

Bayesian Computation for Stochastic Kinetic Models

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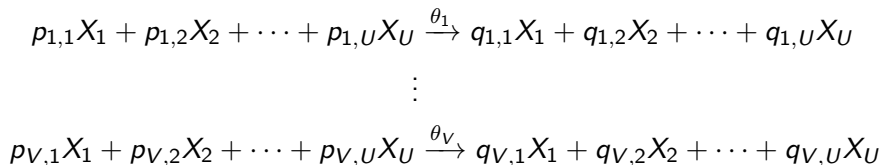
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Definition (Stochastic Kinetic Model)

A Stochastic Kinetic Model (SKM) is a mathematical framework used to describe the time evolution of the species in a system where reactions occur randomly.

Reaction Networks

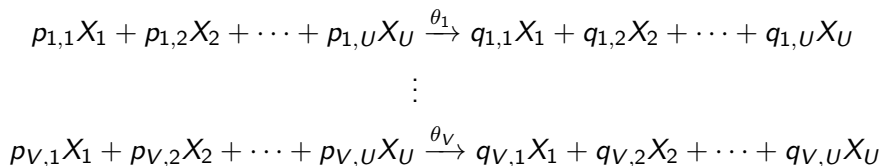
Consider species X_1, \dots, X_U and reaction rates $\theta_1, \dots, \theta_V$. The corresponding reaction network is as follows:



for P and Q ($V \times U$) matrices.

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Definition (Stoichiometric Matrix)

A stoichiometric matrix S can be defined by a $U \times V$ matrix such that $S = (Q - P)$.

Markov Jump Process

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A Markov jump process is a stochastic process that evolves with continuous time in a discrete state space with transitions occurring at random times governed by the Markov property.

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Hazard function

For a reaction i taking place in the time interval $[t, t + dt]$, the probability of this reaction is given by

$$h_i(X_t, \theta_i)dt + \mathcal{O}(dt)$$

where the instantaneous rate/hazard can be defined as

$$h_i(X_t, \theta_i) = \theta_i \prod_{j=1}^U \binom{X_{j,t}}{p_{i,j}}.$$

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Assumptions

- Large number of molecules.
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Methodology

- Number of reactions i in a time step dt is distributed by $Po(h_i(X_t, \theta_i) \cdot dt)$.
- Approximation for large X_t gives $dR_t = h(X_t, \theta) \cdot dt + \text{diag}\{\sqrt{h(X_t, \theta)}\} \cdot dW_t$.
- $dW_t \sim N(0, I \cdot dt)$ for I the identity matrix $d \times d$.
- $dX_t = S \cdot dR_t$.
- $dX_t = S \cdot h(X_t, \theta) \cdot dt + \sqrt{S \text{diag}\{h(X_t, \theta)\} S'} dW_t$.

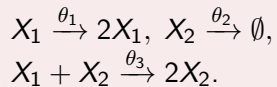
Lotka-Volterra

Let X_1 represent the prey and X_2 represent the predator.

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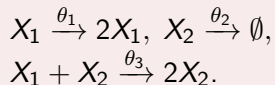
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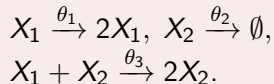
Lotka-Volterra SDE

$$\begin{aligned} \begin{bmatrix} dX_1 \\ dX_2 \end{bmatrix} &= \underbrace{\begin{bmatrix} \theta_1 X_1 - \theta_3 X_1 X_2 \\ \theta_3 X_1 X_2 - \theta_2 X_2 \end{bmatrix}}_{\mu(x)} dt + \\ &\underbrace{\begin{bmatrix} \theta_1 X_1 + \theta_3 X_1 X_2 & \theta_3 X_1 X_2 \\ \theta_3 X_1 X_2 & \theta_3 X_1 X_2 + \theta_2 X_2 \end{bmatrix}}_{\sigma(x)}^{\frac{1}{2}} dW_t \end{aligned}$$

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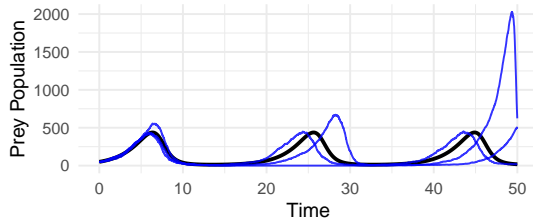
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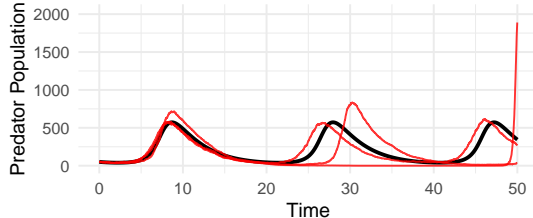
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Prey Population Over Time



Predator Population Over Time



Inference Problem

Aim

Produce parameter estimates for θ given some partially observed data.

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Definition (Euler-Maruyama approximation)

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Discussion

- Lotka-Volterra SDE intractable.
- Euler-Maruyama approximation too crude for large time.
- Use importance sampling and further partitioning of time intervals to overcome the Euler-Maruyama inadequacies.

Posterior

Given data x_0, \dots, x_N , we can find that the posterior takes the form

$$\pi(\theta) \propto \pi_0(\theta) \prod_{t=0}^{T-1} p(x_{t+1}|x_t, \theta).$$

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Transition density

$$p(x_{t+1}|x_t, \theta) \approx \text{Pe}^{(M)}(x_{t+1}|x_t, \theta) = \int \prod_{i=0}^{M-1} \text{Pe}(x_{t_{i+1}}|x_{t_i}, \theta) dx_{t_1:M-1}.$$

Importance Sampling

Durham and Gallant bridge proposal

$$q(x_{t_1}, \dots, x_{t_{M-1}} | x_{t_0}, x_{t_M}) = \prod_{i=0}^{M-2} q(x_{t_{i+1}} | x_{t_i}, x_{t_M}).$$

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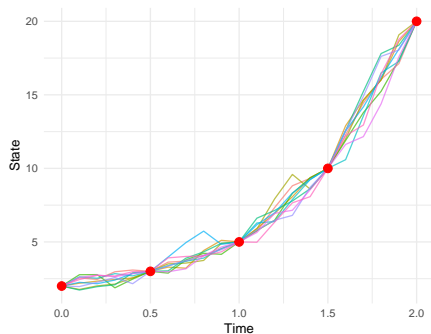


Figure: Durham and Gallant bridges between synthetic observed data points with $M = 10$.

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Estimator

$$\text{Pe}^{(M)}(x_{t+1} | x_t, \theta) = \frac{1}{N} \sum_{j=1}^N \frac{\prod_{i=0}^{M-1} \text{Pe}(x_{t_{i+1}}^{(j)} | x_{t_i}^{(j)}, \theta)}{\prod_{i=0}^{M-2} q(x_{t_{i+1}}^{(j)} | x_{t_i}^{(j)}, x_{t_M})}.$$

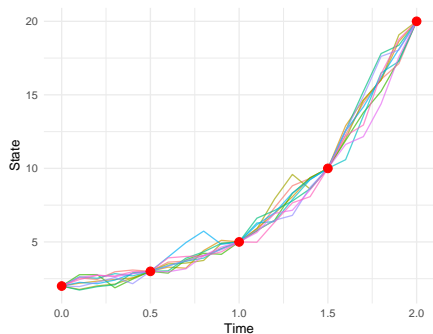


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Pseudo-Marginal Metropolis-Hastings

Outline

- Let u be an auxiliary variable that represents randomness with distribution $g(u)$.
- $\hat{\pi}(\theta, u)$ is the target in the Metropolis-Hastings algorithm.
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Marginalisation

$$\int \hat{\pi}(\theta, u) du \propto \pi_0(\theta) \int \hat{p}(x|\theta) g(u) du \propto \pi_0(\theta) \mathbb{E}_{u \sim g} [\hat{p}(x|\theta)] \propto \pi_0(\theta) p(x|\theta) \propto \pi(\theta).$$

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Definition (PMMH algorithm)

Carry out standard Metropolis Hastings algorithm with target $\hat{\pi}(\theta, u)$ and proposal density

$$q(\theta^*, u^*|\theta, u) = q(\theta^*|\theta)g(u^*).$$

Outline

- All preceding work has been done under the assumption of partially observed data values for both species.
- Now introduce the idea of only observing one species.
- Implement a proposal mechanism for the unobserved species to produce observations to be accepted or rejected along with the rate parameters.

Thank you for listening!