\epsilon)$$

22 where  $J(\epsilon)$  and  $J$  are projections of  $B(\epsilon)$  and  $B$ . Define  $M > 0$  large enough so that  $\epsilon = \frac{1}{l}$  is small for  $l \geq M$ .  $e_\mu$   
23 denotes an  $m$ -dimensional vector with its  $\mu^{th}$  entry equal to 1 and all others equal to zero. Since  $B\left(\frac{1}{l}\right) = \mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}$ ,  
24  $J\left(\frac{1}{l}\right) = \tilde{Q}_1$ , and  $J = e_\mu * e_\mu$ , for  $l \geq M$  we estimate the upper bound of  $\|\tilde{Q}_1\|_E$  as

$$\begin{aligned}
25 \quad \|\tilde{Q}_1\|_E &\leq \|\tilde{Q}_1 - Q\|_E + \|Q\|_E \\
26 \quad &= o\left(\frac{1}{l}\right) + \|e_\mu * e_\mu\|_E \\
27 \quad &\leq o\left(\frac{1}{M}\right) + 1, \quad \text{for } l \geq M.
\end{aligned}$$

1 Since  $\tilde{Q}_1$  and  $M$  do not depend on  $N_c$  where  $\mathbf{l} = (l, \mu)$ , we get

$$\begin{aligned} 2 \quad \max_{\mathbf{l}} \|\tilde{Q}_1\|_E &\leq \max \left( \max_{l < M, \mu} \|\tilde{Q}_{(l, \mu)}\|_E, o\left(\frac{1}{M}\right) + 1 \right) \\ 3 \quad &= O(1). \end{aligned} \quad (119)$$

4 **Step 4:**

5 Next, we estimate  $\sum_{\mu} |\tilde{\lambda}_1^{-1}|$ . Since  $\tilde{\lambda}_1$  is an eigenvalue of  $\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}}$ ,  $\tilde{\lambda}_1^{-1}$  is an eigenvalue of  $(\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1}$ . For  $l > 0$ ,  
6  $\tilde{\alpha}_l \neq 0$  and we expand the inverse matrix as

$$\begin{aligned} 7 \quad (\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} &= \left[ (\tilde{\alpha}_l \tilde{\mathcal{D}}) \left( I + (\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \mathcal{K} \right) \right]^{-1} \\ 8 \quad &= \left( I + (\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \mathcal{K} \right)^{-1} (\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \\ 9 \quad &= \left[ \sum_{p=0}^{\infty} \left( -(\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \mathcal{K} \right)^p \right] (\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1}. \end{aligned} \quad (120)$$

10 We estimate the upper and lower bounds of the spectrum of  $-(\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \mathcal{K}$  to show convergence of the matrix series  
11 in (120). Using the fact that all diagonal elements of  $\tilde{\mathcal{D}}$  are positive and using the definition of  $\tilde{\alpha}_l$ , we get

$$\begin{aligned} 12 \quad \left| \sigma \left( -(\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \mathcal{K} \right) \right| &\leq |\tilde{\alpha}_l^{-1}| \frac{1}{\min_x \tilde{\mathcal{D}}_{xx}} |\sigma(\mathcal{K})| \\ 13 \quad &= \left( \frac{L_x}{l\pi} \right)^2 \frac{1}{\min_x \tilde{\mathcal{D}}_{xx}} |\sigma(\mathcal{K})|. \end{aligned}$$

14 Therefore, there exists  $A > 0$  such for  $l \geq A$

$$15 \quad \left| \sigma \left( -(\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \mathcal{K} \right) \right| \leq 1.$$

16 Since the spectrum of  $-(\tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \mathcal{K}$  is bounded by  $\pm 1$  for  $l \geq A$ , the matrix series in (120) converges. From Levy-  
17 Hadamard theorem [8], each eigenvalue,  $\tilde{\lambda}_1^{-1}$ , of  $(\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1}$  satisfies

$$18 \quad \left| \tilde{\lambda}_1^{-1} - \left[ (\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \right]_{xx} \right| < \sum_{\gamma \neq x} \left| \left[ (\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \right]_{x\gamma} \right|$$

19 and the inequality can be rewritten as

$$20 \quad \left| \left[ (\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \right]_{xx} - \sum_{\gamma \neq x} \left| \left[ (\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \right]_{x\gamma} \right| \right| < |\tilde{\lambda}_1^{-1}| < \left| \left[ (\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \right]_{xx} \right| + \sum_{\gamma \neq x} \left| \left[ (\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \right]_{x\gamma} \right|. \quad (121)$$

21 Using (121) and (120), for each  $l \geq A$  we get

$$\begin{aligned} 22 \quad \sum_{\mu} |\tilde{\lambda}_1^{-1}| &< \sum_{\mu=1}^m \max_x \sum_{\gamma} \left| \left[ (\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}})^{-1} \right]_{x\gamma} \right| \\ 23 \quad &< m \left( \frac{L_x}{l\pi} \right)^2 \max_x \sum_{\gamma} \sum_{p=0}^{\infty} \left| \left[ \left( \left( \frac{L_x}{l\pi} \right)^2 \tilde{\mathcal{D}}^{-1} \mathcal{K} \right)^p \right]_{x\gamma} \right| \tilde{\mathcal{D}}_{\gamma\gamma}^{-1} \\ 24 \quad &\leq m \left( \frac{L_x}{l\pi} \right)^2 \max_x \sum_{\gamma} \left[ I + O\left(\frac{1}{l^2}\right) \right]_{x\gamma} \tilde{\mathcal{D}}_{\gamma\gamma}^{-1} \end{aligned}$$

$$\leq \tilde{C} \left( \frac{1}{l} \right)^2,$$

where  $m$  is the number of diffusing species. Define  $C \equiv \max \left( \tilde{C}, \max_{l < A} \left( mA^2 \left| \tilde{\lambda}_1^{-1} \right| \right) \right)$ . Then, for each  $l > 0$  the eigenvalues are bounded as

$$\sum_{\mu} |\tilde{\lambda}_1^{-1}| \leq \frac{C}{l^2} = O \left( \frac{1}{l^2} \right). \quad (122)$$

Therefore, using (119) and (122) in (114), we get

$$\begin{aligned} \|(ii)\|_E &\leq O \left( \frac{1}{N_c} \right) \sum_{1,0 < l < N_c} l \cdot |\tilde{\lambda}_1^{-1}| \max_l \|Q_l\|_E \\ &\leq O \left( \frac{1}{N_c} \right) \sum_{1,0 < l < N_c} l \times O \left( \frac{1}{l^2} \right) \times O(1) \\ &\leq O \left( \frac{1}{N_c} \right) \sum_{1,0 < l < N_c} O \left( \frac{1}{l} \right) \\ &\leq O \left( \frac{1}{N_c} \right) O \left( 1 + \int_1^{N_c-1} \frac{1}{y} dy \right) \\ &\leq O \left( \frac{\ln N_c}{N_c} \right). \end{aligned} \quad (123)$$

Now, we estimate an upper bound for  $\|(iii)\|_E$ . Using (105), for  $l > 0$  we get

$$\begin{aligned} |\tilde{q}_l(\mathbf{x})| &= \left| \sqrt{\frac{2}{L_x}} \cos \left( \frac{l\pi \mathbf{x}}{L_x} \right) \right| \\ &\leq O(1). \end{aligned} \quad (124)$$

Since the estimates in (111), (113), and (124) do not depend on  $\mathbf{x}$  and since the integrands in (ii) and (iii) are the same vice versa when we exchange  $\mathbf{x}$  and  $\boldsymbol{\xi}$ , using (123) we can get

$$\|(iii)\|_E \leq O \left( \frac{\ln N_c}{N_c} \right). \quad (125)$$

### Step 5:

We will show that  $\|(i)\|_E \leq o \left( \frac{1}{N_c} \right)$ . Using (111), (118), and (119), we get

$$\begin{aligned} \|(i)\|_E &= \left\| \int_0^{\frac{L_x}{N_c}} \sum_{1,0 < l < N_c} N_c \left( \frac{1}{\lambda_1} - \frac{1}{\tilde{\lambda}_1} \right) q_l(\mathbf{x}) q_l(\boldsymbol{\xi}) Q_l \mathbf{u}_m d\boldsymbol{\xi} \right\|_E \\ &\leq \left| \int_0^{\frac{L_x}{N_c}} N_c \times O(1) d\boldsymbol{\xi} \sum_{1,0 < l < N_c} \frac{\tilde{\lambda}_1 - \lambda_1}{\lambda_1 \tilde{\lambda}_1} \right| \times O(1) \\ &= O(1) \times \sum_{1,0 < l < N_c} \left| \frac{\tilde{\lambda}_1 - \lambda_1}{\lambda_1 \tilde{\lambda}_1} \right|. \end{aligned} \quad (126)$$

Then using (117) and (122) in (126), we get

$$\|(i)\|_E \leq \sum_{1,0 < l < N_c} \left| \frac{o \left( \frac{1}{N_c} \right)}{\tilde{\lambda}_1 \left( \tilde{\lambda}_1 + o \left( \frac{1}{N_c} \right) \right)} \right|$$

$$\begin{aligned}
& \leq \sum_{0 < l < N_c} \left| \frac{o\left(\frac{1}{N_c}\right)}{\left(l^4 + o\left(\frac{1}{N_c}\right)\right)} \right| \\
& \leq \frac{o\left(\frac{1}{N_c}\right)}{\left(1 + o\left(\frac{1}{N_c}\right)\right)} + \int_1^{N_c-1} \frac{o\left(\frac{1}{N_c}\right)}{\left(y^4 + o\left(\frac{1}{N_c}\right)\right)} dy \\
& \leq \frac{o\left(\frac{1}{N_c}\right)}{\left(1 + o\left(\frac{1}{N_c}\right)\right)} + \frac{o\left(\frac{1}{N_c}\right)}{(N_c - 1)^3} \leq o\left(\frac{1}{N_c}\right).
\end{aligned} \tag{127}$$

Next, We will show that  $\|(iv)\|_E \leq o\left(\frac{1}{N_c}\right)$ . Using (111), (124), and (118), we get

$$\begin{aligned}
\|(iv)\|_E &= \left\| \int_0^{\frac{L_x}{N_c}} \sum_{1,0 < l < N_c} N_c \frac{1}{\tilde{\lambda}_l} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \left(Q_1 - \tilde{Q}_1\right) \mathbf{u}_m d\boldsymbol{\xi} \right\|_E \\
&= \left| \int_0^{\frac{L_x}{N_c}} N_c \times O(1) d\boldsymbol{\xi} \sum_{1,0 < l < N_c} \frac{1}{\tilde{\lambda}_l} \right| \times o\left(\frac{1}{N_c}\right) \\
&= \sum_{1,0 < l < N_c} \left| \tilde{\lambda}_l^{-1} \right| \times o\left(\frac{1}{N_c}\right).
\end{aligned} \tag{128}$$

Using (122) in (128), we get

$$\begin{aligned}
\|(iv)\|_E &\leq \sum_{l,0 < l < N_c} O\left(\frac{1}{l^2}\right) \times o\left(\frac{1}{N_c}\right) \\
&\leq O\left(1 + \int_1^{N_c-1} \frac{1}{y^2} dy\right) \times o\left(\frac{1}{N_c}\right) \\
&\leq o\left(\frac{1}{N_c}\right).
\end{aligned} \tag{129}$$

### Step 6:

We will show that  $\|(II)\|_E \leq O\left(\frac{1}{N_c}\right)$ . Using (107), (99), and (97), we get

$$\begin{aligned}
(II) &= \int_0^{L_x} \pi_{N_c} \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \left(k^s(\boldsymbol{\xi}) - \tilde{k}^s \delta(\boldsymbol{\xi}, 0)\right) d\boldsymbol{\xi} \\
&= \int_0^{L_x} \sum_{1,l < N_c} \frac{1}{\tilde{\lambda}_l} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \tilde{Q}_1 \left(\frac{1}{\mathcal{N}_A \mathbf{V}_c} [\mathbf{k}^s]^{(1)} I_{\left\{[0, \frac{L_x}{N_c}]\right\}}(\boldsymbol{\xi}) - \frac{L_x}{\mathcal{N}_A \mathbf{V}} [\mathbf{k}^s]^{(1)} \delta(\boldsymbol{\xi}, 0)\right) d\boldsymbol{\xi}.
\end{aligned} \tag{130}$$

In (130), the term with  $l = 0$  in the series becomes zero. Using (122), (124), (119) in (130), we get

$$\begin{aligned}
\|(II)\|_E &\leq \sum_{l,0 < l < N_c} O\left(\frac{1}{l^2}\right) \times O(1) \times \frac{L_x}{\mathcal{N}_A \mathbf{V}} \max_i \mathbf{k}_{i(1,i)}^s \times \left| \int_0^{L_x} \tilde{q}_l(\boldsymbol{\xi}) \left(\frac{N_c}{L_x} I_{\left\{[0, \frac{L_x}{N_c}]\right\}}(\boldsymbol{\xi}) - \delta(\boldsymbol{\xi}, 0)\right) d\boldsymbol{\xi} \right| \\
&\leq \sum_{l,0 < l < N_c} O\left(\frac{1}{l^2}\right) \times \left| \int_0^{L_x} \tilde{q}_l(\boldsymbol{\xi}) \left(\frac{N_c}{L_x} I_{\left\{[0, \frac{L_x}{N_c}]\right\}}(\boldsymbol{\xi}) - \delta(\boldsymbol{\xi}, 0)\right) d\boldsymbol{\xi} \right|.
\end{aligned} \tag{131}$$

Using (105) in (131), we estimate

$$\|(II)\|_E \leq \sum_{l,0 < l < N_c} O\left(\frac{1}{l^2}\right) \times \sqrt{\frac{2}{L_x}} \left| \frac{N_c}{l\pi} \sin\left(\frac{l\pi}{N_c}\right) - 1 \right|. \tag{132}$$

1 By approximating the series in (132), we get

$$\begin{aligned}
2 \quad \|(II)\|_E &\leq O(1) \times \sum_{l, 0 < l < N_c} \frac{\left| 1 - \frac{\sin\left(\frac{l\pi}{N_c}\right)}{\left(\frac{l\pi}{N_c}\right)} \right|}{l^2} \\
3 \quad &\leq O(1) \times \sum_{l, 0 < l < N_c} \frac{1 - \frac{\left(\frac{l\pi}{N_c}\right) - \frac{\left(\frac{l\pi}{N_c}\right)^3}{3!}}{\left(\frac{l\pi}{N_c}\right)}}{l^2} \\
4 \quad &\leq O\left(\frac{1}{N_c}\right). \tag{133}
\end{aligned}$$

5 Now, we will show that  $\|(III)\|_E \leq O\left(\frac{1}{N_c}\right)$ . Using (108) and (97), we compute

$$\begin{aligned}
6 \quad (III) &= \int_0^{L_x} \left( I_m - \pi_{N_c} \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) \right) \tilde{k}^s \delta(\boldsymbol{\xi}, 0) d\boldsymbol{\xi} \\
7 \quad &= \int_0^{L_x} \left( \sum_{l, l \geq N_c} \frac{1}{\lambda_l} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(\boldsymbol{\xi}) \tilde{Q}_1 \right) \frac{L_x}{\mathcal{N}_A \mathbf{V}} [\mathbf{k}^s]^{(1)} \delta(\boldsymbol{\xi}, 0) d\boldsymbol{\xi} \\
8 \quad &= \sum_{l, l \geq N_c} \left( \frac{1}{\lambda_l} \tilde{q}_l(\mathbf{x}) \tilde{q}_l(0) \tilde{Q}_1 \right) \frac{L_x}{\mathcal{N}_A \mathbf{V}} [\mathbf{k}^s]^{(1)}. \tag{134}
\end{aligned}$$

9 Using (124), (122), and (119) in (134), we get

$$\begin{aligned}
10 \quad \|(III)\|_E &\leq \sum_{l \geq N_c} O\left(\frac{1}{l^2}\right) \times O(1) \times \max_i \mathbf{k}_{i(1,i)}^s \\
11 \quad &\leq O(1) \times \int_{N_c-1}^{\infty} \frac{1}{y^2} dy \\
12 \quad &\leq O\left(\frac{1}{N_c}\right). \tag{135}
\end{aligned}$$

13 In conclusion, we prove that for any  $\mathbf{x} \in [0, L_x]$ ,  $\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_E \leq O\left(\frac{\ln N_c}{N_c}\right)$  as shown in (123), (125), (127),  
14 (129), (133), and (135). Since the upper bound is independent of  $\mathbf{x}$ , we get  $\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_{L_2} = O\left(\frac{\ln N_c}{N_c}\right)$ .  $\square$

15 Using Lemma 1, we prove the convergence of  $U_\infty(\mathbf{x})$  to  $u_\infty(\mathbf{x})$ , and this implies the convergence of  $\frac{1}{\mathcal{N}_A \mathbf{V}_c} X_\infty$  as  
16  $N_c \rightarrow \infty$ . Since  $Y_\infty$  is expressed in terms of  $X_\infty$ ,  $\overline{M}_\infty$  converges as  $N_c \rightarrow \infty$ , and this implies the convergence of  
17  $\frac{CV^*}{CV^*}$ .

## 18 E. Proof of Theorem 2

19 Before proving the theorem, we first state and prove a lemma which is used to prove the theorem. The proof of the  
20 theorem is given at the end of this appendix. In the following lemma, we prove that each component of  $\overline{M}_\infty/\mathcal{N}_A$   
21 converges to the corresponding value of the steady-state concentration in the continuum deterministic reaction-  
22 diffusion system.

23 The mean vector of the stochastic system with no inputs is governed by

$$24 \quad \frac{dM(t)}{dt} = \Omega M(t), \tag{136}$$

25 where  $\Omega = \Delta \otimes \mathcal{D} + I_{N_c} \otimes \mathcal{K}$ . The corresponding deterministic reaction-diffusion system is governed by

$$\begin{aligned}
26 \quad \frac{\partial u(\mathbf{x}, t)}{\partial t} &= \tilde{D} \Delta u(\mathbf{x}, t) + \mathcal{K} u(\mathbf{x}, t), \quad \mathbf{x} \in [0, L_x], \\
27 \quad \frac{\partial u(\mathbf{x}, t)}{\partial x} &= 0, \quad \mathbf{x} = 0, L_x,
\end{aligned} \tag{137}$$

$$u(\mathbf{x}, 0) = u_0(\mathbf{x}),$$

where  $u(\mathbf{x}, t)$  is a concentration vector for species in  $\mathbf{x} \in [0, L_x]$  at time  $t$ . In (137), the diffusion matrix is given as  $\tilde{\mathcal{D}} = (L_x/N_c)^2 \mathcal{D}$ . The integrated initial concentration,  $u_0(\mathbf{x})$ , between  $\mathbf{x} \in \left[ \frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c} \right)$  is expressed in terms of  $M(0)$  as follows:

$$\int_{\frac{(k-1)L_x}{N_c}}^{\frac{kL_x}{N_c}} (u_0(\boldsymbol{\xi}))_i d\boldsymbol{\xi} = \frac{L_x}{N_A \mathbf{V}} [M(0)]_{i(k,i)}. \quad (138)$$

**Lemma 2.** Let  $M_\infty$  be the steady-state solution of (136) and let  $u_\infty(\mathbf{x})$  be the steady-state solution of (137). Define  $U_\infty(\mathbf{x})$  as a vector with each component satisfying

$$(U_\infty(\mathbf{x}))_i \equiv \sum_{k=1}^{N_c} \frac{1}{N_A V_c} [M_\infty]_{i(k,i)} I_{\left\{ \left[ \frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c} \right] \right\}}(\mathbf{x}), \quad i = 1, \dots, s,$$

where  $s$  is the number of species. Then,  $\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_{L_2} = O\left(\frac{1}{N_c}\right)$  and converges to 0 as  $N_c \rightarrow \infty$ .

*Proof.* The proof of Lemma 2 is given in 2 steps:

- **Step 1** Express  $U_\infty(\mathbf{x})$  and  $u_\infty(\mathbf{x})$  in terms of the Green's function;
- **Step 2** Show  $\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_{L_2} = O\left(\frac{1}{N_c}\right)$ .

**Step 1:**

We first express  $U_\infty(\mathbf{x})$  in terms of the Green's function. Recall that  $\phi_{N_c}$  and  $\phi_{N_c}^*$  are the eigenvector and the adjoint eigenvector of  $\Delta$  corresponding to  $\alpha_{N_c} = 0$ . Rearrange the order of species in the mean vector,  $M(t)$ , so that that  $\varphi_{(N_c,s)}$  and  $\varphi_{(N_c,s)}^*$  become an eigenvector and an adjoint eigenvector corresponding to the zero eigenvalue of  $\Omega$ . Define

$$Q_{(N_c,s)} \equiv \varphi_{(N_c,s)} * \overline{\varphi_{(N_c,s)}^*}. \quad (139)$$

Then, the projection corresponding to the zero eigenvalue is expressed as

$$P_s = (\phi_{N_c} * \overline{\phi_{N_c}^*}) \otimes Q_{(N_c,s)}. \quad (140)$$

Letting  $t \rightarrow \infty$  in (90) and using (140), the steady-state mean vector for species numbers is expressed as

$$\begin{aligned} M_\infty &= P_s M(0) \\ &= (\phi_{N_c} * \overline{\phi_{N_c}^*}) \otimes Q_{(N_c,s)} M(0). \end{aligned}$$

Here, note that

$$\begin{aligned} \phi_{N_c} &= \overline{\phi_{N_c}^*} \\ &= \sqrt{\frac{1}{N_c}} [1, 1, \dots, 1]^T. \end{aligned} \quad (141)$$

We define a spatial variable  $q_{N_c}(\mathbf{x})$  as

$$\begin{aligned} q_{N_c}(\mathbf{x}) &\equiv \sqrt{\frac{N_c}{L_x}} \sum_{l'=1}^{N_c} (\phi_{N_c})_{l'} I_{\left\{ \left[ \frac{(l'-1)L_x}{N_c}, \frac{l'L_x}{N_c} \right] \right\}}(\mathbf{x}) \\ &= \sqrt{\frac{1}{L_x}}. \end{aligned} \quad (142)$$

where the last equality comes from (141). Define a matrix for a discrete Green's function as follows, and using (142) we get

$$G(\mathbf{x}, \boldsymbol{\xi}) \equiv q_{N_c}(\mathbf{x}) q_{N_c}(\boldsymbol{\xi}) Q_{(N_c,s)}$$

$$= \frac{1}{L_x} Q_{(N_c, s)} \equiv G. \quad (143)$$

Then,  $U_\infty(\mathbf{x})$  is expressed as

$$U_\infty(\mathbf{x}) = GU_0(\mathbf{x}). \quad (144)$$

Here,  $U_0(\mathbf{x})$  is defined as follows:

$$(U_0(\mathbf{x}))_i \equiv \sum_{k=1}^{N_c} \frac{1}{\mathcal{N}_A V_c} [M(0)]_{i(k, i)} I_{\left\{ \left[ \frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c} \right] \right\}}(\mathbf{x}), \quad 1 = 1, \dots, s. \quad (145)$$

Using (138), (145) is written as

$$(U_0(\mathbf{x}))_i = \sum_{k=1}^{N_c} \left( \frac{N_c}{L_x} \int_{\frac{(k-1)L_x}{N_c}}^{\frac{kL_x}{N_c}} (u_0(\boldsymbol{\xi}))_i d\boldsymbol{\xi} \right) I_{\left\{ \left[ \frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c} \right] \right\}}(\mathbf{x}), \quad 1 = 1, \dots, s. \quad (146)$$

Next, we express  $u_\infty(\mathbf{x})$  in terms of the Green's function. Let  $\tilde{\alpha}_0 = 0$  and  $\tilde{q}_0(\mathbf{x})$  be a solution of the scalar problem

$$\begin{aligned} \Delta \tilde{q}(\mathbf{x}) &= \tilde{\alpha}_0 \tilde{q}(\mathbf{x}), & \mathbf{x} &\in [0, L_x], \\ \tilde{q}'(\mathbf{x}) &= 0, & \mathbf{x} &= 0, L_x, \end{aligned}$$

satisfying  $\|\tilde{q}(\mathbf{x})\|_{L_2} = 1$ . Then, we get

$$\tilde{q}_0(\mathbf{x}) = \sqrt{\frac{1}{L_x}}. \quad (147)$$

Let  $\tilde{\lambda}_{(0, s)} = 0$  and  $\tilde{\varphi}_{(0, s)}$  be a solution of the algebraic eigenvalue problem

$$(\mathcal{K} + \tilde{\alpha}_l \tilde{\mathcal{D}} - \tilde{\lambda}_l I_s) \tilde{\varphi}_l = 0,$$

and let  $\tilde{\varphi}_{(0, s)}^*$  be the solution of the corresponding adjoint eigenvalue problem. Define

$$\tilde{Q}_{(0, s)} \equiv \tilde{\varphi}_{(0, s)} * \overline{\tilde{\varphi}_{(0, s)}^*},$$

and the Green's function as

$$\begin{aligned} \tilde{G}(\mathbf{x}, \boldsymbol{\xi}) &\equiv \tilde{q}_0(\mathbf{x}) \tilde{q}_0(\boldsymbol{\xi}) \tilde{Q}_{(0, s)} \\ &= \frac{1}{L_x} \tilde{Q}_{(0, s)} \equiv \tilde{G}, \end{aligned}$$

where (147) is used to compute  $\tilde{G}(\mathbf{x}, \boldsymbol{\xi})$ . Note that

$$\begin{aligned} Q_{(N_c, s)} &= \tilde{Q}_{(0, s)}, \\ G &= \tilde{G}. \end{aligned} \quad (148)$$

Then, the steady-state concentration vector is given as

$$u_\infty(\mathbf{x}) = \tilde{G}u_0(\mathbf{x}). \quad (149)$$

## Step 2:

We will show that  $\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_{L_2}$  is  $O\left(\frac{1}{N_c}\right)$ . Subtracting (149) from (144) and using (148), we get

$$\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_{L_2} = \left( \int_0^{L_x} \left\| GU_0(\mathbf{x}) - \tilde{G}u_0(\mathbf{x}) \right\|_E^2 d\mathbf{x} \right)^{1/2}$$



$$\leq \|G\|_E \left( \int_0^{L_x} \|U_0(\mathbf{x}) - u_0(\mathbf{x})\|_E^2 d\mathbf{x} \right)^{1/2}. \quad (150)$$

Using (143) and (139) we get

$$\begin{aligned} \|G\|_E &= \left\| \frac{1}{L_x} Q_{(N_c, s)} \right\|_E \\ &= \left\| \frac{1}{L_x} \left( \varphi_{(N_c, s)} * \overline{\varphi_{(N_c, s)}^*} \right) \right\|_E \\ &\leq O(1), \end{aligned} \quad (151)$$

where the last inequality comes from the fact that  $\varphi_{(N_c, s)}$  and  $\varphi_{(N_c, s)}^*$  do not depend on  $N_c$ , which are the eigenvector and adjoint eigenvector of  $\mathcal{K} + \alpha_l \mathcal{D}$  corresponding to  $\alpha_{N_c} = 0$ . Using (146) for  $\mathbf{x} \in \left[ \frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c} \right)$ , we have

$$\begin{aligned} |(U_0(\mathbf{x}))_i - (u_0(\mathbf{x}))_i|^2 &\leq \left| \frac{1}{L_x/N_c} \int_{\frac{(k-1)L_x}{N_c}}^{\frac{kL_x}{N_c}} [(u_0(\boldsymbol{\xi}))_i - (u_0(\mathbf{x}))_i] d\boldsymbol{\xi} \right|^2 \\ &\leq \max_{b \in \left[ \frac{(k-1)L_x}{N_c}, \frac{kL_x}{N_c} \right)} (u_0(b))_i^2 \left( \frac{L_x}{N_c} \right)^2 \\ &= O\left( \frac{1}{N_c^2} \right). \end{aligned} \quad (152)$$

When  $A$  is an  $m \times n$  matrix,  $\|A\|_E \leq \sqrt{mn} \|A\|_{\max}$ . Therefore, using (152) we get

$$\begin{aligned} \|U_0(\mathbf{x}) - u_0(\mathbf{x})\|_E &\leq \sqrt{s} \|U_0(\mathbf{x}) - u_0(\mathbf{x})\|_{\max} \\ &= \sqrt{s} \max_i |(U_0(\mathbf{x}))_i - (u_0(\mathbf{x}))_i| \\ &\leq O\left( \frac{1}{N_c^2} \right). \end{aligned} \quad (153)$$

Using (150), (151), and (153), we get

$$\|U_\infty(\mathbf{x}) - u_\infty(\mathbf{x})\|_{L_2} = O\left( \frac{1}{N_c} \right). \square$$

In Lemma 2, we prove the convergence of  $U_\infty(\mathbf{x})$  to  $u_\infty(\mathbf{x})$ . Now, we show the convergence of  $\overline{CV}^*$  as  $N_c \rightarrow \infty$ . Define  $M_d(0)^{(k)}$  and  $M_{d, \infty}^{(k)}$  be  $s \times s$  diagonal matrices where diagonal elements are given as

$$\begin{aligned} \left( M_d(0)^{(k)} \right)_{ii} &= [M_d(0)]_{i(k, i)}, \\ \left( M_{d, \infty}^{(k)} \right)_{ii} &= [M_{d, \infty}]_{i(k, i)}, \end{aligned}$$

for  $k = 1, \dots, N_c$  and for  $i = 1, \dots, s$ . Among the terms in (15), we first compute  $P_s M_d(0) P_s^T$  using (140) and (141) as follows:

$$\begin{aligned} P_s M_d(0) P_s^T &= \left( (\phi_{N_c} * \overline{\phi_{N_c}^*}) \otimes Q_{(N_c, s)} \right) M_d(0) \left( (\phi_{N_c} * \overline{\phi_{N_c}^*}) \otimes Q_{(N_c, s)}^T \right) \\ &= \frac{1}{N_c^2} (\mathbf{u}_{N_c} * \mathbf{u}_{N_c}) \otimes \sum_{p=1}^{N_c} Q_{(N_c, s)} M_d(0)^{(p)} Q_{(N_c, s)}^T, \end{aligned}$$

1 where  $\mathbf{u}_{N_c}$  is an  $N_c$ -dimensional vector with each element equal to 1. Then, each component of  $M_{d,\infty}^{-1} P_s M_d(0) P_s^T M_{d,\infty}^{-1}$   
 2 is written as

$$\begin{aligned} & \left[ M_{d,\infty}^{-1} P_s M_d(0) P_s^T M_{d,\infty}^{-1} \right]_{\mathbf{i}(k,i), \mathbf{i}(q,j)} \\ &= \frac{1}{N_c^2} \left[ \left( M_{d,\infty}^{(k)} \right)^{-1} \left( \sum_{p=1}^{N_c} Q_{(N_c,s)} M_d(0)^{(p)} Q_{(N_c,s)}^T \right) \left( M_{d,\infty}^{(q)} \right)^{-1} \right]_{ij}. \end{aligned} \quad (154)$$

4 Using the fact that sum of the initial mean species numbers in all compartments is bounded and using (151), we get

$$\begin{aligned} & \left\| \sum_{p=1}^{N_c} Q_{(N_c,s)} M_d(0)^{(p)} Q_{(N_c,s)}^T \right\|_E \leq \left\| \sum_{p=1}^{N_c} M_d(0)^{(p)} \right\|_E \|Q_{(N_c,s)}\|_E \|Q_{(N_c,s)}^T\|_E \\ & \leq O(1). \end{aligned} \quad (155)$$

7 Since each component of  $\left( M_{d,\infty}^{(k)} \right)^{-1}$  is bounded by  $O(N_c)$ , using (155) (154) is bounded by  $O(1)$ . Therefore, we get

$$\begin{aligned} \overline{CV}^* &= \sqrt{V_c \times \lambda_{\max} \left( M_{d,\infty}^{-1} - M_{d,\infty}^{-1} P_s M_d(0) P_s^T M_{d,\infty}^{-1} \right)} \\ &= \sqrt{\sigma \left( \left( \frac{M_{d,\infty}}{V_c} \right)^{-1} - O \left( \frac{1}{N_c} \right) \mathbf{u}_{sN_c} * \mathbf{u}_{sN_c} \right)} \\ &\rightarrow \sqrt{\frac{1}{\min_{i,\mathbf{x}} (u_\infty(\mathbf{x}))_i}} \end{aligned}$$

11 as  $N_c \rightarrow \infty$  where  $\mathbf{u}_{sN_c}$  is an  $sN_c$ -dimensional vector with each element equal to 1. Therefore, this gives the conver-  
 12 gence of  $\overline{CV}^*$ .

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