

State and Parameter Estimation from Partial State Observations in Stochastic Reaction Networks

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- Stochastic reaction networks
- Filtering problem: Continuous in time observations.
 - Evolution equations for conditional probabilities.
 - Unnormalized equations and stochastic interpretation.
 - Monte Carlo algorithm: Particle Filter.
 - Numerical examples.
- Filtering problem: Discrete in time observations.

Stochastic reaction networks

n molecular **species** and m **reaction channels**. $Z(t) \in \mathbb{Z}_+^n$: the copy number vector of the species at time t is a CTMC.

- **Stoichiometric vector**: $\nu_j \in \mathbb{Z}^n$, $j = 1, \dots, m$.
- **Propensity functions** $a_j(z)$:

$a_j(z)h + o(h)$ = Probability that reaction j will occur in infinitesimal time interval $(t, t + h]$, given $Z(t) = z$.

Example: $A + B \longrightarrow C$, $C \longrightarrow A + B$, $\emptyset \longrightarrow C$.

$$\nu_1 = \begin{bmatrix} -1 \\ -1 \\ 1 \end{bmatrix}, \quad \nu_2 = \begin{bmatrix} 1 \\ 1 \\ -1 \end{bmatrix}, \quad \nu_3 = \begin{bmatrix} 0 \\ 0 \\ 1 \end{bmatrix},$$

Evolution of the p.m.f.

The time evolution of the probability mass function $p(t, z) = P\{Z(t) = z\}$ is given by the *Kolmogorov's forward equations* also known as the *chemical master equations* (CME):

$$p'(t, z) = \sum_{j=1}^m a_j(z - \nu_j) p(t, z - \nu_j) - \sum_{j=1}^m a_j(z) p(t, z), \quad \forall t > 0, z \in \mathbb{Z}_+^n$$

It is hard to compute when state space is large or infinite. Necessitates Monte Carlo methods.

Stochastic Simulation Algorithm [Gillespie,1977]

- 1: **Input:** Initial state z_0 , terminal time T
- 2: Initialize $z = z_0$, $t = 0$
- 3: **while** $t < T$ **do**
- 4: Set $\lambda_i = a_j(z)$ for $j = 1, \dots, m$, and $\lambda_0 = \sum_{j=1}^m \lambda_j$
- 5: Generate $\tau \sim \text{Exponential}(\lambda_0)$
- 6: Generate $u \sim \text{Uniform}[0, 1]$
- 7: **if** $t + \tau < T$ **then**
- 8: Choose j that j th reaction occurs with probability $\frac{\lambda_j}{\lambda_0}$
- 9: Set $t = t + \tau$, $z = z + \nu_j$
- 10: **else**
- 11: Set $t = T$
- 12: **end if**
- 13: **end while**

Continuous observations: Problem statement

- **Assumption:** Exact (noiseless) observations of the copy number of the last n_2 species in continuous time.
- $Z(t) = (X(t), Y(t))$, where $X(t) \in \mathbb{Z}_+^{n_1}$ is the **unobserved state** and $Y(t) \in \mathbb{Z}_+^{n_2}$ is the **observed state**.
- **Goal:** Estimate the conditional probability distribution

$$\pi(t, x) = P\{X(t) = x \mid Y(s) = y(s), 0 \leq s \leq t\} \quad \forall x \in \mathbb{Z}_+^{n_1}.$$

Observable and unobservable reactions

- $\nu_j = (\nu'_j, \nu''_j)$, where $\nu'_j \in \mathbb{Z}^{n_1}$, $\nu''_j \in \mathbb{Z}^{n_2}$, $j = 1, \dots, m$.

- **Observable reactions:** $\mathcal{O} = \{j : \nu''_j \neq 0\}$.

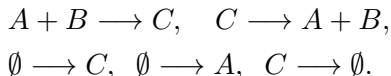
Unobservable reactions: $\mathcal{U} = \{j : \nu''_j = 0\}$.



$$a^{\mathcal{O}}(x, y) = \sum_{j \in \mathcal{O}} a_j(x, y), \quad a^{\mathcal{U}}(x, y) = \sum_{j \in \mathcal{U}} a_j(x, y).$$

Example of observable and unobservable reactions

Example:



- Suppose we observe species A . Then observable reactions $\mathcal{O} = \{1, 2, 4\}$. Unobservable reactions $\mathcal{U} = \{3, 5\}$.
- Suppose we observe species B . Then observable reactions $\mathcal{O} = \{1, 2\}$. Unobservable reactions $\mathcal{U} = \{3, 4, 5\}$.

Evolution of $\pi(t, x)$

It can be shown that $\pi(t, x)$ satisfies **nonlinear** system of ODEs with jumps:

- Let $t_k, k = 1, 2, \dots$ be the successive jump times of $y(t)$.
- On interval $[t_k, t_{k+1})$

$$\begin{aligned} \pi'(t, x) = & \sum_{j \in \mathcal{U}} \pi(t, x - \nu'_j) a_j(x - \nu'_j, y(t_k)) - \sum_{j \in \mathcal{U}} \pi(t, x) a_j(x, y(t_k)) \\ & - \pi(t, x) \left(a^\mathcal{O}(x, y(t_k)) - \sum_{\tilde{x}} a^\mathcal{O}(\tilde{x}, y(t_k)) \pi(t, \tilde{x}) \right) \quad \forall x \in \mathbb{Z}_+^{n_1}. \end{aligned} \quad (1)$$

- At jump times t_k

$$\pi(t_k, x) = \frac{\sum_{l \in \mathcal{O}_k} a_l(x - \nu'_l, y(t_{k-1})) \pi(t_{k-}, x - \nu'_l)}{\sum_{\tilde{x}} \sum_{l \in \mathcal{O}_k} a_l(\tilde{x}, y(t_{k-1})) \pi(t_{k-}, \tilde{x})} \quad \forall x \in \mathbb{Z}_+^{n_1}, \quad (2)$$

where $\mathcal{O}_k = \{j \in \mathcal{O} \mid \nu''_j = y(t_k) - y(t_{k-})\}$.

Issues:

- Hard to compute when state space of X is large or infinite.
- Nonlinear evolution does not correspond to a Markov process.

Unnormalized conditional distribution $\rho(t, x)$

Define $\rho(t, x)$ which satisfies linear ODEs with jumps:

- On interval $[t_k, t_{k+1})$,

$$\begin{aligned} \rho'(t, x) = & \sum_{j \in \mathcal{U}} \rho(t, x - \nu'_j) a_j(x - \nu'_j, y(t_k)) - \sum_{j \in \mathcal{U}} \rho(t, x) a_j(x, y(t_k)) \\ & - \rho(t, x) a^{\mathcal{O}}(x, y(t_k)) \quad \forall x \in \mathbb{Z}_+^{n_1}. \end{aligned} \quad (3)$$

- At jump times t_k ,

$$\rho(t_k, x) = \frac{1}{|\mathcal{O}_k|} \sum_{j \in \mathcal{O}_k} a_j(x - \nu'_j, y(t_{k-1})) \rho(t_k-, x - \nu'_j), \quad x \in \mathbb{Z}_+^{n_1}, \quad (4)$$

where $\mathcal{O}_k = \{j \in \mathcal{O} \mid \nu''_j = y(t_k) - y(t_k-)\}$.

Advantages of $\rho(t, x)$

- $\pi(t, x) = \rho(t, x) / \sum_{\tilde{x}} \rho(t, \tilde{x})$.
 $\rho(t, x)$ is called **unnormalized conditional distribution**.
- We can relate $\rho(t, x)$ to a Markov process.

More on $\pi(t, x)$ and $\rho(t, x)$

- Filtering formula for discrete time Markov chains [*Fristedt, Jain and Krylov, 2007*].
- Rigorous derivation of evolution equation for $\pi(t, x)$ in the finite state CTMC case is in [*Confortola and Fuhrman, 2013*].
- Intuitive derivation for reaction networks based on the discrete time formula was derived in [*Rathinam and Yu, 2021*].
- The unnormalized equations were introduced in [*Rathinam and Yu, 2021*]. Can be derived by setting $\rho(t, x) = g(t) \pi(t, x)$ and looking for a linear evolution equation for ρ .

Monte Carlo Algorithms

We can relate $\rho(t, x)$ to a Markov process $(V(t), w(t))$:

- $V(t)$ is a proxy for $X(t)$,
- $w(t)$ is a weight process,
- $\rho(t, x) = \mathbb{E}[w(t) 1_{\{x\}}(V(t))]$.

In practice, we use an estimator

$$\hat{\rho}(t, x) = \frac{1}{N_s} \sum_{i=1}^{N_s} 1_{\{x\}}(V^{(i)}(t)) w^{(i)}(t), \quad (5)$$

with i.i.d. samples $V^{(i)}(t), w^{(i)}(t), i = 1, \dots, N_s$.

Evolution of $(V(t), w(t))$

In between jump times, $t_k \leq t < t_{k+1}$.

- The process V is evolved according to the underlying reaction network with **all observable reactions removed** (Gillespie Algorithm).
- The weight process $w(t)$ is evolved according to

$$w(t) = w(t_k) \exp \left\{ - \int_{t_k}^t a^{\mathcal{O}}(V(s), y(t_k)) ds \right\}. \quad (6)$$

Evolution at jump times t_k

The process $(V(t), w(t))$ jumps as follows. Randomly pick one $j \in \mathcal{O}_k$ with uniform probability $1/|\mathcal{O}_k|$, and set

$$V(t_k) = V(t_k-) + \nu'_j, \quad (7)$$

$$w(t_k) = w(t_k-) a_j(V(t_k-), y(t_{k-1})), \quad (8)$$

where $\mathcal{O}_k = \{j \in \mathcal{O} \mid \nu''_j = y(t_k) - y(t_{k-1})\}$.

Main Result [Rathinam and Yu, 2021]

The process $(V(t), w(t))$ described by the above procedure satisfies

$$\mathbb{E}[1_{\{x\}}(V(t)) w(t)] = \rho(t, x),$$

for all $t \geq 0$ and x , where $\rho(t, x)$ is the solution to the unnormalized filtering equations (3) and (4).

An analytically tractable example

We consider the simple reaction network



with mass action form of propensities $a_1(z) = c_1 z_2$, $a_2(z) = c_2$, $a_3(z) = c_3 z_2$. Species S is observed exactly while the species A is unobserved. Thus, $Z(t) = (X(t), Y(t)) = (\#A(t), \#S(t))$.

$$\pi(t, x) = \frac{\lambda^x e^{-\lambda}}{x!}$$

where $\lambda = \int_0^t c_1 y(t) dt$.

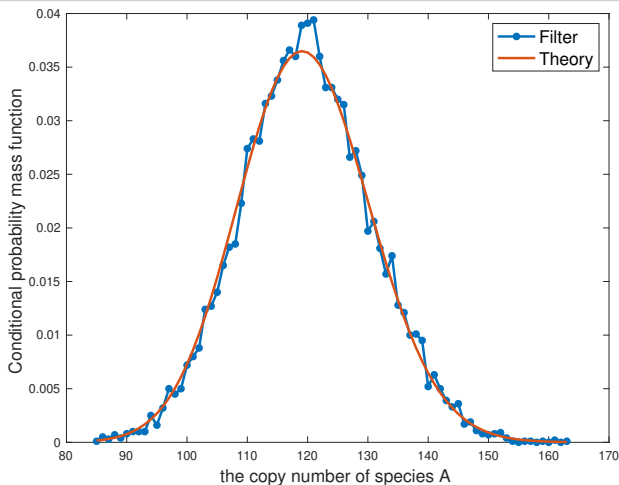


Figure: Estimated conditional probability mass function of the copy number of species A compared with the exact theoretical function at time $T = 20$. $z_0 = (0, 5)$, parameter vector $c = (1, 5, 1)$, filter sample size $N_s = 10,000$.

A point estimate of conditional expectation $\mathbb{E}[X(t) | \mathcal{Y}_t]$

$$\mathcal{E}(t) = \frac{\sum_{i=1}^{N_s} V^{(i)}(t) w^{(i)}(t)}{\sum_{i=1}^{N_s} w^{(i)}(t)}.$$

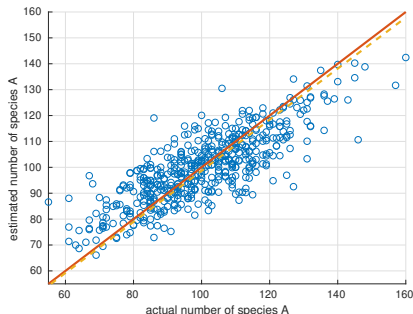
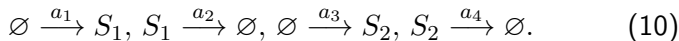


Figure: Scatter plot of the point estimates $\mathcal{E}(T)$ of the copy number of species A (based on the observation of species S) against the actual copy numbers $X(T)$ with $T = 20$. Trial sample size $N_r = 500$, filter particle number $N_s = 1000$.

Genetic toggle switch example



Let $Z(t) = (X(t), Y(t)) = (\#S_1(t), \#S_2(t))$,

with $a_1(z) = \frac{\alpha_1}{1+y^\beta}$, $a_2(z) = x$, $a_3(z) = \frac{\alpha_2}{1+x^\gamma}$, $a_4(z) = y$.

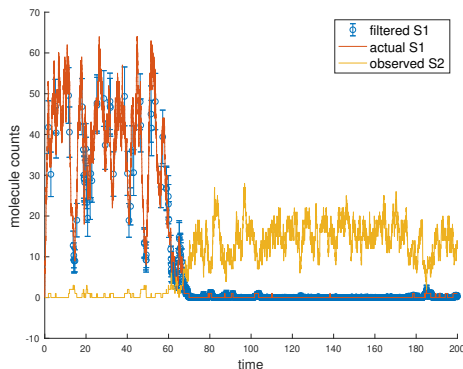


Figure: Estimation of copy number of S_1 at time t based on one observed trajectory of S_2 over $[0, t]$. Estimates are shown at jump times of the observed species S_2 along with their 68% confidence intervals. Filter sample size $N_s = 10,000$. $Z(0) = (0, 0)$, $\alpha_1 = 50$, $\alpha_2 = 16$, $\beta = 2.5$, $\gamma = 1$.

Parameter estimation

Propensity functions $a_j = a_j(x, y, c)$
where $c \in \mathbb{R}^p$.

Bayesian parameter estimation
framework:

- Start with a prior distribution $\bar{\mu}$ on the parameter space
- Treat c as a constant-valued stochastic process $C(t)$
- Apply our algorithm to Markov process $Z(t) = (X(t), C(t), Y(t))$

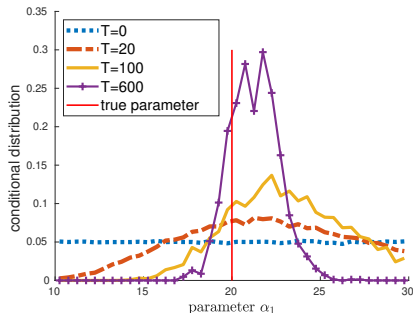


Figure: The posterior conditional probability density function of parameters α_1 and in the genetic toggle example. The nominal parameter values were $\alpha_1 = 20, \alpha_2 = 9, \beta = 2.5, \gamma = 1$.

Discrete time observations: Problem statement

- **Assumption:** Exact (noiseless) observations of the copy number of the last n_2 species at (discrete) snapshots in time.
- $Z(t) = (X(t), Y(t))$, where $X(t) \in \mathbb{Z}_+^{n_1}$ is the **unobserved state** and $Y(t_k) \in \mathbb{Z}_+^{n_2}$ for

$$0 = t_0 < t_1 < t_2 < \dots$$

are the **observed states**.

- **Focus on two observations** at $t_0 = 0$ and $t_1 = T$.
- **Goal:** Estimate the conditional probability distribution

$$\pi(t, z) = P\{Z(t) = z \mid Y(0) = y_0, Y(T) = y\} \quad \forall z \in \mathbb{Z}_+^n.$$

- **Special case:** **Chemical bridge process.** Given end states $Z(0) = z_0$ and $Z(T) = z_T$, generate sample trajectories according to the conditional distribution.

Approach outline: Focus on reaction count vector $R(T)$

- Observations: $Y_0 = y_0$ and $Y(T) = y$. Assume x_0 is known.
 $R(t) = (R_1(t), \dots, R_m(t))$ is the reaction event count vector.

$$X(t) = x_0 + \nu' R(t),$$

$$Y(t) = y_0 + \nu'' R(t).$$

- Change in observed species: $y - y_0 = \nu'' R(T)$.
- Split $R(T) = (R'(T), R''(T))$, where $R'(T)$ is “free” and $R''(T)$ is determined by $R'(T)$.
- Assume that (under simulation probability measure P_0) process R is m -variate Poisson (independent components) with possibly time dependent intensity $\lambda(t, x_0, y_0, y)$.
- **Proxy processes**: I is a proxy for R , $\tilde{Z}(t) = z_0 + \nu I(t)$.
 $\tilde{Z} = (U, V)$ is a proxy for $Z = (X, Y)$.

Main Result

Compute the Girsanov change of measure process $\tilde{L}(t)$:

$$\begin{aligned}\tilde{L}_j(t) = & \left(\prod_{i \geq 1} \frac{a_j(U(\tilde{T}_i^j-), V(\tilde{T}_i^j-))}{\lambda_j(T_i^j-, y_0, y)} 1_{\{\tilde{T}_i^j \leq t\}} \right) \\ & \times \exp \left(\int_0^t (\lambda_j(s, y_0, y) - a_j(U(s), V(s))) ds \right),\end{aligned}$$

where $\tilde{T}_i^j$ for $i = 1, 2, \dots$ are the i th jump time of I_j .

Main Result : Rathinam and Yu

$$P\{Z(t) = z \mid O\} = \frac{\mathbb{E}_0[1_{\{z\}}(\tilde{Z}(t)) \tilde{L}(T) W \mid O]}{\mathbb{E}_0[\tilde{L}(T) W \mid O]}$$

where $O = \{Y_0 = y_0, Y(T) = y\}$ is the observed event.

References

- F. Confortola and M. Fuhrman, *Filtering of continuous-time Markov chains with noise-free observation and applications*, Stochastics, 85:2, 216-251, 2013.
- B. Fristedt, N. Jain and N. Krylov, *Filtering and prediction: A primer*, American Mathematical Society, 2007.
- M. Rathinam and M. Yu, *State and parameter estimation from exact partial state observation in stochastic reaction networks*, Journal of Chemical Physics, 154, 034103, 2021.