

An Introduction to Particle Methods (a.k.a. Sequential Monte Carlo) for Filtering and Smoothing

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- 1 Bayesian Dynamic Models
 - Hidden Markov Models and State-Space Models
 - Extensions
- 2 The Filtering and Smoothing Recursions
- 3 Sequential Importance Sampling
- 4 Sequential Importance Sampling with Resampling

Hidden Markov Model (HMM)

The Hidden State Process $\{X_k\}_{k \geq 0}$ is a Markov chain with initial probability density function (pdf) $t_0(x)$ and transition density function $t(x, x')$ such that*

$$p(x_{0:k}) = t_0(x_0) \prod_{l=0}^{k-1} t(x_l, x_{l+1}) .$$

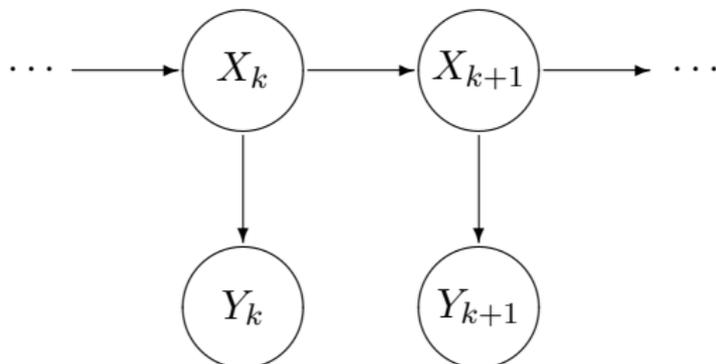
The Observed Process $\{Y_k\}_{k \geq 0}$ is such that the conditional joint density of $y_{0:k}$ given $x_{0:k}$ has the conditional independence (product) form

$$p(y_{0:k} | x_{0:k}) = \prod_{l=0}^k \ell(x_l, y_l) .$$

* $x_{0:k}$ denotes the collection x_0, \dots, x_k .

Graphical Representation of the Dependence Structure

The HMM can be represented pictorially by a **Bayesian network** which depicts the conditional independence relations:



State-Space Form

Here the model is described in a functional form:

$$\begin{aligned}X_{k+1} &= a(X_k, U_k) , \\ Y_k &= b(X_k, V_k) ,\end{aligned}$$

where $\{U_k\}_{k \geq 0}$ and $\{V_k\}_{k \geq 0}$ are mutually independent i.i.d. sequences of random variables (also independent of X_0).

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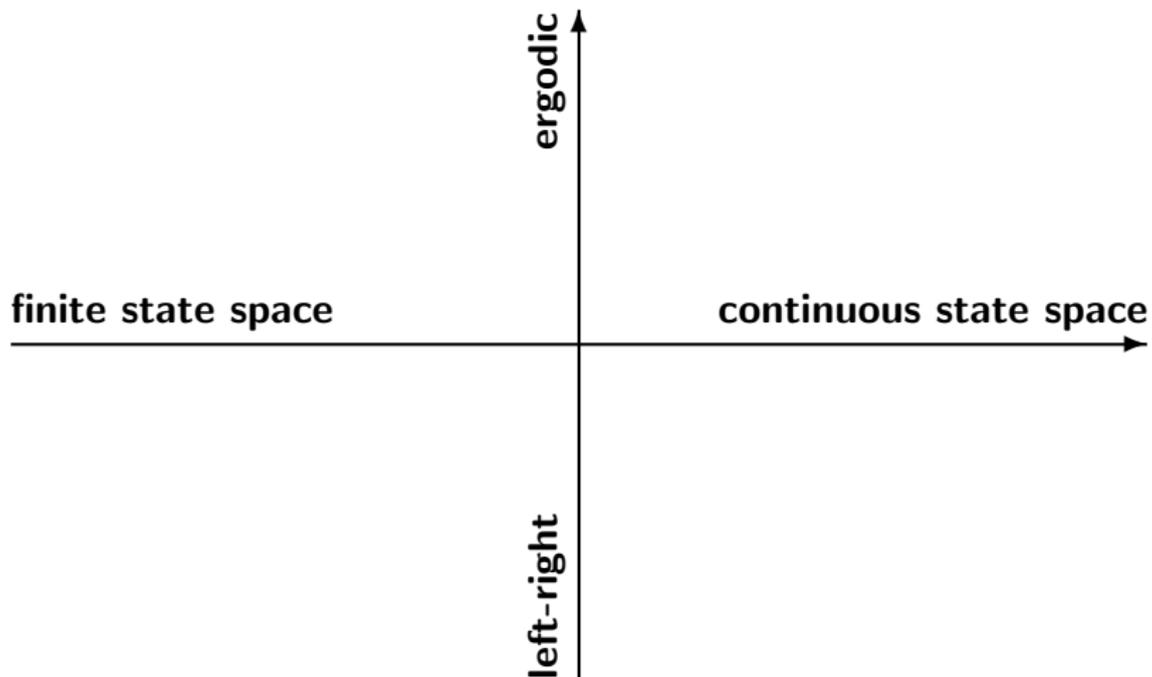
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Remark

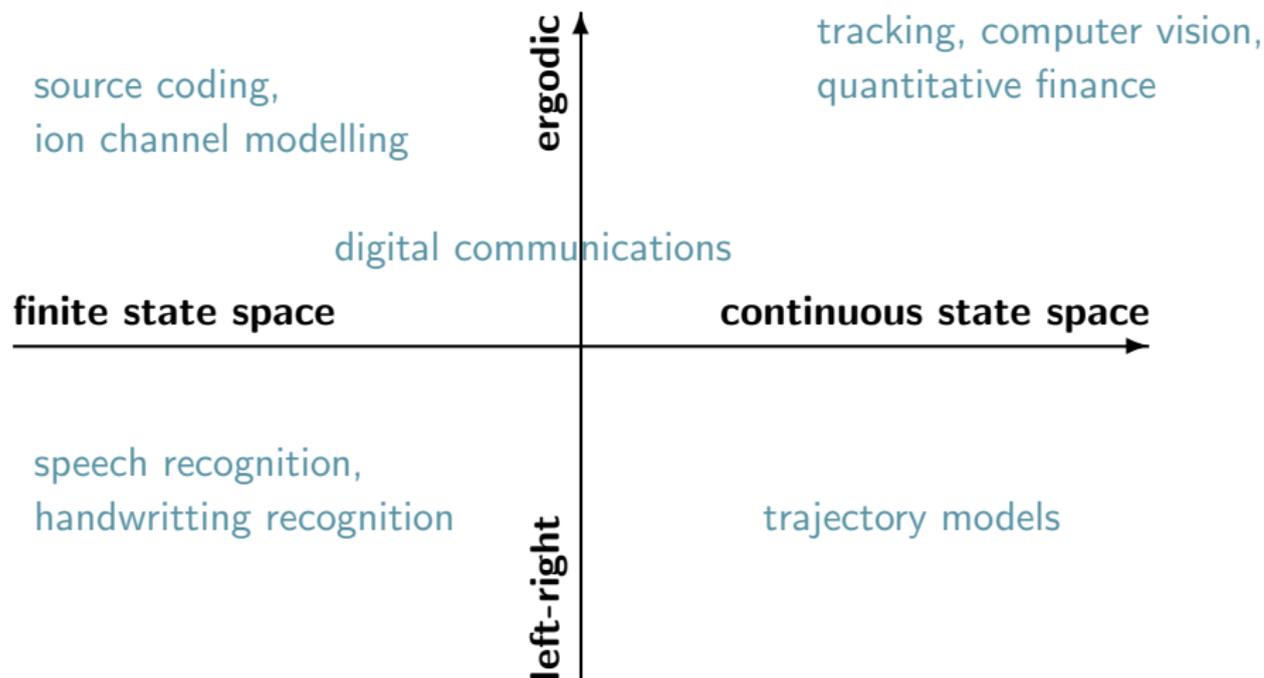
The term *state-space model* often refers to the case where a and b are linear functions of their arguments (and $\{U_k\}$, $\{V_k\}$, X_0 are jointly Gaussian).

Likewise, the term *HMM* is sometimes used (**not in this talk!**) more restrictively for the case where X is a finite set.

HMM Examples



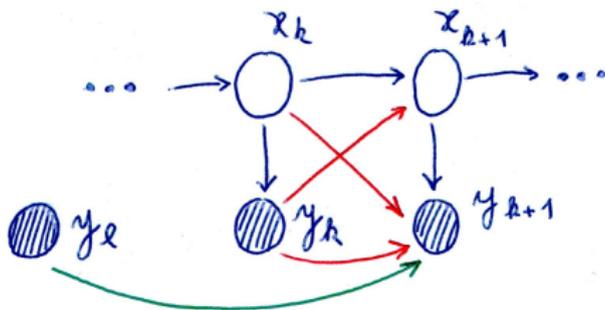
HMM Examples



Beyond HMMs

For sequential Monte Carlo methods, the key point is the **structure of the joint conditional** $p(x_{0:k}|y_{0:k})$. The methods described in this talk directly apply in cases where the joint conditional may be factored as

$$p(x_{0:k}|y_{0:k}) = p(x_0|y_0) \prod_{l=0}^{k-1} p(x_{l+1}|x_l, y_{0:l+1})$$



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 - Basic Recursions
 - Computational Filtering and Smoothing Approaches
- 3 Sequential Importance Sampling
- 4 Sequential Importance Sampling with Resampling

Tasks of interest for HMMs

State Inference How to make probabilistic statements on the state sequence given the model **and the observations?**

Filtering $\pi_{k|k}(x_k) = p(x_k|Y_{0:k})$

Prediction $\pi_{k+1|k}(x_{k+1}) = p(x_{k+1}|Y_{0:k})$

Smoothing $\pi_{0:k|k}(x_{0:k}) = p(x_{0:k}|Y_{0:k})$
(*fixed-interval*: $\pi_{l|k}$ for $l = 0, \dots, k$;
fixed-lag: $\pi_{k|k+\Delta}$ for $k = 0, \dots$)

Parameter Inference How to tune the model parameters based on the observations?

Recursive Structure of the Joint Smoothing Density

By Bayes' rule

$$\begin{aligned}\pi_{0:k+1|k+1}(x_{0:k+1}) &= (\mathbf{L}_{k+1}(Y_{0:k+1}))^{-1} t_0(x_0) \prod_{l=0}^k t(x_l, x_{l+1}) \prod_{l=0}^{k+1} \ell(x_l, Y_l) \\ &= \left(\frac{\mathbf{L}_{k+1}(Y_{0:k+1})}{\mathbf{L}_k(Y_{0:k})} \right)^{-1} \pi_{0:k|k}(x_{0:k}) t(x_k, x_{k+1}) \ell(x_{k+1}, Y_{k+1}),\end{aligned}$$

where the normalization constants \mathbf{L}_k , i.e., the **likelihood** of the observations, is usually not computable.

The Joint Smoothing Recursion

$$\pi_{0:k+1|k+1}(x_{0:k+1}) = \left(\frac{L_{k+1}}{L_k} \right)^{-1} \pi_{0:k|k}(x_{0:k}) t(x_k, x_{k+1}) \ell(x_{k+1}, Y_{k+1})$$

The **marginal recursion** may be decomposed in two steps:

Prediction

$$\pi_{k+1|k}(x_{k+1}) = \int \pi_{k|k}(x_k) t(x_k, x_{k+1}) dx_k$$

Filtering

$$\pi_{k+1|k+1}(x_{k+1}) = \left(\frac{L_{k+1}}{L_k} \right)^{-1} \pi_{k+1|k}(x_{k+1}) \ell(x_{k+1}, Y_{k+1})$$

Exact Implementation of the Filtering and Smoothing Recursions

When X is finite (Baum *et al.*, 1970) The computational cost of filtering is $|X|^2$ per time index.

In linear Gaussian state-space models (Kalman & Bucy, 1961) The filtering and prediction recursion is implemented by the *Kalman filter* (L_{k+1}/L_k is interpreted as the likelihood of the $(k+1)$ -th innovation).

Such *finite dimensional filters* exist only in very specific models (see, e.g., Runggaldier & Spizzichino, 2001).

The Finite Case

Forward (Filtering) – Backward (Smoothing)

Forward For $k = 0$ up to $n - 1$,

$$\pi_{k+1|k+1}(x_{k+1}) = \frac{\ell(x_{k+1}, Y_{k+1}) \sum_{x_k} \pi_{k|k}(x_k) t(x_k, x_{k+1})}{\sum_{x'} \ell(x', Y_{k+1}) \sum_x \pi_{k|k}(x) t(x, x')}$$

Backward For $k = n - 1$ down to 0 ,

$$\pi_{k|n}(x_k) = \sum_{x_{k+1}} b_k(x_k | x_{k+1}) \pi_{k+1|n}(x_{k+1})$$

where

$$\begin{aligned} b_k(x_k | x_{k+1}) &= \frac{\pi_{k|k}(x_k) t(x_k, x_{k+1})}{\sum_x \pi_{k|k}(x) t(x, x_{k+1})} \\ &= P(X_k = x_k | X_{k+1} = x_{k+1}, Y_{0:k}) \end{aligned}$$

Approximate Implementations of the Filtering and Smoothing Recursions

- EKF (Extended Kalman Filter) Linearization-based approach (for non-linear Gaussian state space models);
- UKF (Unscented Kalman Filter, Julier & Uhlmann, 1997) Point-based approach;
- and more Gaussian or Assumed Density Filters (ADF) (Wu, Hu, Xu & Hu, 2006).
- Variational Methods (e.g., Valpola & Karhunen, 2002) Based on parametric density approximation arguments.
- Exact Suboptimal Filters In particular, Kalman filter viewed as minimum mean square error **linear** filtering.

Sequential Monte Carlo (SMC)

- **Sequential Monte Carlo** (sometimes called *particle filtering*) is a method which uses pseudo-random simulations to produce successive populations of “particles” $X_k^{1:n}$ and associated weights $W_k^{1:n}$ such that

$$\sum_{i=1}^n W_k^i f(X_k^i) \approx \int f(x) \pi_{k|k}(x) dx ,$$

for all functions f of interest.

- The SMC process is sequential in the sense that given $X_k^{1:n}$, $W_k^{1:n}$ and the observations $Y_{0:k+1}$, $X_{k+1}^{1:n}$ and $W_{k+1}^{1:n}$ are conditionally independent of previous populations of particles.
- SMC is based on importance sampling and resampling.

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 - Self-Normalized Importance Sampling
 - Sequential Importance Sampling (SIS)
 - Weight Degeneracy
 - SIS: Summary
- 4 Sequential Importance Sampling with Resampling

Self-Normalized Importance Sampling, or IS (Hammersley & Handscomb, 1964)

IS is a weighted form of Monte Carlo approximation, in which expectations under **the target pdf π**

$$\pi(f) = \mathbb{E}_{\pi}[f(X)]$$

are estimated as

$$\hat{\pi}_q^n(f) = \sum_{i=1}^n \frac{\omega^i}{\underbrace{\sum_{j=1}^n \omega^j}_{W^i}} f(X^i) = \frac{\frac{1}{n} \sum_{i=1}^n \omega^i f(X^i)}{\frac{1}{n} \sum_{j=1}^n \omega^j},$$

where

- $X^i \sim \text{iid } q$, where q is an instrumental pdf
- $\omega^i = \frac{\pi}{q}(X^i)$.

This form of IS (sometimes also called Bayesian IS) does not necessitate that π be properly normalized.

Performance of IS

Assuming that $E_{\pi}[\frac{\pi}{q}(X)(1 + f^2(X))] < \infty$, $\hat{\pi}_q^n(f)$ is consistent and asymptotically normal, with **asymptotic variance** given by

$$v_q(f) = E_{\pi} \left[\frac{\pi}{q}(X) (f(X) - \pi(f))^2 \right].$$

The asymptotic variance can be estimated from the IS sample by

$$\hat{v}_q^n(f) = n \sum_{i=1}^n (W^i)^2 \{f(X^i) - \hat{\pi}_q^n(f)\}^2,$$

where $W^i = \omega^i / \sum_{j=1}^n \omega^j$ are the **normalized weights**.

Back to the Filtering and Smoothing Problem

How to estimate expectations under the posterior

$\pi_{0:k|k}(x_{0:k}) = p(x_{0:k}|Y_{0:k})$ in the model

$$p(x_{0:k}) = t_0(x_0) \prod_{l=0}^{k-1} t(x_l, x_{l+1}) ,$$

$$p(y_{0:k}|x_{0:k}) = \prod_{l=0}^k \ell(x_l, y_l) ,$$

using a sequential algorithm ?

Sequential Smoothing through IS, or SIS (Handschin & Mayne, 1969-1970)

- 1 Propose n independent particle trajectories $\{X_{0:k+1}^i\}_{1 \leq i \leq n}$ under a Markovian scheme such that

$$p(x_{0:k+1}) = \rho_{0:k+1}(x_{0:k+1}) = q_0(x_0) \prod_{l=1}^k q_l(x_l, x_{l+1}) .$$

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- 2 Compute importance weights sequentially:

$$\omega_{k+1}^i = \frac{\pi_{0:k+1|k+1}(X_{0:k+1}^i)}{\rho_{0:k+1}(X_{0:k+1}^i)} = \omega_k^i \times \frac{t(X_k^i, X_{k+1}^i) \ell(X_{k+1}^i, Y_{k+1})}{q_k(X_k^i, X_{k+1}^i)}.$$

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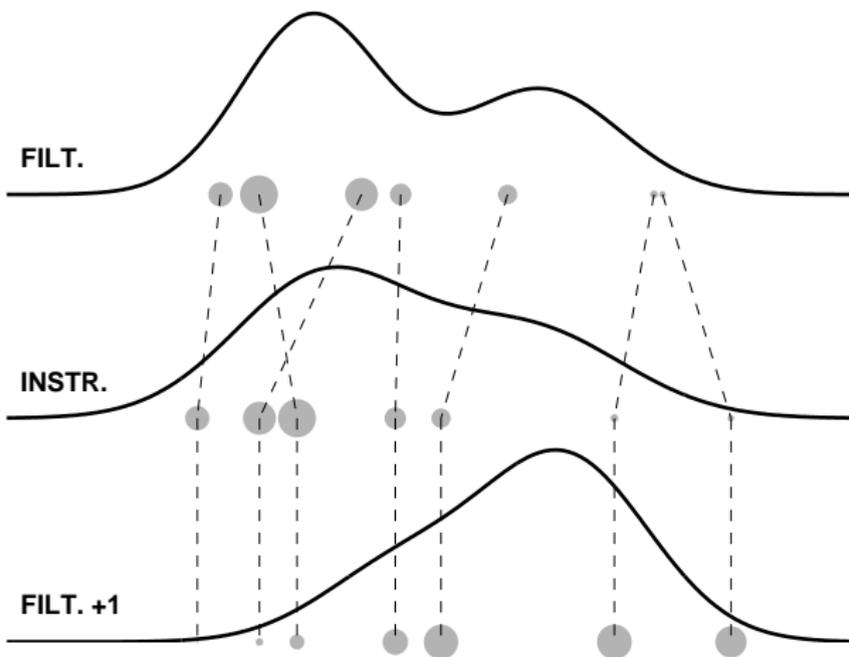
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Then,

$$\sum_{i=1}^n \frac{\omega_{k+1}^i}{\sum_{j=1}^n \omega_{k+1}^j} f(X_{0:k+1}^i)$$

is an estimate of $\mathbb{E}[f(X_{0:k+1}) | Y_{0:k+1}]$.



One step of the SIS algorithm with just seven particles.

Weight Degeneracy

Empirically, the SIS approach always fail when the time-horizon k is more than a few tens; the IS weights $\omega_k^{1:n}$ usually become very unbalanced with a few weights dominating all the other

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To understand why it is the case, consider the (silly) model where

$$\begin{cases} t(x, x') = t(x') = t_0(x'), & \text{(Independent states)} \\ \ell(x, y) = \ell(y), & \text{(Non-informative observations)} \end{cases}$$

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Then

$$\omega_{k+1}^i = \omega_k^i \times \frac{t(X_{k+1}^i)}{q(X_{k+1}^i)}$$

Weight Degeneracy (Contd.)

For a function of interest f that only depends on the last coordinate x_k of the trajectory $x_{0:k}$, the asymptotic variance of the SIS approximation to $\pi_{k|k}(f) = \mathbb{E}_{\pi_{k|k}}[f(X)]$ is given by

$$\begin{aligned}
 v_k(f) &= \\
 &\int \cdots \int \left(\prod_{l=0}^k \frac{t}{q}(x_l) \right)^2 (f(x_k) - \pi_{k|k}(f))^2 \prod_{l=0}^k q(x_l) dx_0 \dots dx_k \\
 &= \underbrace{\left(\int \frac{t}{q}(x) t(x) dx \right)^k}_{>1} \int \frac{t}{q}(x) (f(x) - \pi_{k|k}(f))^2 t(x) dx .
 \end{aligned}$$

In practise, this situation can usually be detected by monitoring the *effective sample size* or *entropy* criteria, which become abnormally small.

Summary

- Sequential Importance Sampling (SIS) is based on simulating independent Markovian trajectories.
- SIS is bound to degenerate in the long-term (depends on everything, including n , but typically between 10 to 100 observations).

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 - Sampling Importance Resampling
 - Sequential Importance Sampling with Resampling (SISR)
 - Marginal and Trajectory-Wise Approximations
 - SISR: Summary

In IS, it is indeed possible to **reset the weights to a constant value** at the price of a, usually moderate, increase in variance.

Sampling Importance Resampling (Rubin, 1987)

Replace $\{X^{1:n}, W^{1:n}\}$ by $\{\tilde{X}^{1:\tilde{N}}, \tilde{W}^{1:\tilde{N}}\}$ such that the discrepancy between the resampled weights $\{\tilde{W}^{1:\tilde{N}}\}$ is reduced and $\sum_{i=1}^{\tilde{N}} \tilde{W}^i \delta_{\tilde{X}^i}$ is a good approximation to $\sum_{i=1}^n W^i \delta_{X^i}$.

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In general the resampling is random and subject to the constraints

$$\begin{cases} \tilde{N} = n, \\ \tilde{W}^i = 1/\tilde{N}, \\ \mathbb{E} \left[\sum_{i=1}^{\tilde{N}} \mathbb{1}\{\tilde{X}^i = X^j\} \mid X^{1:n}, W^{1:n} \right] = \tilde{N} W^j \quad (1 \leq j \leq n). \end{cases}$$

The last condition is often referred to as *unbiasedness* or *proper weighting*.

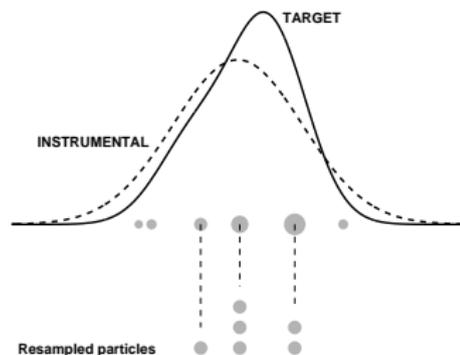
Multinomial Resampling

- 1 Draw, conditionally independently given $\{X^{1:n}, W^{1:n}\}$, n discrete random variables (J^1, \dots, J^n) taking their values in the set $\{1, \dots, n\}$ with probabilities (W^1, \dots, W^n) .
- 2 Set, for $i = 1, \dots, n$, $\tilde{X}^i = X^{J^i}$ and $\tilde{W}^i = 1/n$.

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Let $C^i = \sum_{j=1}^n \mathbb{1}\{\tilde{X}^j = X^i\}$ ($i = 1, \dots, n$) denote the number of times each particle is duplicated in the resampling process. The counts (C^1, \dots, C^n) follow a multinomial distribution with parameters n , (W^1, \dots, W^n) , conditionally to $\{X^{1:n}, W^{1:n}\}$.



Some Results on SIR

- 1 $\tilde{X}^i \xrightarrow{\mathcal{D}} \pi$ as $n \rightarrow \infty$ (some extensions of this result)
- 2 $\frac{1}{n} \sum_{i=1}^n f(\tilde{X}^i)$ is an asymptotically normal estimator of $\pi(f)$ (assuming $\mathbb{E}_\pi[\frac{\pi}{q}(X)(1 + f^2(X)) + f^2(X)] < \infty$) with asymptotic variance given by

$$\tilde{v}_q(f) = \underbrace{\mathbb{E}_\pi \left[\frac{\pi}{q}(X) (f(X) - \pi(f))^2 \right]}_{v_q(f)} + \underbrace{\mathbb{E}_\pi \left[(f(X) - \pi(f))^2 \right]}_{\text{Var}_\pi[f(X)]}$$

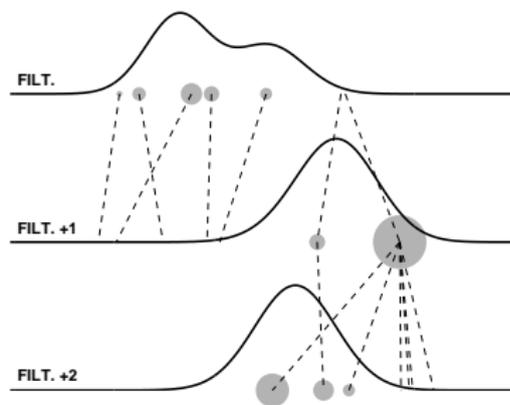
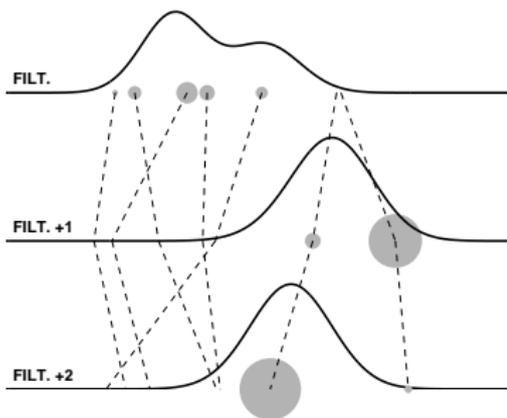
If n is sufficiently large, the cost of resampling is very moderate in situation that are challenging for IS, i.e., when $v_q(f) \gg \text{Var}_\pi[f(X)]$.

The Simplest Functional Algorithm (Gordon et al., 1993)

Regular resampling is added to avoid **weight degeneracy** and to guarantee the long-term ($k \rightarrow \infty$) stability of the particle filter.

The Bootstrap filter

- 1 Given $\tilde{X}_k^{1:n}$, propose new positions X_{k+1}^i independently **under the prior dynamic** $t(\tilde{X}_k^i, \cdot)$, for $i = 1, \dots, n$;
- 2 Compute the **weights** $\omega_{k+1}^i = \ell(X_{k+1}^i, Y_{k+1})$, for $i = 1, \dots, n$ and normalize them ($W_{k+1}^i = \omega_{k+1}^i / \sum_{j=1}^n \omega_{k+1}^j$);
- 3 **Resample** to obtain $\tilde{X}_{k+1}^{1:n}$, e.g., by drawing independent indices J_{k+1}^i such that $P(J_{k+1}^i = j | W_{k+1}^{1:n}) = W_{k+1}^j$ and setting $\tilde{X}_{k+1}^i = X_{k+1}^{J_{k+1}^i}$ (Multinomial Resampling).



SIS (left) and SISR (right).

Marginal and Trajectory-Wise Approximations

SMC is expected to approximate the filtering pdfs in the sense that

$$\sum_{i=1}^n W_k^i f(X_k^i) \longrightarrow \int f(x) \pi_{k|k}(x) dx ,$$

as n increases, for arbitrary functions f .

But recalling our original SIS interpretation, one should also have

$$\sum_{i=1}^n W_k^i f(X_{0:k}^i) \longrightarrow \int \cdots \int f(x_{0:k}) \pi_{0:k|k}(x_{0:k}) dx_0 \cdots dx_k .$$

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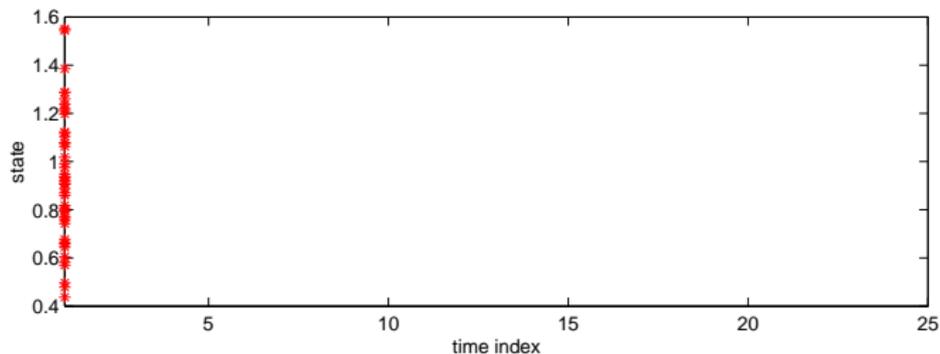
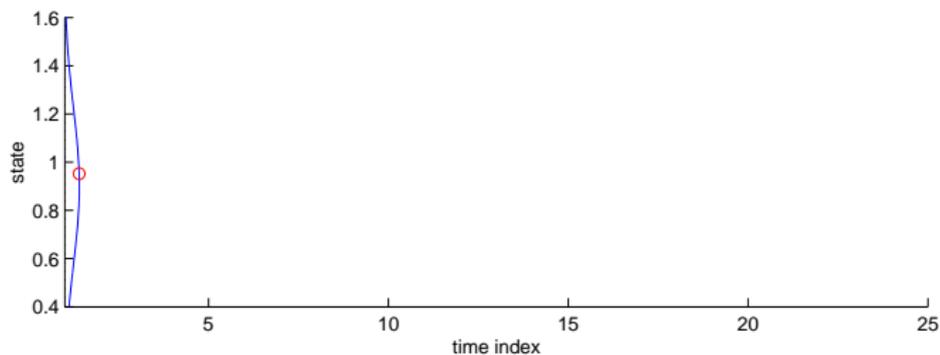
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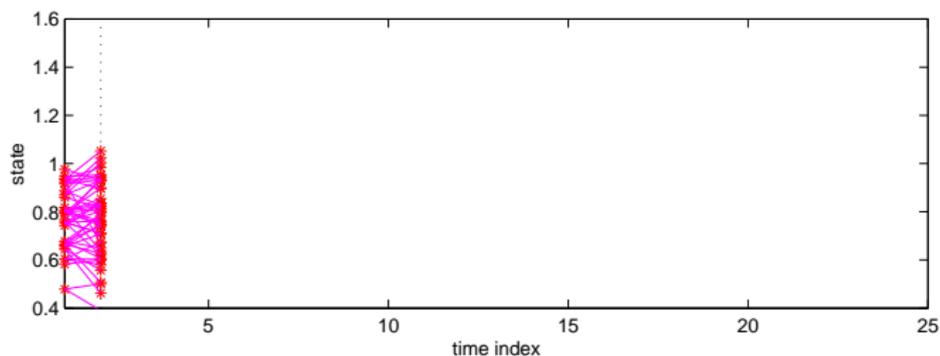
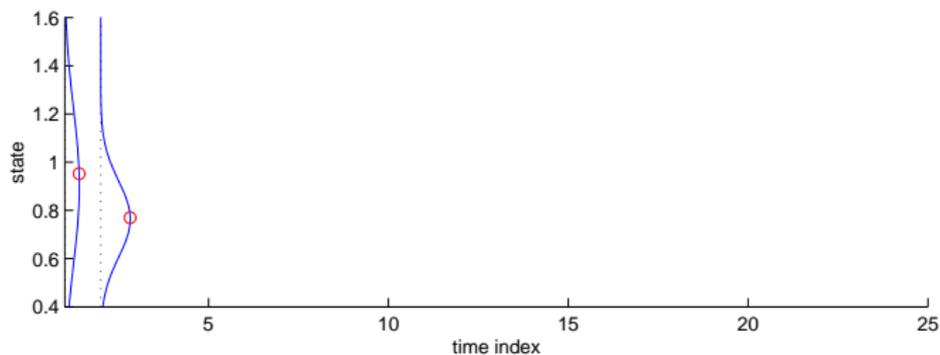
- 1 In what sense is this true? [Several: Consistency, central limit theorem, L^p bounds, convergence in distribution of subpopulations (“propagation of chaos”)]
- 2 What is the influence of n ? [Easy: $1/\sqrt{n}$]
- 3 What is the influence of k ? [Harder: depends on forgetting properties of the model and whether one considers marginal or trajectory-wise approximations]

The Particle Paths



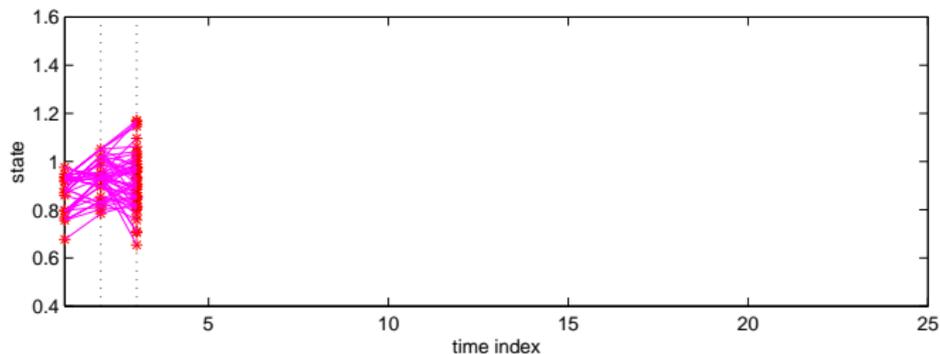
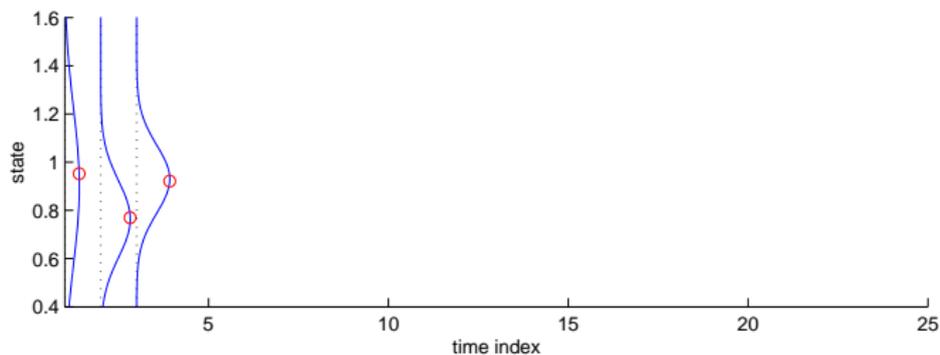
Predictive densities and evolution of the particle ancestry tree

The Particle Paths



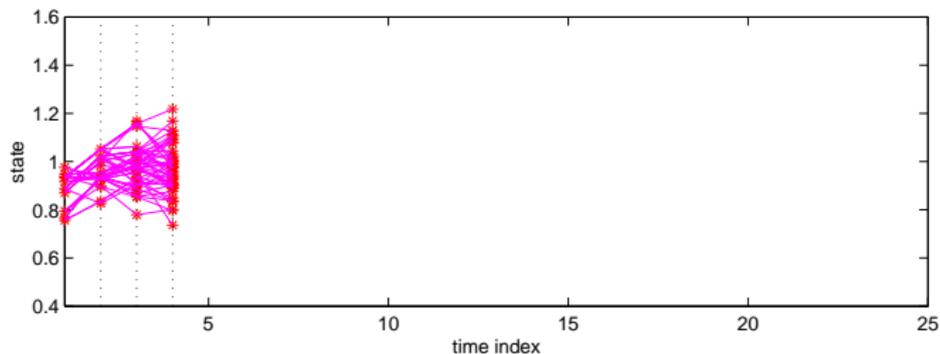
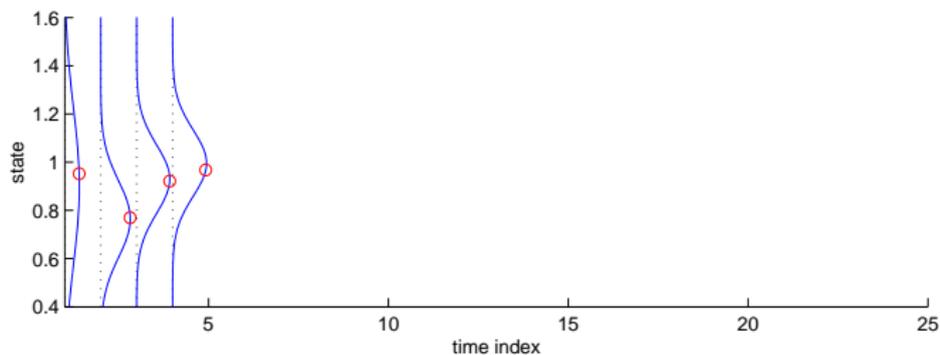
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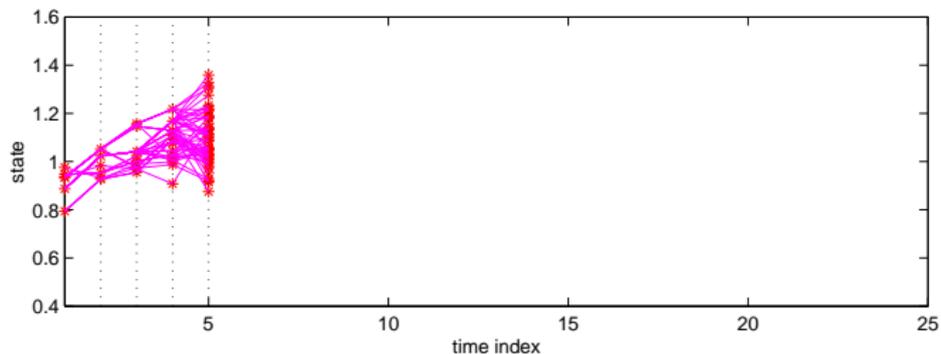
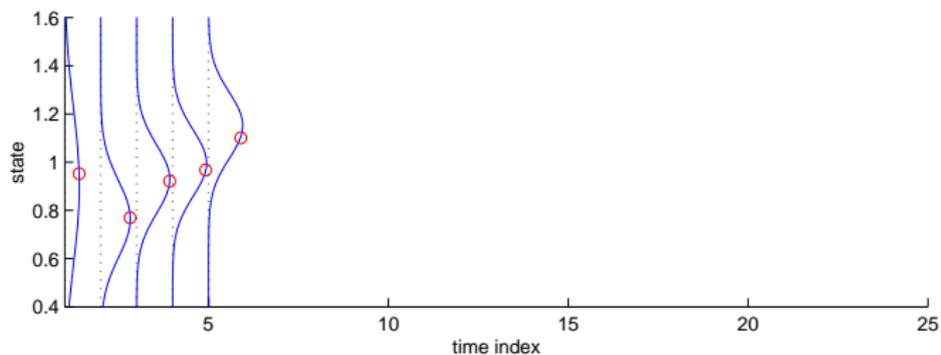
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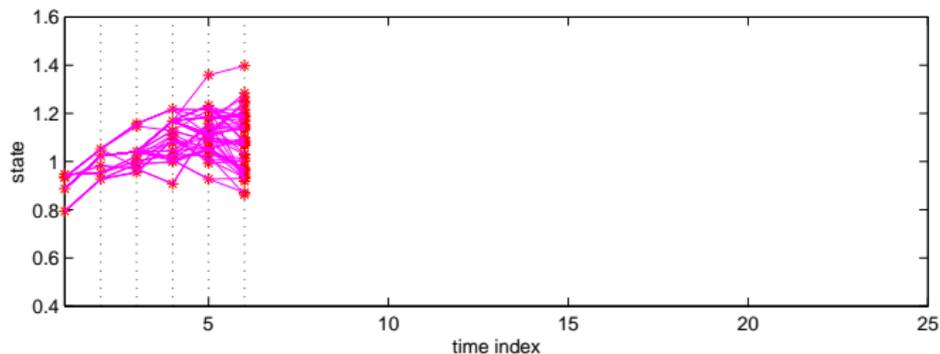
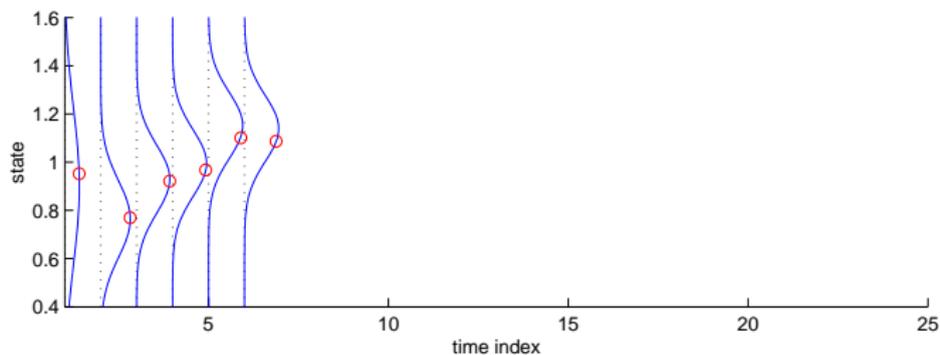
Predictive densities and evolution of the particle ancestry tree

The Particle Paths



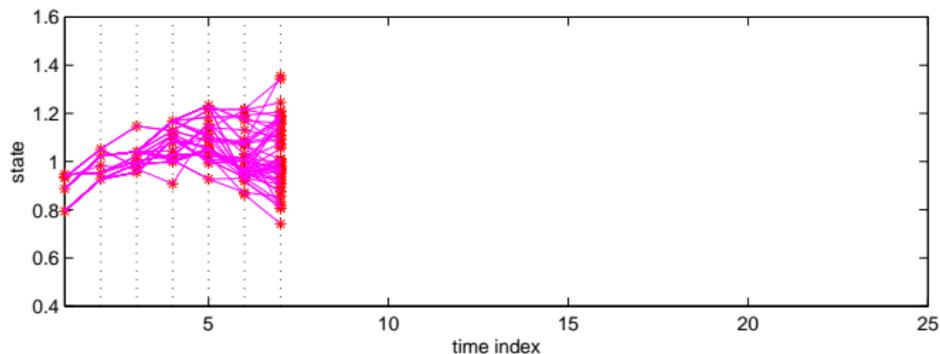
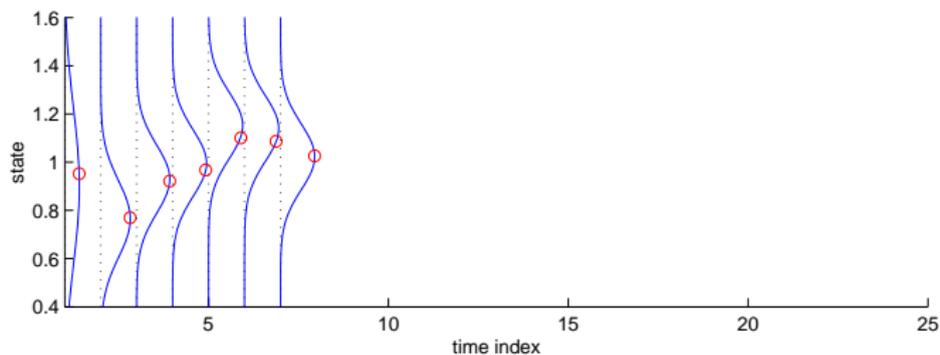
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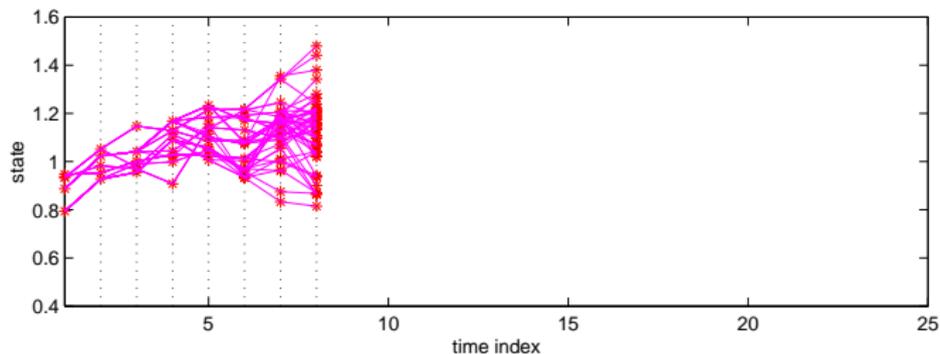
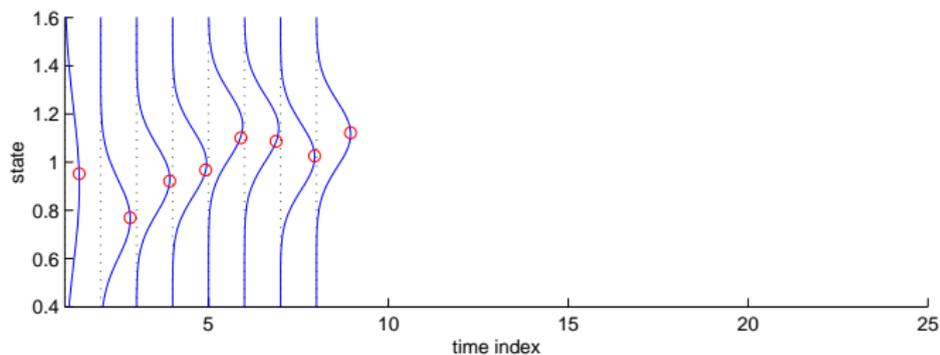
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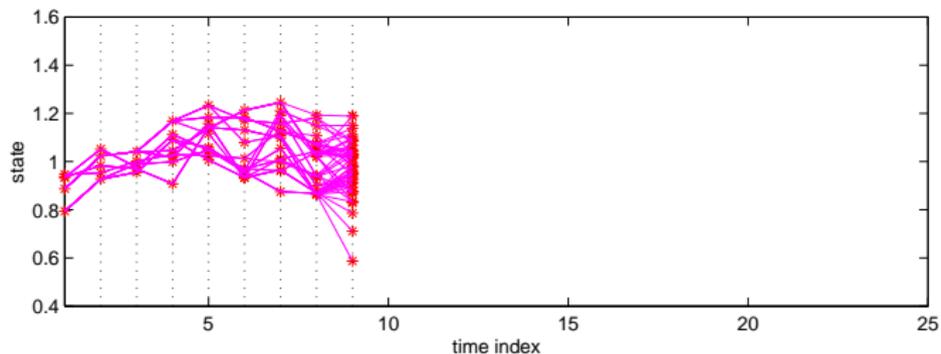
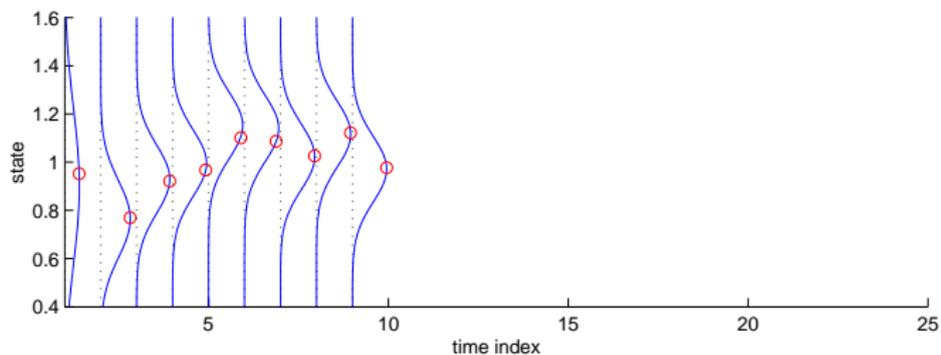
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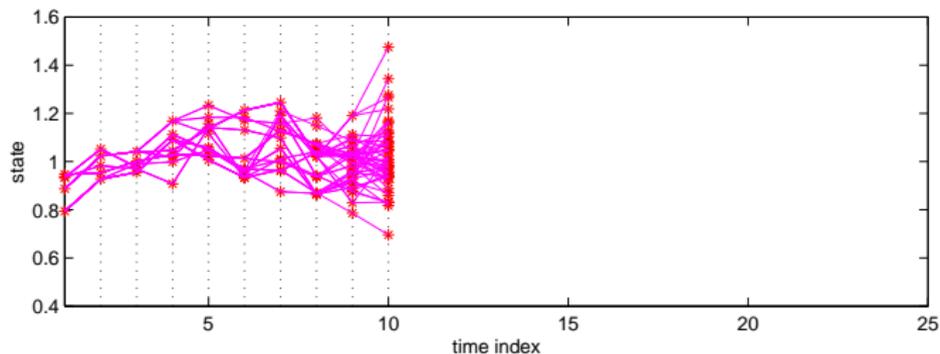
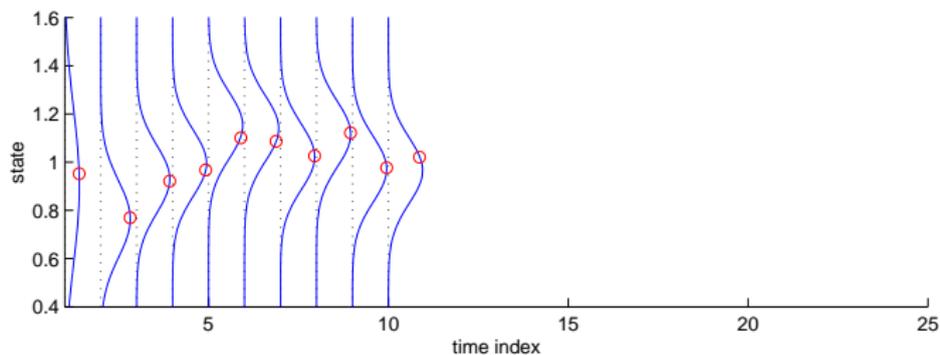
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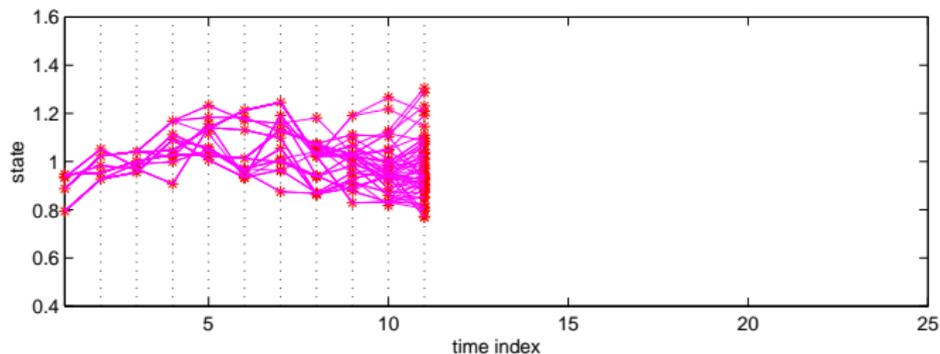
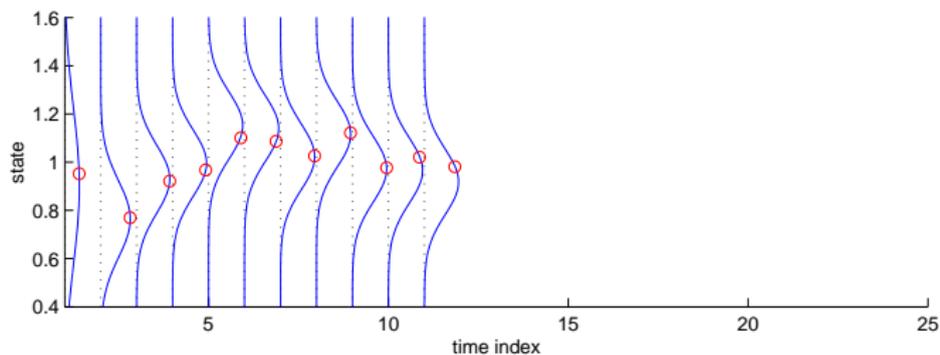
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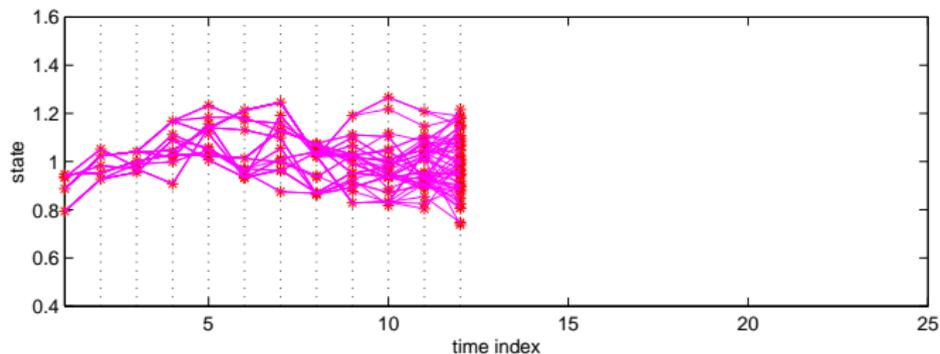
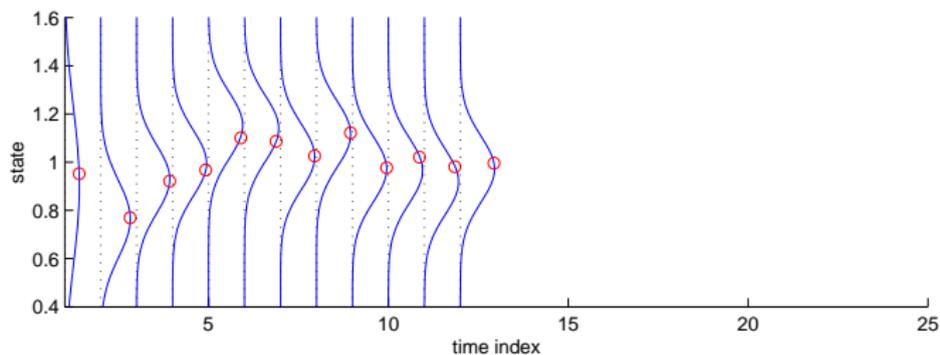
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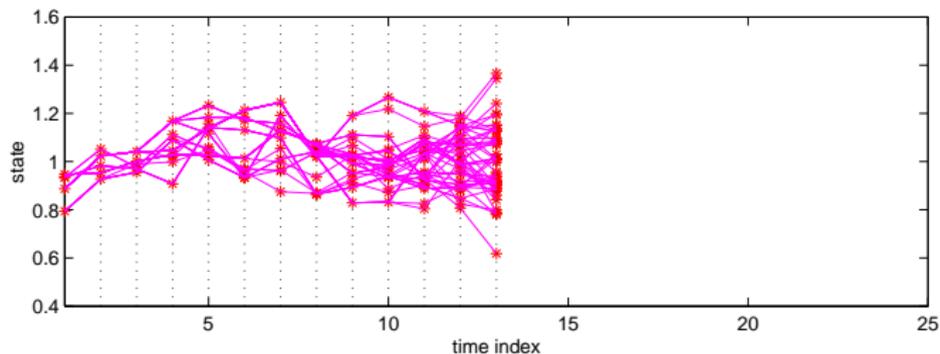
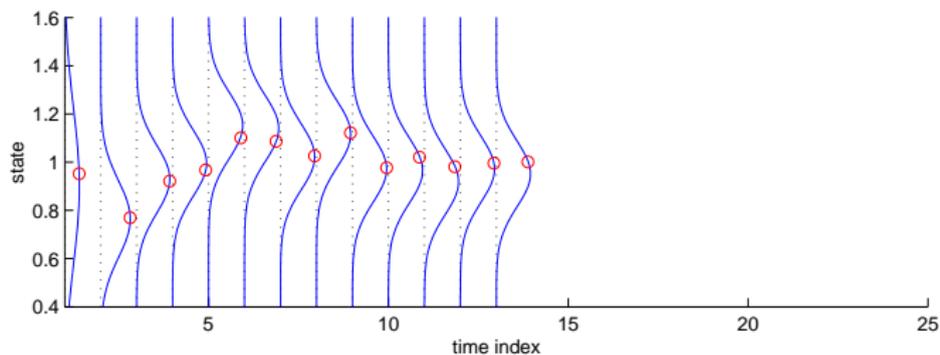
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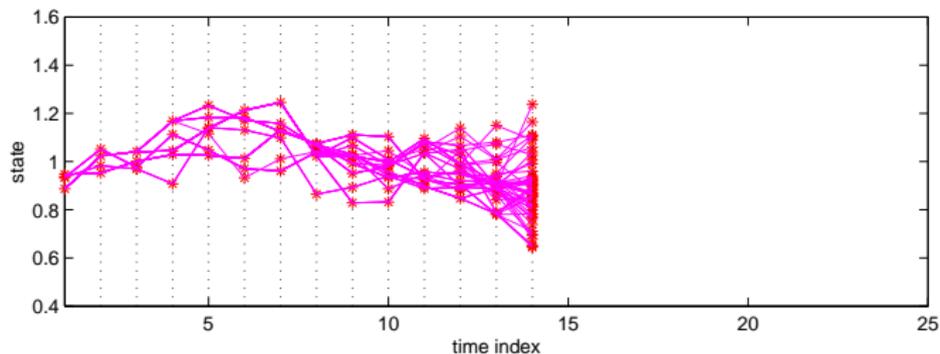
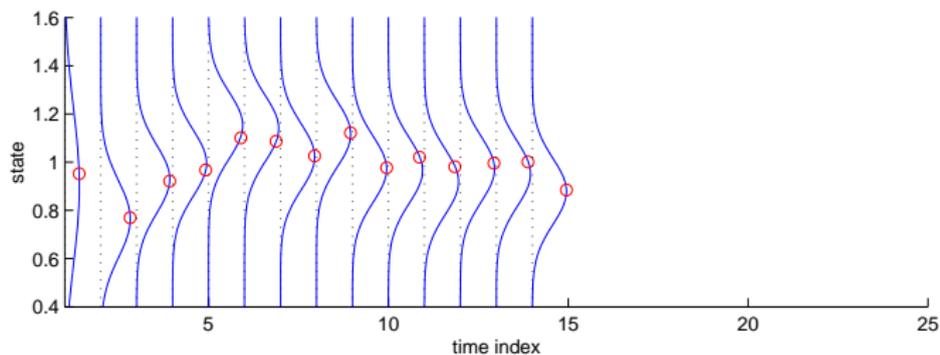
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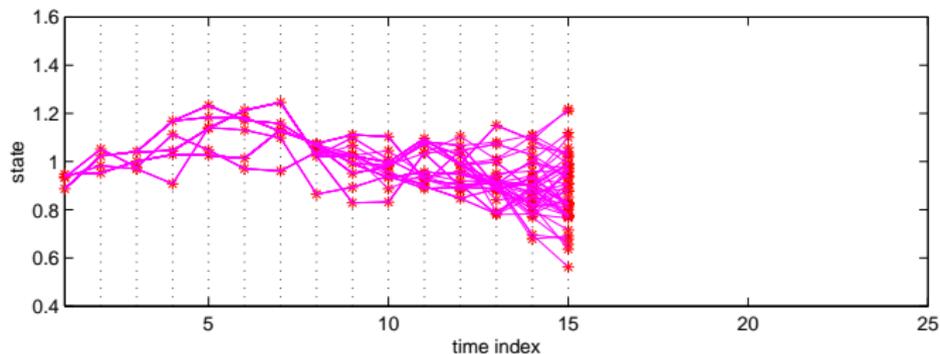
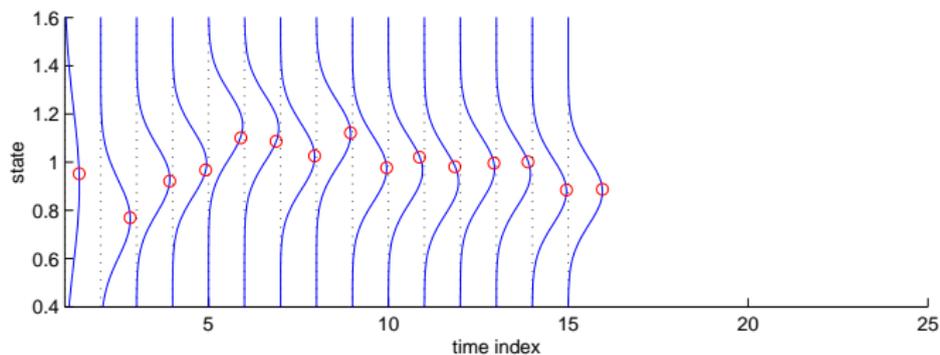
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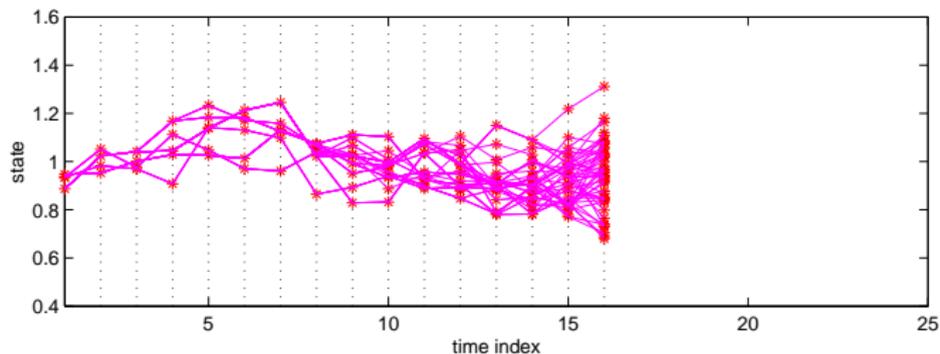
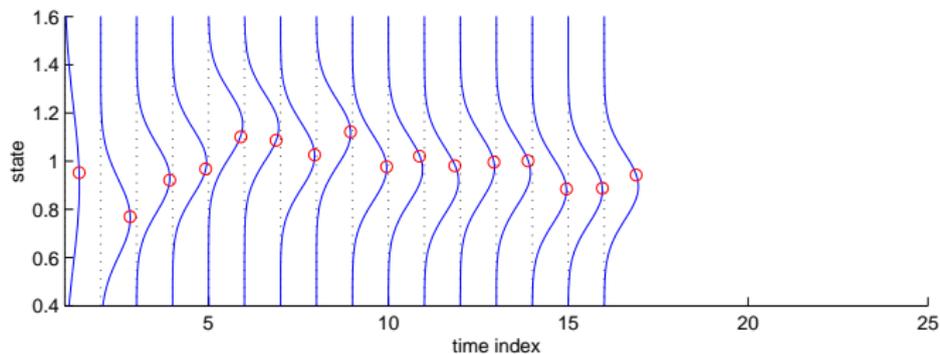
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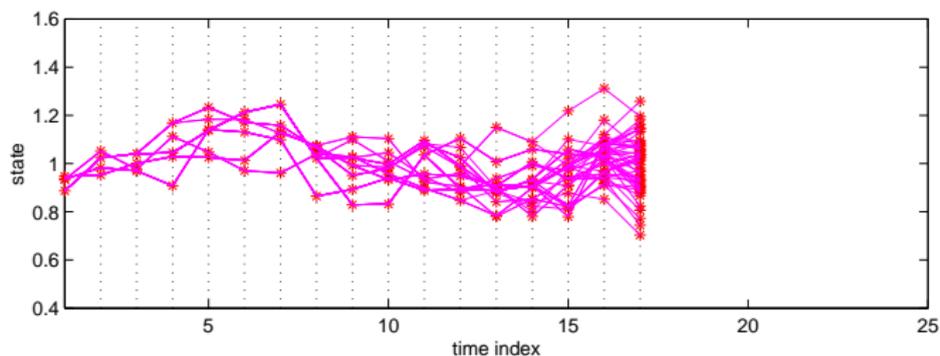
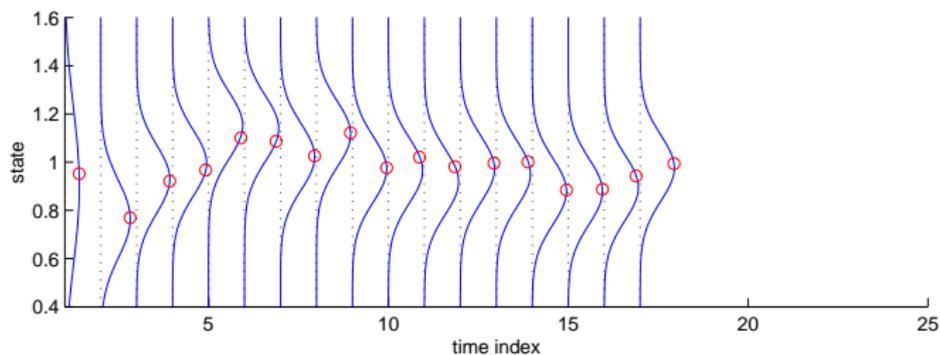
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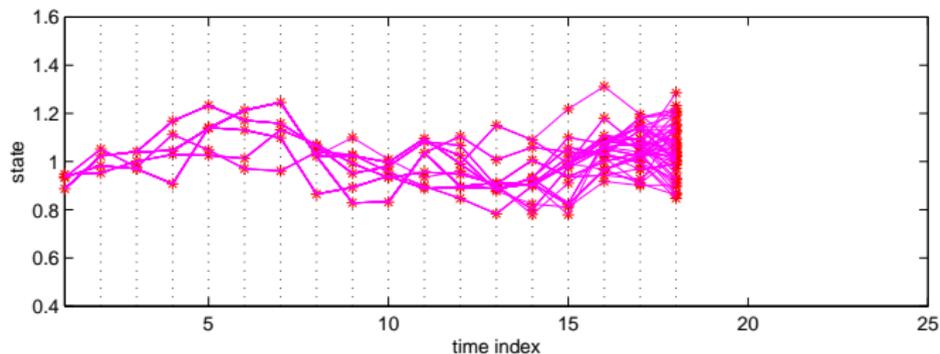
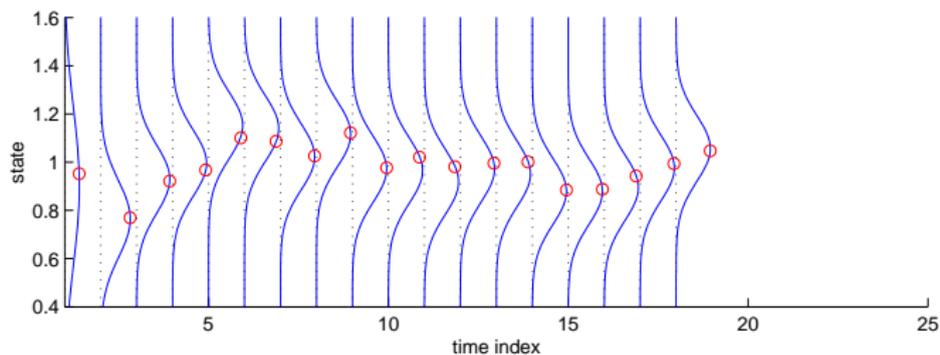
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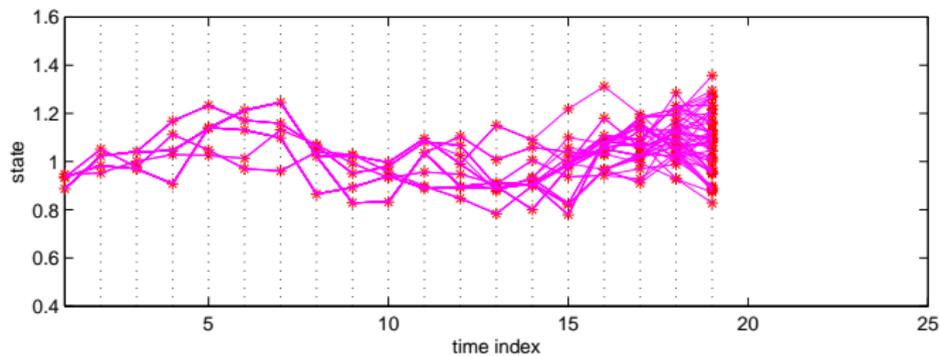
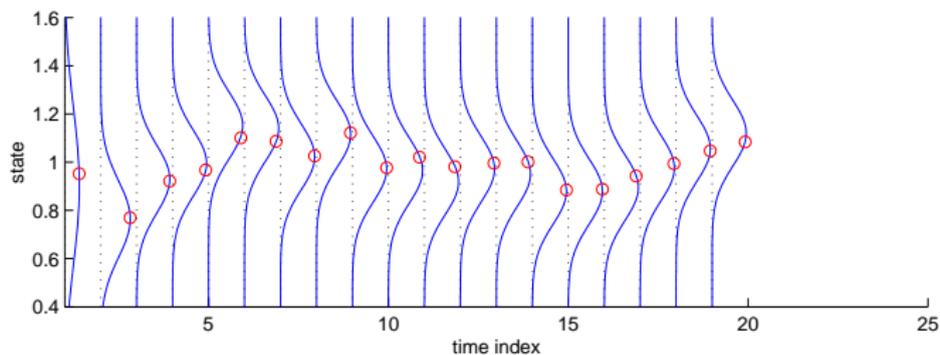
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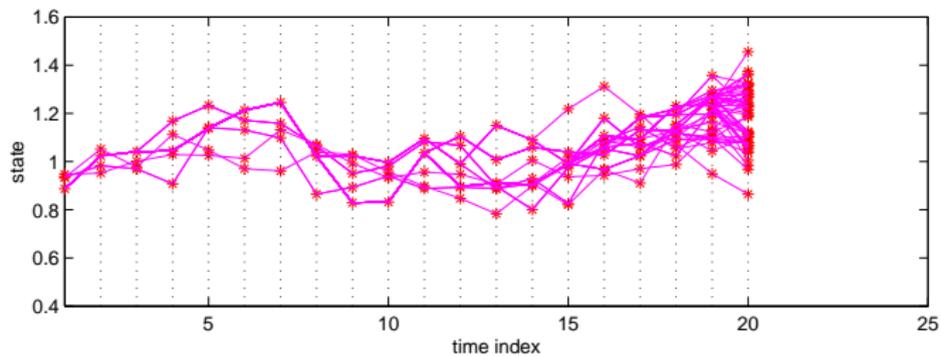
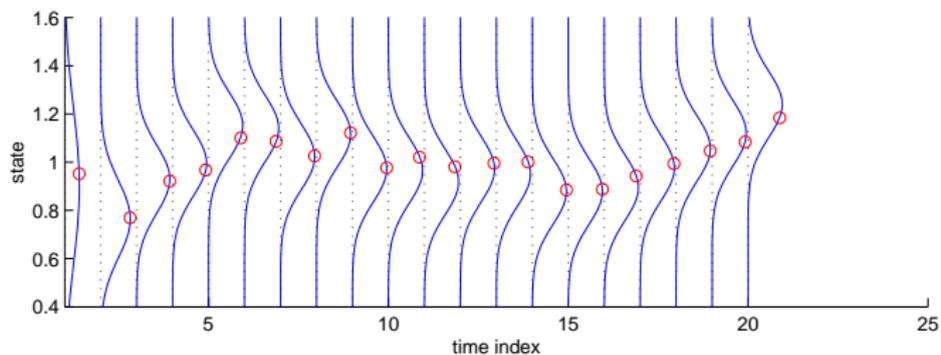
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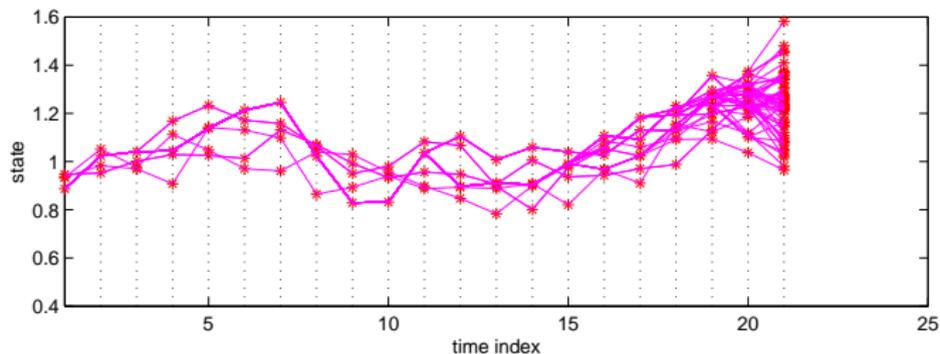
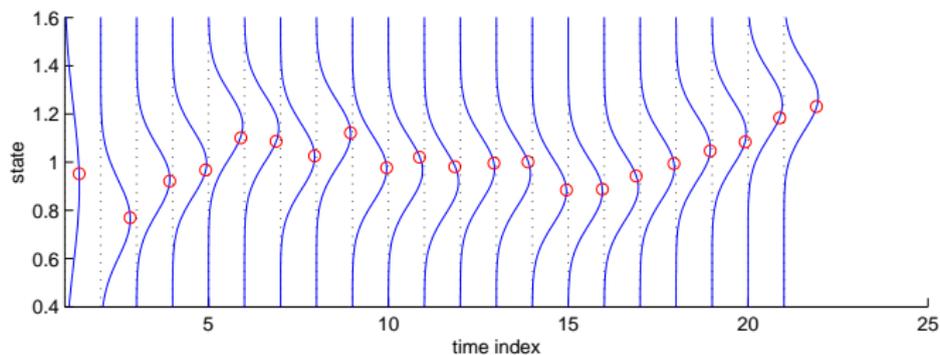
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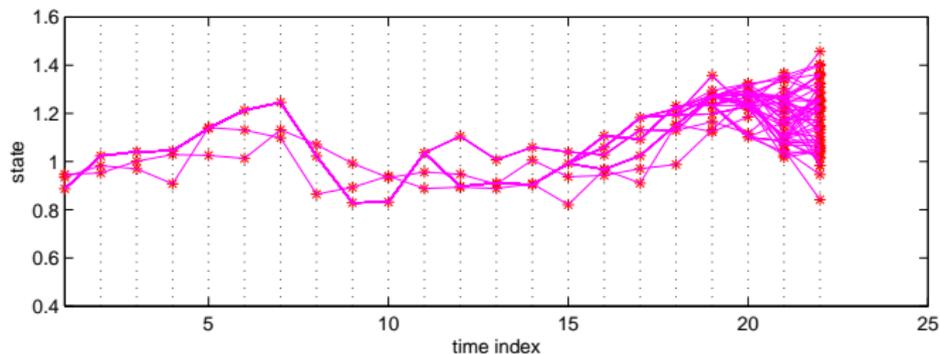
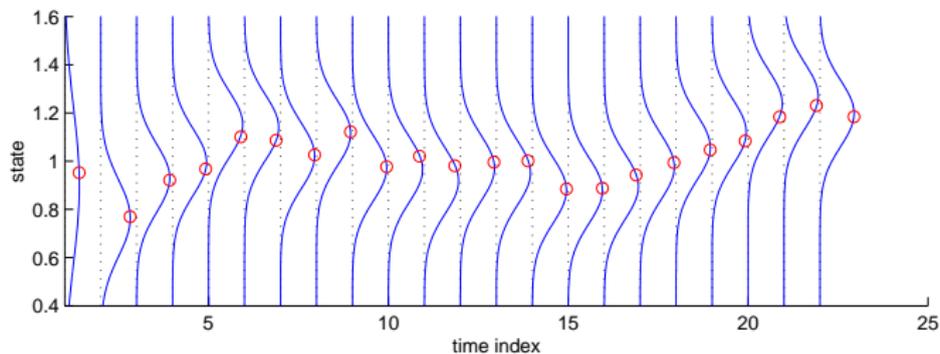
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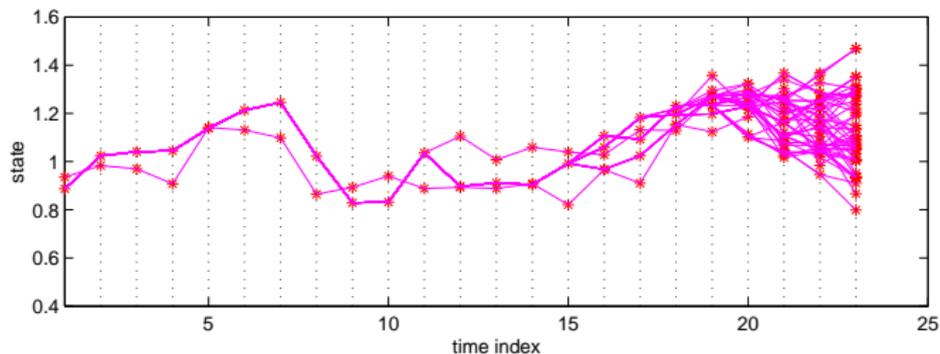
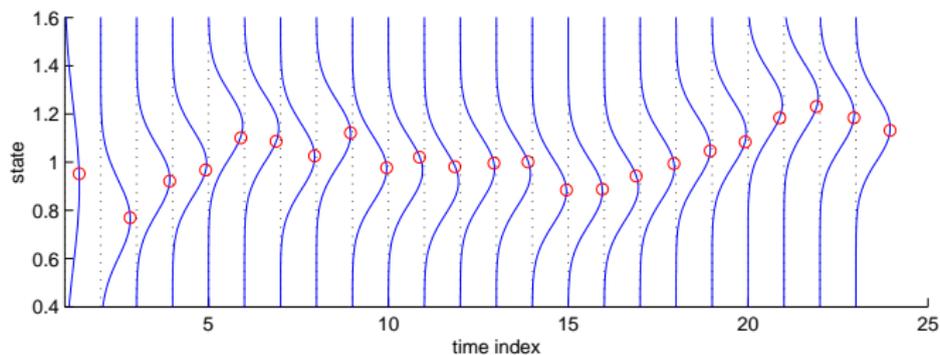
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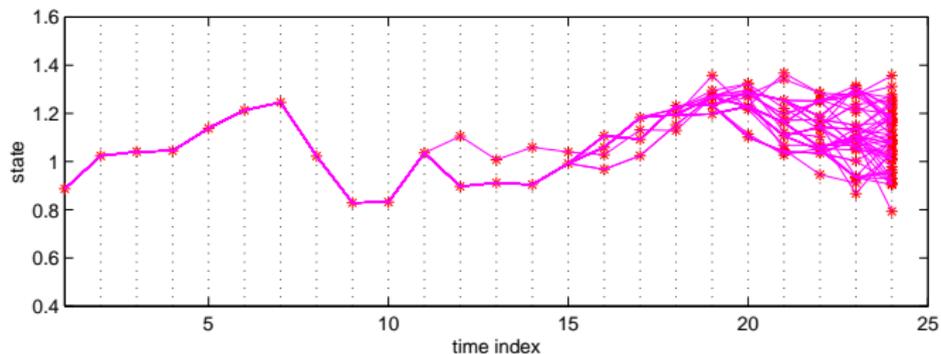
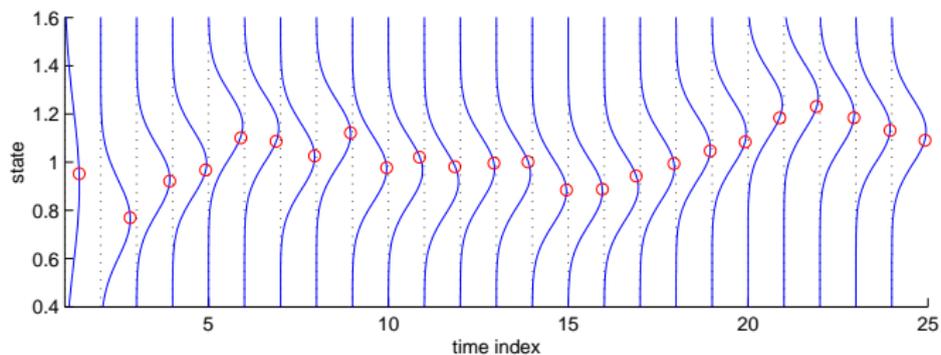
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Summary

- With **resampling**, SISR can achieve **long-term stability**.
- The increase in variance due to resampling is moderate, especially when resampling is applied only when needed.
- The method is still sensitive to outliers, model misspecification, etc., which may necessitate the use of more elaborate strategies (clever choices of the instrumental kernel, adaptive strategies, etc.)
- Accurate smoothing approximations require more elaborate techniques

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