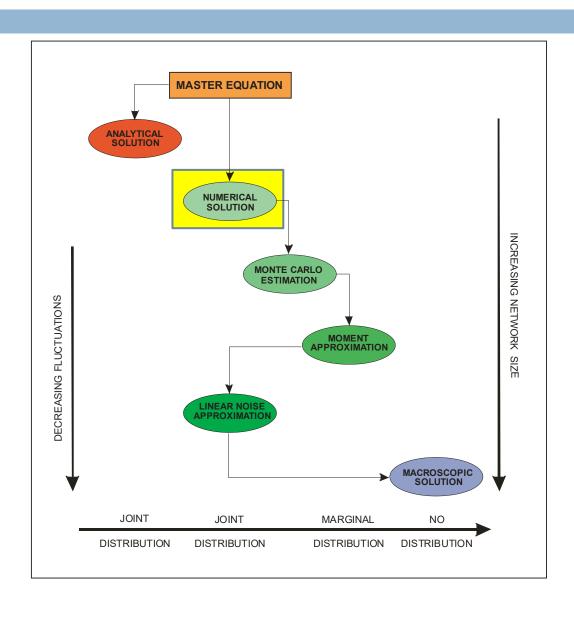
LECTURE #5

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Available Methods



Numerical Solution

The ME

$$\frac{\partial p_X(\mathbf{x};t)}{\partial t} = \sum_{m=1}^{M} \left\{ \pi_m(\mathbf{x} - \mathbf{s}_m) p_X(\mathbf{x} - \mathbf{s}_m;t) - \pi_m(\mathbf{x}) p_X(\mathbf{x};t) \right\}, \quad t > 0$$

of the population process can be expressed as a <u>linear</u> system of coupled first-order differential equations:

$$\frac{d\mathbf{p}(t)}{dt} = \mathbf{P}\mathbf{p}(t), \ t > 0$$

- $\mathbf{p}(t)$ is a $K \times 1$ vector that contains the nonzero probabilities $p_X(\mathbf{x};t)$.
- $lackbox{ extbf{P}}$ is a large $K \times K$ <u>sparse</u> matrix whose structure can be inferred from the ME.

Numerical Solution

- lacktriangle For large networks, matrix f P is <u>highly</u> sparse.
- When the columns of the <u>net</u> stoichiometric matrix S are all different from each other, the i-th column of P contains <u>at most</u> M+1 <u>nonzero</u> elements.
- The off-diagonal elements are given by

$$\pi_m(\mathbf{x}_i) > 0$$
, for $m = 1, 2, ..., M$,

where \mathbf{x}_i is the i-th state.

The diagonal element is given by

$$-\sum_{m=1}^{M} \pi_m(\mathbf{x}_i) < 0$$

 Consequently, the elements of each column of matrix P sum to zero.

Example

Consider a Markovian reaction network comprised of five species and the following five reactions (N = M = 5):

$$X_{1} + X_{2} \xrightarrow{\kappa_{1}x_{1}x_{2}} X_{3}$$

$$X_{3} \xrightarrow{\kappa_{2}x_{3}} X_{1} + X_{2}$$

$$X_{4} \xrightarrow{\kappa_{3}x_{4}} X_{3}$$

$$X_{4} \xrightarrow{\kappa_{4}x_{4}} X_{5}$$

$$X_{5} \xrightarrow{\kappa_{5}x_{5}} X_{4}$$

$$\mathbf{S} = \begin{bmatrix} -1 & 1 & 0 & 0 & 0 \\ -1 & 1 & 0 & 0 & 0 \\ 1 & -1 & 1 & 0 & 0 \\ 0 & 0 & -1 & -1 & 1 \\ 0 & 0 & 0 & 1 & -1 \end{bmatrix} \begin{cases} \mathbf{x}_1 & \mathbf{x}_2 \\ \mathbf{x}_3 & \mathbf{x}_4 \\ \mathbf{x}_4 & \mathbf{x}_5 \\ \mathbf{x}_4 & \mathbf{x}_5 \\ \mathbf{x}_6 & \mathbf{x}_7 \\ \mathbf{x}_8 & \mathbf{x}_9 \\ \mathbf{x}_9 & \mathbf{x}_9 \\ \mathbf{x}_{1} & \mathbf{x}_{1} \\ \mathbf{x}_{2} & \mathbf{x}_{1} \\ \mathbf{x}_{1} & \mathbf{x}_{2} \\ \mathbf{x}_{1} & \mathbf{x}_{2} \\ \mathbf{x}_{2} & \mathbf{x}_{3} \\ \mathbf{x}_{1} & \mathbf{x}_{2} \\ \mathbf{x}_{2} & \mathbf{x}_{3} \\ \mathbf{x}_{1} & \mathbf{x}_{2} \\ \mathbf{x}_{3} & \mathbf{x}_{4} \\ \mathbf{x}_{1} & \mathbf{x}_{2} \\ \mathbf{x}_{2} & \mathbf{x}_{3} \\ \mathbf{x}_{3} & \mathbf{x}_{4} \\ \mathbf{x}_{1} & \mathbf{x}_{2} \\ \mathbf{x}_{3} & \mathbf{x}_{4} \\ \mathbf{x}_{4} & \mathbf{x}_{4} \\ \mathbf{x}_{5} & \mathbf{x}_{5} \\ \mathbf{x}_{1} & \mathbf{x}_{2} \\ \mathbf{x}_{2} & \mathbf{x}_{3} \\ \mathbf{x}_{4} & \mathbf{x}_{5} \\ \mathbf{x}_{5} & \mathbf{x}_{5} \\ \mathbf{x}_{1} & \mathbf{x}_{2} \\ \mathbf{x}_{3} & \mathbf{x}_{4} \\ \mathbf{x}_{4} & \mathbf{x}_{5} \\ \mathbf{x}_{5} \\ \mathbf{x}_{5} & \mathbf{x}_{5} \\ \mathbf{x}_{5}$$

$$\begin{array}{c} \boldsymbol{X}_{1} + \boldsymbol{X}_{2} \xrightarrow{\kappa_{1} x_{1} x_{2}} \boldsymbol{X}_{3} \\ \boldsymbol{X}_{3} \xrightarrow{\kappa_{2} x_{3}} \boldsymbol{X}_{1} + \boldsymbol{X}_{2} \\ \boldsymbol{X}_{4} \xrightarrow{\kappa_{3} x_{4}} \boldsymbol{X}_{3} \\ \boldsymbol{X}_{4} \xrightarrow{\kappa_{4} x_{4}} \boldsymbol{X}_{5} \\ \boldsymbol{X}_{5} \xrightarrow{\kappa_{5} x_{5}} \boldsymbol{X}_{4} \end{array}$$

$$\begin{array}{c} \boldsymbol{S} = \begin{bmatrix} -1 & 1 & 0 & 0 & 0 \\ -1 & 1 & 0 & 0 & 0 \\ 1 & -1 & 1 & 0 & 0 \\ 0 & 0 & -1 & -1 & 1 \\ 0 & 0 & 0 & 1 & -1 \end{bmatrix} \begin{bmatrix} \boldsymbol{x}_{1} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 1 \end{bmatrix}, \boldsymbol{x}_{2} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 1 \end{bmatrix}, \boldsymbol{x}_{3} = \begin{bmatrix} 1 \\ 1 \\ 0 \\ 0 \\ 0 \end{bmatrix}, \boldsymbol{x}_{4} = \begin{bmatrix} 1 \\ 1 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

$$\begin{array}{c} \boldsymbol{x}_{1} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 1 \end{bmatrix}, \boldsymbol{x}_{2} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 1 \end{bmatrix}, \boldsymbol{x}_{3} = \begin{bmatrix} 1 \\ 1 \\ 0 \\ 0 \end{bmatrix}, \boldsymbol{x}_{4} = \begin{bmatrix} 1 \\ 1 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

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$$\mathbf{P} = \begin{bmatrix} -\kappa_5 & \kappa_4 & 0 & 0 \\ \kappa_5 & -(\kappa_3 + \kappa_4) & 0 & 0 \\ 0 & \kappa_3 & -\kappa_2 & \kappa \\ 0 & 0 & \kappa_2 & -\kappa_3 \end{bmatrix}$$

$$\mathbf{p}(t) = \begin{bmatrix} p(\mathbf{x}_1; t) \\ p(\mathbf{x}_2; t) \\ p(\mathbf{x}_3; t) \\ p(\mathbf{x}_4; t) \end{bmatrix}$$

Example

$$\pi_1(\mathbf{x}) = \kappa_1 x_1 x_2$$

$$\pi_2(\mathbf{x}) = \kappa_2 x_3$$

$$\pi_3(\mathbf{x}) = \kappa_3 x_4$$

$$\pi_4(\mathbf{x}) = \kappa_4 x_4$$

$$\pi_5(\mathbf{x}) = \kappa_5 x_5$$

$$\mathbf{x}_{1} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 1 \end{bmatrix}$$

$$\mathbf{x}_{1} - \mathbf{s}_{4} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 1 \end{bmatrix} - \begin{bmatrix} 0 \\ 0 \\ 0 \\ -1 \\ 1 \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 1 \\ 0 \end{bmatrix} = \mathbf{x}_{2}$$

impossible states

$$\frac{\partial p_{X}(\mathbf{x}_{1};t)}{\partial t} = \pi_{1}(\mathbf{x}_{1} - \mathbf{s}_{1}) p_{X}(\mathbf{x}_{1} - \mathbf{s}_{1};t) - \pi_{1}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
+ \pi_{2}(\mathbf{x}_{1} - \mathbf{s}_{2}) p_{X}(\mathbf{x}_{1} - \mathbf{s}_{2};t) - \pi_{2}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
+ \pi_{3}(\mathbf{x}_{1} - \mathbf{s}_{2}) p_{X}(\mathbf{x}_{1} - \mathbf{s}_{3};t) - \pi_{3}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
+ \pi_{4}(\mathbf{x}_{1} - \mathbf{s}_{4}) p_{X}(\mathbf{x}_{1} - \mathbf{s}_{4};t) - \pi_{4}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
+ \pi_{5}(\mathbf{x}_{1} - \mathbf{s}_{3}) p_{X}(\mathbf{x}_{1} - \mathbf{s}_{4};t) - \pi_{5}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
= \pi_{4}(\mathbf{x}_{1} - \mathbf{s}_{4}) p_{X}(\mathbf{x}_{1} - \mathbf{s}_{4};t) - \pi_{1}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
- \pi_{2}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t) - \pi_{3}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
- \pi_{4}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t) - \pi_{5}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
= \pi_{4}(\mathbf{x}_{1} - \mathbf{s}_{4}) p_{X}(\mathbf{x}_{1} - \mathbf{s}_{4};t) - \pi_{5}(\mathbf{x}_{1}) p_{X}(\mathbf{x}_{1};t)
= -\kappa_{5} p_{X}(\mathbf{x}_{1};t) + \kappa_{4} p_{X}(\mathbf{x}_{2};t)$$
 (first row of **P**)

zero propensities

Numerical Solution & FSP

□ Note that (assuming $K < \infty$)

$$\frac{d\mathbf{p}(t)}{dt} = \mathbf{P}\mathbf{p}(t) \iff \mathbf{p}(t) = \exp(t\mathbf{P})\mathbf{p}(0)$$

- Therefore, solving the ME is equivalent to evaluating the matrix exponential $\exp(t\mathbf{P})$.
- $lue{}$ Unfortunately, this computation is not possible, since the size of matrix $\bf P$ is prohibitively large.
- □ To reduce the size, we could employ a method known as <u>finite-state</u> <u>projection</u> (FSP) method.
- □ FSP requires appropriate truncation of the state-space to determine the possible (non-zero probability) set of states and development of a computationally feasible algorithm for calculating the matrix exponential.

- One of the best methods for computing matrix exponentials is known as the <u>Krylov subspace approximation</u> (KSA) method.
- For a sufficiently small time-step $\tau > 0$, this method approximates the vector $\mathbf{p}(t+\tau) = \exp(\tau \mathbf{P})\mathbf{p}(t)$ when \mathbf{P} is a <u>large</u> and <u>sparse</u> matrix.
- This is done by using a polynomial series expansion of the form:

$$\hat{\mathbf{p}}(t+\tau) = c_0 \mathbf{p}(t) + c_1 \tau \mathbf{P} \mathbf{p}(t) + \dots + c_{K_0-1} (\tau \mathbf{P})^{K_0-1} \mathbf{p}(t)$$

$$\hat{\mathbf{p}}(t+\tau) = c_0 \mathbf{p}(t) + c_1 \tau \mathbf{P} \mathbf{p}(t) + \dots + c_{K_0-1} (\tau \mathbf{P})^{K_0-1} \mathbf{p}(t)$$

- The coefficients $c_0, c_1, \dots, c_{K_0-1}$ are estimated by minimizing the <u>least-squares error</u> (LSE) $\|\mathbf{p}(t+\tau) \hat{\mathbf{p}}(t+\tau)\|_2^2$.
- $lue{}$ The optimal K_0 -order polynomial approximation of

 $\mathbf{p}(t+\tau)$ is a point in the K_0 -dimensional Krylov subspace:

$$\mathcal{K}(t) = \operatorname{span}\left\{\mathbf{p}(t), \tau \mathbf{P}\mathbf{p}(t), \dots, (\tau \mathbf{P})^{K_0 - 1}\mathbf{p}(t)\right\}$$

This element can be approximated by

$$\hat{\mathbf{p}}(t+\tau) = \|\mathbf{p}(t)\|_{2} \mathbf{V}(t) \exp\{\tau \mathbf{H}(t)\} \mathbf{e}_{1}$$

- oxdot e₁ is the first column of the $K_0 \times K_0$ identity matrix.
- $lackbox{$lackbox{V}(t)$ is a $K imes K_0$ matrix whose columns form an <u>orthonormal basis</u> for the Krylov subspace <math>\Re(t)$.
- \blacksquare $\mathbf{H}(t)$ is a $K_0 \times K_0$ Hessenberg matrix (upper triangular with an extra subdiagonal).
- Both matrices are computed by a well-known procedure in linear algebra, known as the <u>Arnoldi procedure</u>.

- The KSA method reduces the problem of calculating the exponential of a large and sparse $K \times K$ matrix \mathbf{P} to the problem of calculating the exponential of the much smaller and dense $K_0 \times K_0$ matrix \mathbf{H} ($K_0 = 30 50$ is usually sufficient for many applications).
- Computation of the reduced-size problem can be done by standard methods.
- The KSA method can be recursively implemented using:

$$\hat{\mathbf{p}}((j+1)\tau) = \exp\{\tau \mathbf{P}\} \hat{\mathbf{p}}(j\tau)$$

$$= ||\hat{\mathbf{p}}(j\tau)||_2 \mathbf{V}(j\tau) \exp\{\tau \mathbf{H}(j\tau)\} \mathbf{e}_1$$

- Practical implementation of the FSP algorithm is difficult.
- The main issue here is the size of the truncated state-space, which is usually very large.
- For this reason, these methods are most often limited to <u>relatively</u> <u>small</u> reaction networks.
- The KSA method is based on several approximations whose cumulative effect may appreciably affect its accuracy, numerical stability and computational efficiency.
- The KSA method <u>does not</u> guarantee that $\hat{\mathbf{p}}(j au)$ is a probability vector !!

The ME

$$\frac{\partial p_Z(\mathbf{z};t)}{\partial t} = \sum_{m=1}^{M} \left\{ a_m(\mathbf{z} - \mathbf{e}_m) p_Z(\mathbf{z} - \mathbf{e}_m;t) - a_m(\mathbf{z}) p_Z(\mathbf{z};t) \right\}, \quad t > 0$$

associated with the DA process can be expressed as a <u>linear</u> system of coupled first-order differential equations:

$$\frac{d\mathbf{q}(t)}{dt} = \mathbf{Q}\mathbf{q}(t), \ t > 0$$

- $\mathbf{q}(t)$ is a $Q \times 1$ vector that contains the nonzero probabilities $p_Z(\mathbf{z};t)$.
- $lackbox{ Q is a large } Q imes Q ext{ sparse matrix whose structure can be inferred from the ME.}$

- \square Ordering the elements in the Z state-space <u>lexicographically</u> results in a matrix \mathbf{Q} that is <u>lower triangular</u>.
- In this case, and for a given time step $\tau > 0$, we can use the <u>implicit</u> <u>Euler method</u> for solving differential equations to estimate $\mathbf{q}(t)$ at discrete time points $j\tau$.
- Given an estimate $\hat{\mathbf{q}}(j\tau)$ of $\mathbf{q}(j\tau)$, we can obtain an estimate $\hat{\mathbf{q}}((j+1)\tau)$ of $\mathbf{q}((j+1)\tau)$ by solving the following system of linear equations: $(\mathbf{I}-\tau\mathbf{Q})\hat{\mathbf{q}}((j+1)\tau)=\hat{\mathbf{q}}(j\tau)$

 $Q \times Q$ identity matrix

https://en.wikipedia.org/wiki/Lexicographic order https://en.wikipedia.org/wiki/Backward Euler method

This is because, from the differential equation

$$\frac{d\mathbf{q}(t)}{dt} = \mathbf{Q}\mathbf{q}(t)$$

we have that

$$\frac{d\mathbf{q}((j+1)\tau)}{dt} = \mathbf{Q}\mathbf{q}((j+1)\tau)$$

$$\Rightarrow \frac{\mathbf{q}((j+1)\tau) - \mathbf{q}(j\tau)}{\tau} \simeq \mathbf{Q}\mathbf{q}((j+1)\tau)$$

$$\Rightarrow (\mathbf{I} - \tau\mathbf{Q})\mathbf{q}((j+1)\tau) \simeq \mathbf{q}(j\tau)$$

- The previous step is possible for <u>any</u> value of τ and can be efficiently done by a standard <u>forward substitution algorithm</u>.
- The resulting method is <u>always</u> stable, producing a <u>valid</u> probability vector at each iteration.
- Its accuracy can be controlled by a <u>single</u> parameter, the step size τ .

https://en.wikipedia.org/wiki/Triangular matrix#Forward and back substitution

- The implicit Euler (IE) method is computationally superior to KSA when the cardinality of the DA state-space is not much larger than the cardinality of the population state-space.
- This is not always possible: the DA process is <u>non-decreasing</u> as opposed to the population numbers that can either increase or decrease.
- The IE method can only be used when the number of reaction events are sufficiently constrained or remain small during a time interval of interest.

Example: In the case of the simple SIR model of epidemiology, the net stoichiometric matrix is given by

$$\mathbf{S} = \begin{bmatrix} -1 & 0 \\ 1 & -1 \\ 0 & 1 \end{bmatrix}$$

and the matrix S^TS is invertible.

 $\ \square$ Consequently, there is a one to one correspondence between the X and Z state-spaces, since

$$\mathbf{X}(t) = \mathbf{x}_0 + \mathbf{S}\mathbf{Z}(t) \Leftrightarrow \mathbf{Z}(t) = (\mathbf{S}^T\mathbf{S})^{-1}\mathbf{S}^T[\mathbf{X}(t) - \mathbf{x}_0]$$

In this case, the IE method is preferable to the KSA method.

- **Example:** Modeling a well-documented 1978 influenza epidemic in an English boarding school.
- We can use the IE method to compute the <u>exact</u> solution of the underlying ME.
- There is a total of 763 students in the school.
- Stochastic SIR model:

$$\begin{array}{c|c}
X_1 \leftrightarrow S \\
X_2 \leftrightarrow I
\end{array}
\qquad
\begin{array}{c|c}
X_1 + X_2 \rightarrow 2X_2 \\
X_2 \rightarrow X_3
\end{array}$$

$$X_3 \leftrightarrow \mathbb{R}$$

$$\begin{array}{c|c} X_1 \longleftrightarrow S & X_1 + X_2 \to 2X_2 \\ X_1 & X_2 & X_3 \end{array}$$

$$\pi_1(x_1, x_2, x_3) = \kappa_1 x_1 x_2$$
$$\pi_2(x_1, x_2, x_3) = \kappa_2 x_2$$

$$\kappa_1 = 0.00218 / \text{day}$$

$$\kappa_2 = 0.44036 \, / \, \text{day}$$

Initial conditions:

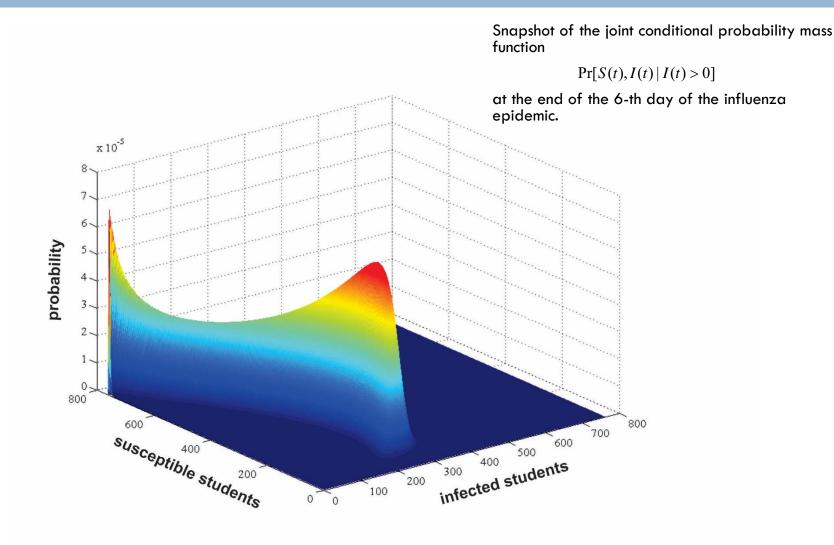
$$X_{1}(0) = 762$$

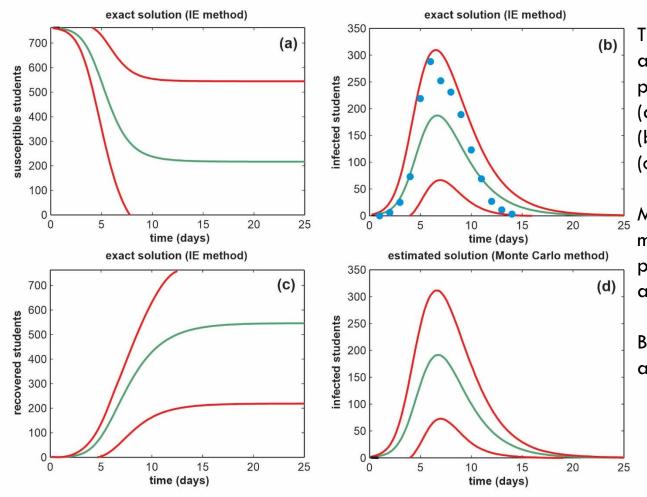
 $X_{2}(0) = 1$
 $X_{3}(0) = 0$
 $X_{1} + X_{2} \to 2X_{2}$
 $X_{2} \to X_{3}$

- The Z state space is a 2-D rectangular grid of points from (0,0) to (762,763).
- \square It contains a total of $763 \times 764 = 582,932$ points.
- \square KSA method \Longrightarrow 4,328 seconds of CPU time.
- □ IE method ⇒ 52 seconds of CPU time.



see video-4-1.mov



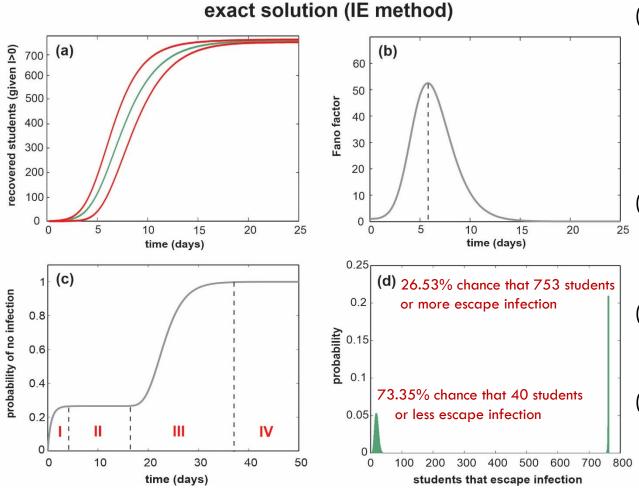


The mean profiles (green lines) and the ± 1 standard deviation profiles (red lines) of

- (a) Susceptible students.
- b) Infected students.
- (c) Recovered students.

Monte Carlo estimates of the mean and standard deviation profiles of the infected students are depicted in (d).

Blue circles in (b) mark available data.



- (a) Evolution of the expected number of recovered students (green line) and the ±1 standard deviations (red lines), given that at least one student is always infected.
 - The Fano factor (variance/mean) associated with the results in (a) as a function of time.
- (c) Dynamic evolution of the probability of no infection Pr[I(t) = 0]
- d) The probability mass function $\Pr[S(\infty) = s, I(\infty) = 0]$ at steady-state.