

On the concentration of interacting particle processes

Part I : Introduction and survey

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Sino-French Summer Institute in Stochastic Modeling and
Applications, Beijing, June 2011

Introduction

- Stochastic particle sampling methods
- Interacting jumps models
- Particle Feynman-Kac models

Feynman-Kac models

- Some basic notation
- Description of the models
- The 3 keys formulae

Interacting particle systems

- A bad tempting idea
- Adaptive interacting jumps
- The 4 particle estimates
- Island particle models

Concentration inequalities

- Current population models
- Particle free energy
- Genealogical tree models
- Backward particle models

Some ideas on how it works

Introduction

Stochastic particle sampling methods

Interacting jumps models

Particle Feynman-Kac models

Feynman-Kac models

Interacting particle systems

Concentration inequalities

Some ideas on how it works

Introduction

Stochastic particle methods
=
Universal adaptive sampling technique

2 types of stochastic interacting particle models:

- ▶ Diffusive particle models with mean field drifts
[McKean-Vlasov style]
- ▶ Interacting jump particle models
[Boltzmann & Feynman-Kac style]

Lectures ⊂ Interacting jumps models

- ▶ Interacting jumps = Recycling transitions =
- ▶ Discrete generation models (\Leftrightarrow geometric jump times)



Equivalent particle algorithms

Sequential Monte Carlo	Sampling	Resampling
Particle Filters	Prediction	Updating
Genetic Algorithms	Mutation	Selection
Evolutionary Population	Exploration	Branching-selection
Diffusion Monte Carlo	Free evolutions	Absorption
Quantum Monte Carlo	Walkers motions	Reconfiguration
Sampling Algorithms	Transition proposals	Accept-reject-recycle

Equivalent particle algorithms

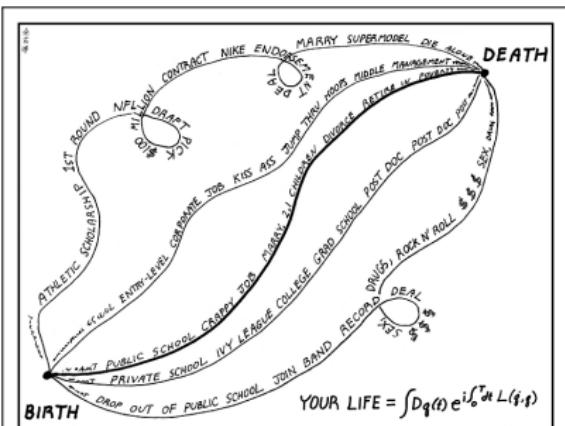
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More botanical names:
bootstrapping, spawning,
cloning, pruning,
replenish, enrichment, go
with the winner, *some
new ideas* ?



A single stochastic model

Particle interpretation of Feynman-Kac path integrals



The Path Integral Formulation of Your Life

Introduction

Feynman-Kac models

Some basic notation

Description of the models

The 3 keys formulae

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Basic notation

$\mathcal{P}(E)$ probability meas., $\mathcal{B}(E)$ bounded functions on E .

- ▶ $(\mu, f) \in \mathcal{P}(E) \times \mathcal{B}(E) \quad \longrightarrow \quad \mu(f) = \int \mu(dx) f(x)$
- ▶ $Q(x_1, dx_2)$ **integral operators** $x_1 \in E_1 \rightsquigarrow x_2 \in E_2$

$$\begin{aligned} Q(f)(x_1) &= \int Q(x_1, dx_2) f(x_2) \\ [\mu Q](dx_2) &= \int \mu(dx_1) Q(x_1, dx_2) \quad (\Rightarrow [\mu Q](f) = \mu[Q(f)]) \end{aligned}$$

- ▶ **Boltzmann-Gibbs transformation**

[Positive and bounded potential function G]

$$\mu(dx) \mapsto \Psi_G(\mu)(dx) = \frac{1}{\mu(G)} G(x) \mu(dx)$$

Basic notation

$E = \{1, \dots, d\} \rightsquigarrow \text{matrix operations}$

$$\mu = [\mu(1), \dots, \mu(d)] \quad Q = (Q(i,j))_{1 \leq i,j \leq d} \quad f = \begin{bmatrix} f(1) \\ \vdots \\ f(d) \end{bmatrix}$$

and

$$\Psi_G(\mu) \propto \mu \operatorname{Diag}(G(1), \dots, G(d))$$

Feynman-Kac measures

- ▶ Markov chain X_n on E_n , with transitions M_n :

$$\mathbb{P}_n(d(x_0, \dots, x_n)) = \eta_0(dx_0) M_1(x_0, dx_1) \dots M_n(x_{n-1}, dx_n)$$

- ▶ Potential functions $G_n(x_n) \in [0, 1]$

$$d\mathbb{Q}_n := \frac{1}{Z_n} \left\{ \prod_{0 \leq p < n} G_p(X_p) \right\} d\mathbb{P}_n$$

Flow of n -marginals

$$\eta_n(f) = \gamma_n(f)/\gamma_n(1) \quad \text{with} \quad \gamma_n(f) := \mathbb{E} \left(f(X_n) \prod_{0 \leq p < n} G_p(X_p) \right)$$

Nonlinear equation:

$$\eta_{n+1} = \Psi_{G_n}(\eta_n) M_{n+1} := \Phi_{n+1}(\eta_n)$$

2 "elementary" illustrations

► Confinements :

Random walk $X_n \in \mathbb{Z}^d$, $X_0 = 0$ & $G_n := 1_{[-L,L]}$, $L > 0$.

$$\mathbb{Q}_n = \text{Law}((X_0, \dots, X_n) \mid X_p \in [-L, L], \forall 0 \leq p < n)$$

and

$$\mathcal{Z}_n = \gamma_n(1) = \text{Proba}(X_p \in [-L, L], \forall 0 \leq p < n)$$

► Self avoiding walks

$$\mathbf{X}_n = (X_0, \dots, X_n) \quad \& \quad \mathbf{G}_n(\mathbf{X}_n) = 1_{X_n \notin \{X_0, \dots, X_{n-1}\}}$$

$$\mathbb{Q}_n = \text{Law}((\mathbf{X}_0, \dots, \mathbf{X}_n) \mid X_p \neq X_q, \forall 0 \leq p < q < n)$$

and

$$\mathcal{Z}_n = \gamma_n(1) = \text{Proba}(X_p \neq X_q, \forall 0 \leq p < q < n)$$

▷ Continuous time models

$$X_n := X'_{[t_n, t_{n+1}[} \quad \& \quad G_n(X_n) = \exp \int_{t_n}^{t_{n+1}} V_s(X'_s) ds$$



$$\prod_{0 \leq p < n} G_p(X_p) = \exp \left\{ \int_{t_0}^{t_n} V_s(X'_s) ds \right\}$$

or using a simple "Euler's scheme" $X'_{t_p} = X_p$

$$e^{\int_{t_0}^{t_n} [V_s(X'_s) ds + W_s(X'_s) dB_s]} \simeq \prod_{0 \leq p < n} e^{V_{t_p}(X_p) \Delta t + W_{t_p}(X_p) \sqrt{\Delta t} N_p(0,1)}$$



The 3 keys formulae

- ▶ Time marginal measures = Path space measures:

$$\gamma_n(f_n) = \mathbb{E} \left(f_n(\mathbf{X}_n) \prod_{0 \leq p < n} \mathbf{G}_p(\mathbf{X}_p) \right)$$

$$[\mathbf{X}_n := (X_0, \dots, X_n) \quad \& \quad \mathbf{G}_n(\mathbf{X}_n) = G_n(X_n)] \implies \eta_n = \mathbb{Q}_n$$

- ▶ Normalizing constants (= Free energy models):

$$\mathcal{Z}_n = \mathbb{E} \left(\prod_{0 \leq p < n} G_p(X_p) \right) = \prod_{0 \leq p < n} \eta_p(G_p)$$

The last key

► Backward Markov models

$$\mathbb{Q}_n(dx_0, \dots, dx_n) \propto \eta_0(dx_0) Q_1(x_0, dx_1) \dots Q_n(x_{n-1}, dx_n)$$

with

$$\begin{aligned} Q_n(x_{n-1}, dx_n) &:= G_{n-1}(x_{n-1}) M_n(x_{n-1}, dx_n) \\ &\stackrel{\text{hyp}}{=} H_n(x_{n-1}, x_n) \nu_n(dx_n) \\ \Rightarrow \eta_{n+1}(dx) &= \frac{1}{\eta_n(G_n)} \eta_n(H_{n+1}(\cdot, x)) \nu_{n+1}(dx) \end{aligned}$$

If we set

$$\mathbb{M}_{n+1, \eta_n}(x_{n+1}, dx_n) = \frac{\eta_n(dx_n) H_{n+1}(x_n, x_{n+1})}{\eta_n(H_{n+1}(\cdot, x_{n+1}))}$$

then we find the backward equation

$$\eta_{n+1}(dx_{n+1}) \mathbb{M}_{n+1, \eta_n}(x_{n+1}, dx_n) = \frac{1}{\eta_n(G_n)} \eta_n(dx_n) Q_{n+1}(x_n, dx_{n+1})$$

The last key (continued)

$$\mathbb{Q}_n(d(x_0, \dots, x_n)) \propto \eta_0(dx_0) Q_1(x_0, dx_1) \dots Q_n(x_{n-1}, dx_n)$$

⊕

$$\eta_{n+1}(dx_{n+1}) \mathbb{M}_{n+1, \eta_n}(x_{n+1}, dx_n) \propto \eta_n(dx_n) Q_{n+1}(x_n, dx_{n+1})$$

↓

Backward Markov chain model :

$$\mathbb{Q}_n(d(x_0, \dots, x_n)) = \eta_n(dx_n) \mathbb{M}_{n, \eta_{n-1}}(x_n, dx_{n-1}) \dots \mathbb{M}_{1, \eta_0}(x_1, dx_0)$$

with the dual/backward Markov transitions

$$\mathbb{M}_{n+1, \eta_n}(x_{n+1}, dx_n) \propto \eta_n(dx_n) H_{n+1}(x_n, x_{n+1})$$

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A bad tempting idea

Adaptive interacting jumps

The 4 particle estimates

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A bad tempting idea = i.i.d. weighted samples X_n^i

$$\mathcal{Z}_n := \mathbb{E} \left(\prod_{0 \leq p < n} G_p(X_p) \right) \simeq \mathcal{Z}_n^N := \frac{1}{N} \sum_{i=1}^N \prod_{0 \leq p < n} G_p(X_p^i)$$

Example :

X_n simple RW $\in \mathbb{Z}^d$, $G_n = 1_{[-10,10]}$ $\Rightarrow \exists n = n(\omega) : \mathcal{Z}_n^N = 0$

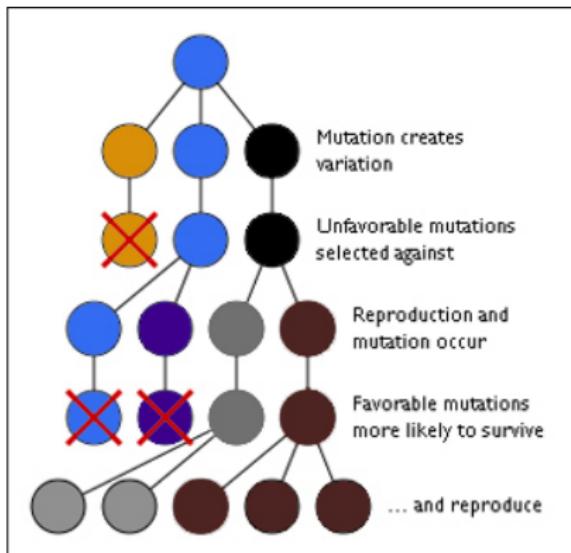
$$N \mathbb{E} \left(\left[\frac{\mathcal{Z}_n^N}{\mathcal{Z}_n} - 1 \right]^2 \right) = \frac{1 - \mathcal{Z}_n}{\mathcal{Z}_n} \simeq \text{Proba}(X_p \in A, \forall 0 \leq p < n)^{-1}$$

Our objective

$$N \mathbb{E} \left(\left[\frac{\mathcal{Z}_n^N}{\mathcal{Z}_n} - 1 \right]^2 \right) \leq c \times n$$

Solution = Genetic style interacting jumps

- ▶ Mutation-Exploration-Proposals w.r.t. transitions M_n .
- ▶ Selection-Recycling-(Accept.-Rejet) w.r.t. potential G_n .



Stochastic population dynamics

$$\begin{bmatrix} \xi_n^1 \\ \vdots \\ \xi_n^i \\ \vdots \\ \xi_n^N \end{bmatrix} \xrightarrow{S_{n,\eta_n^N}} \begin{bmatrix} \widehat{\xi}_n^1 & \xrightarrow{M_{n+1}} & \xi_{n+1}^1 \\ \vdots & & \vdots \\ \widehat{\xi}_n^i & \longrightarrow & \xi_{n+1}^i \\ \vdots & & \vdots \\ \widehat{\xi}_n^N & \longrightarrow & \xi_{n+1}^N \end{bmatrix}$$

Accept/Reject-Recycle=Selection =Geometric recycling interacting jumps

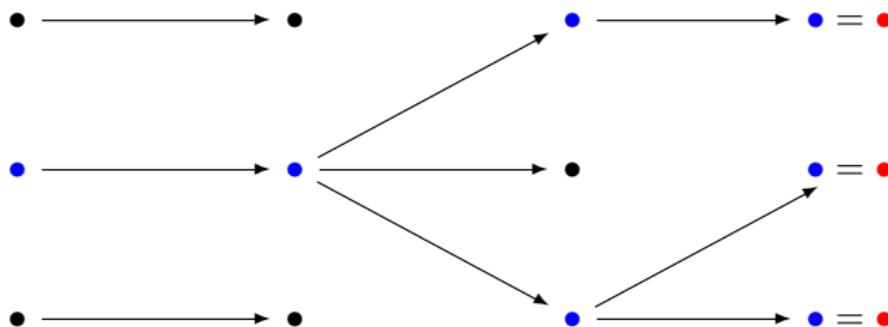
$$S_{n,\eta_n^N}(\xi_n^i, dx)$$

$$:= G_n(\xi_n^i) \delta_{\xi_n^i}(dx) + (1 - G_n(\xi_n^i)) \sum_{j=1}^N \frac{G_n(\xi_n^j)}{\sum_{k=1}^N G_n(\xi_n^k)} \delta_{\xi_n^j}(dx)$$

$$\text{Ex. : } G_n = 1_A \rightsquigarrow G_n(\xi_n^i) = 1_A(\xi_n^i)$$

The 4 particle estimates

Genealogical tree evolution $(N, n) = (3, 3)$

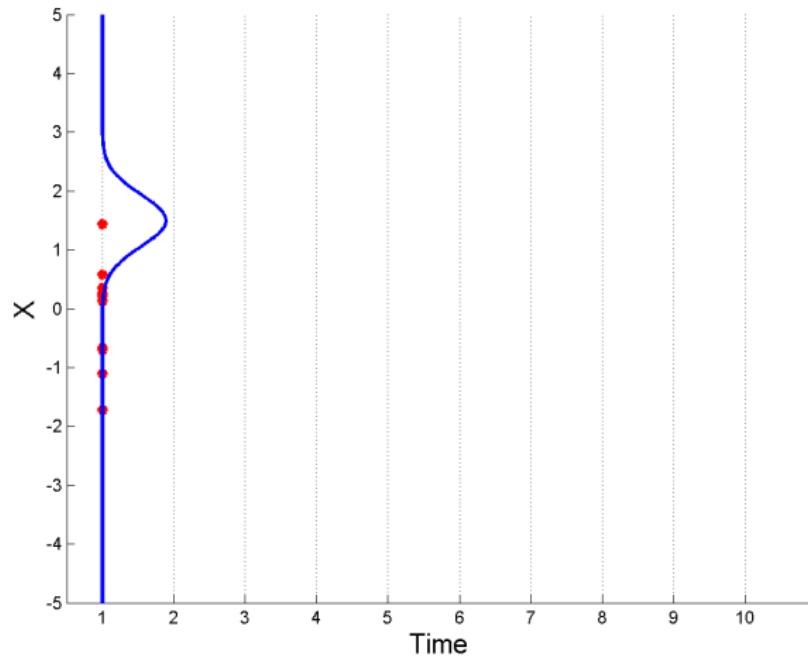


- ▶ Individuals in the current population

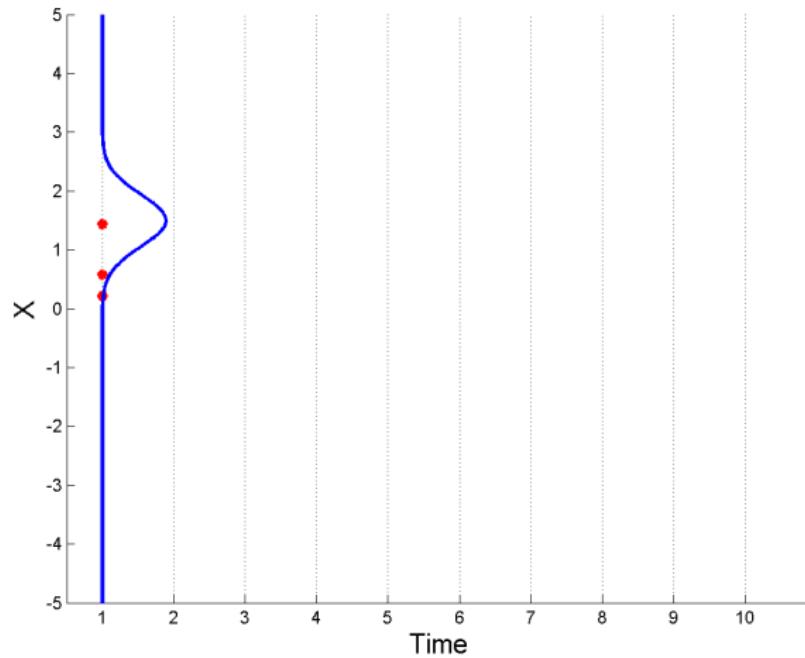
= Almost i.i.d. samples w.r.t. FK marginal meas. η_n

$$\eta_n^N := \frac{1}{N} \sum_{1 \leq i \leq N} \delta_{\xi_n^i} \longrightarrow_{N \rightarrow \infty} \eta_n = \text{solution of a nonlinear m.v.p.}$$

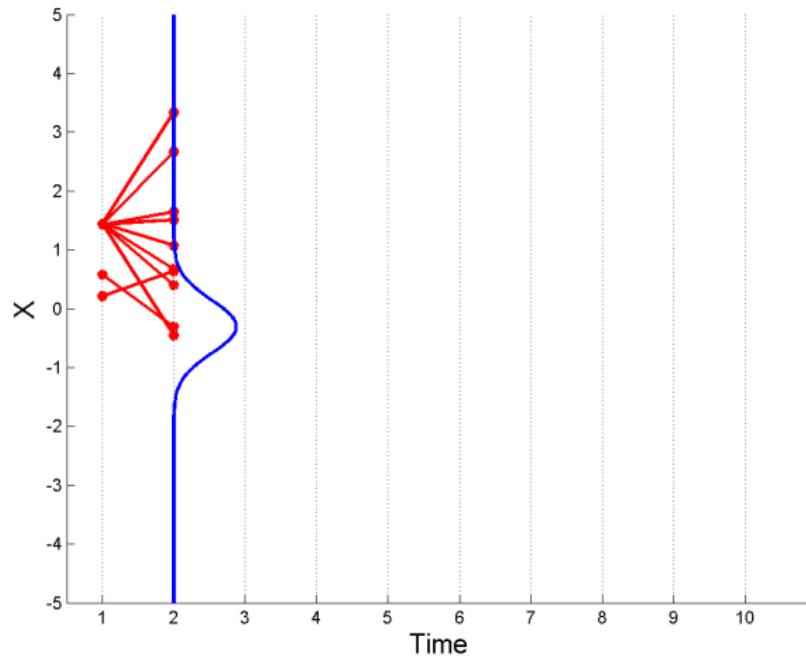
Graphical illustration : $\eta_n^N := \frac{1}{N} \sum_{1 \leq i \leq N} \delta_{\xi_i} \simeq \eta_n$



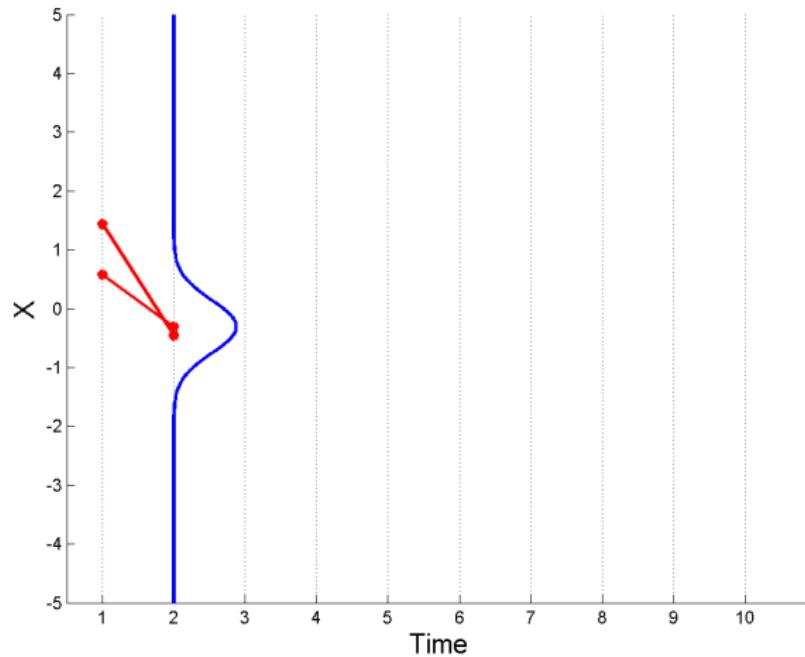
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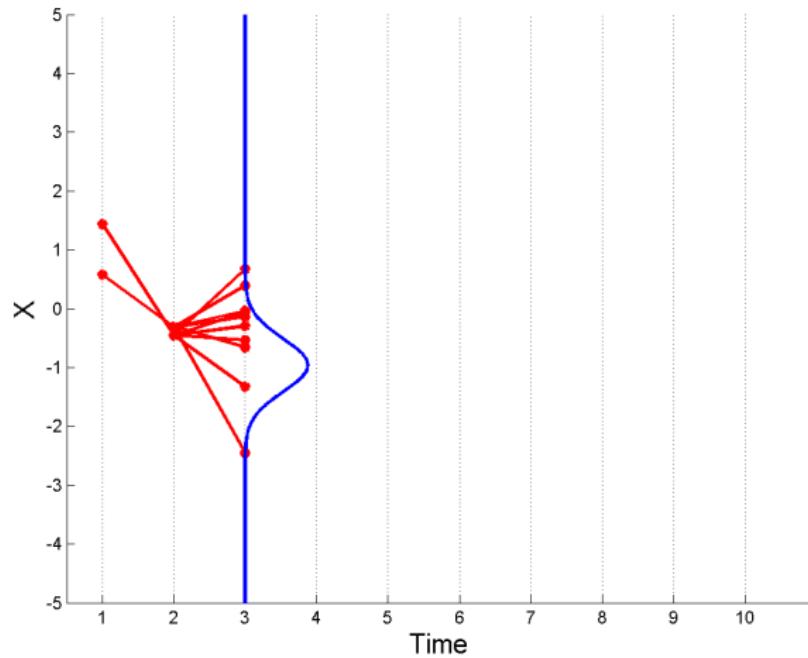
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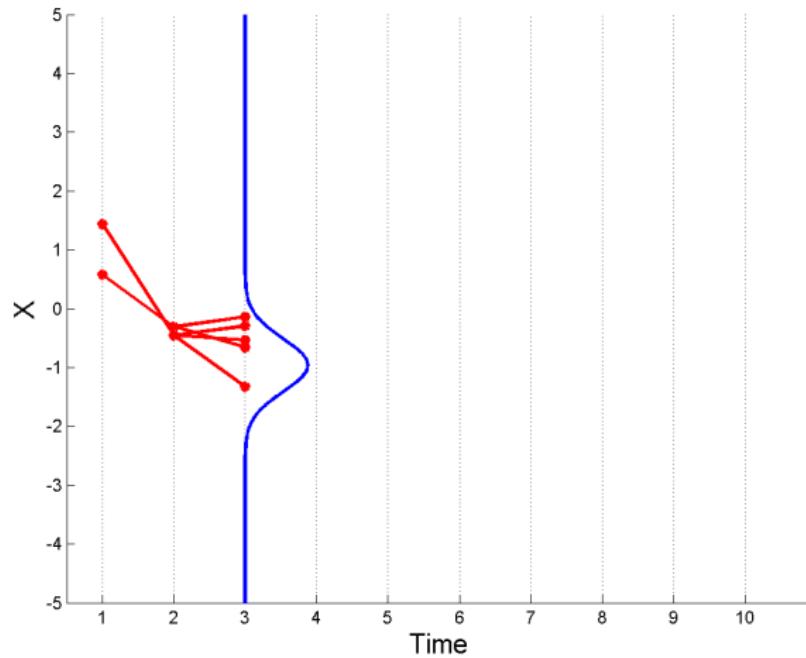
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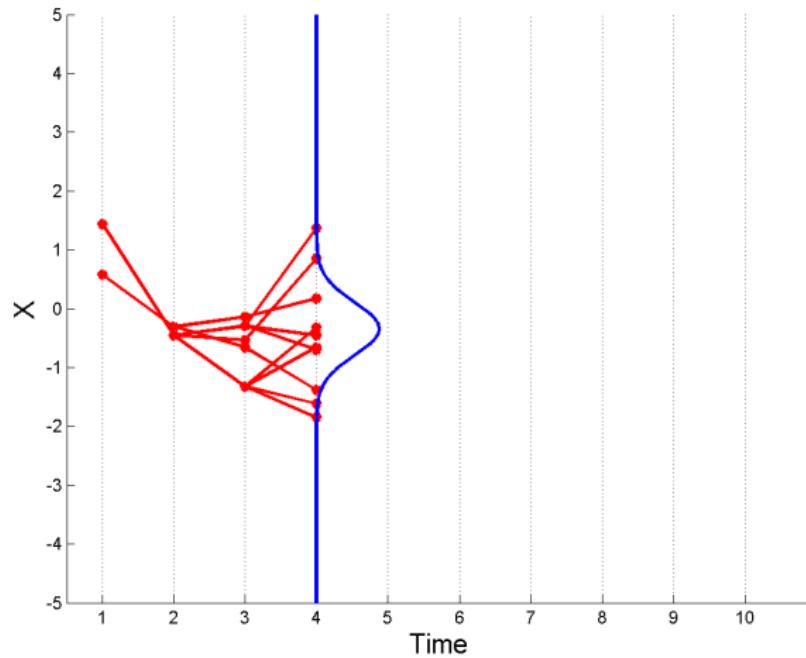
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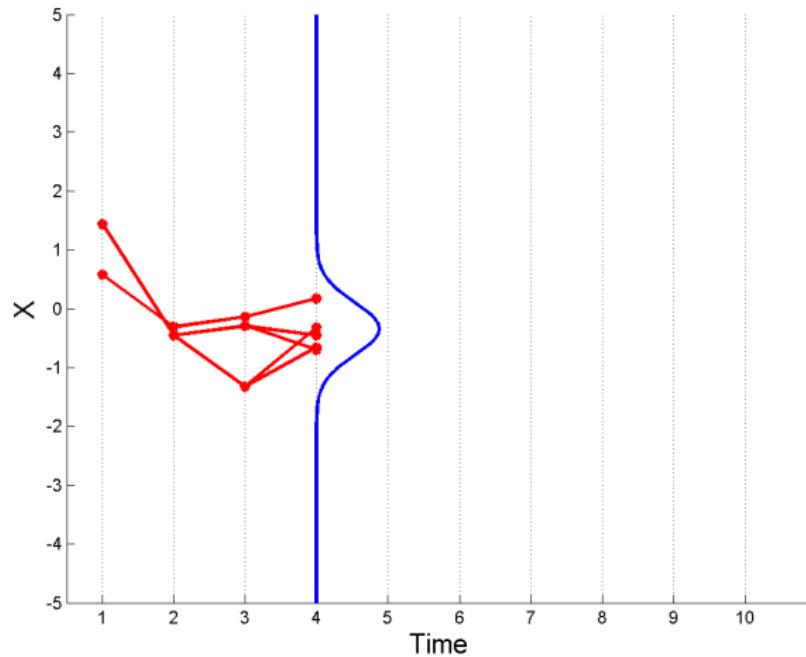
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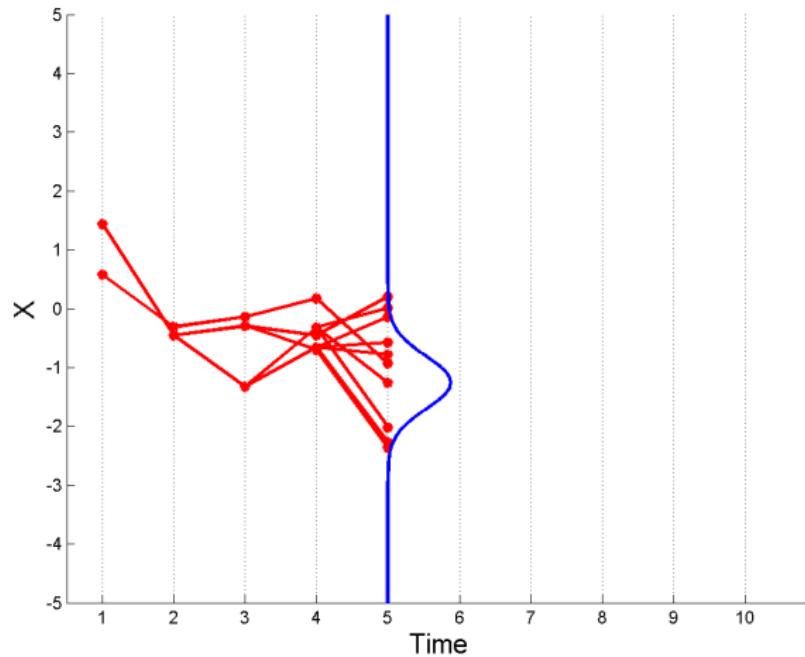
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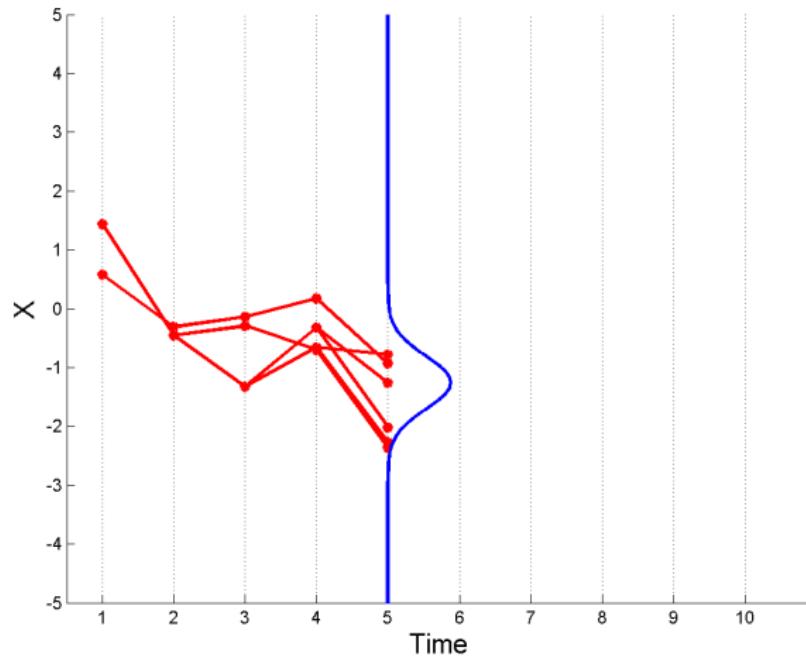
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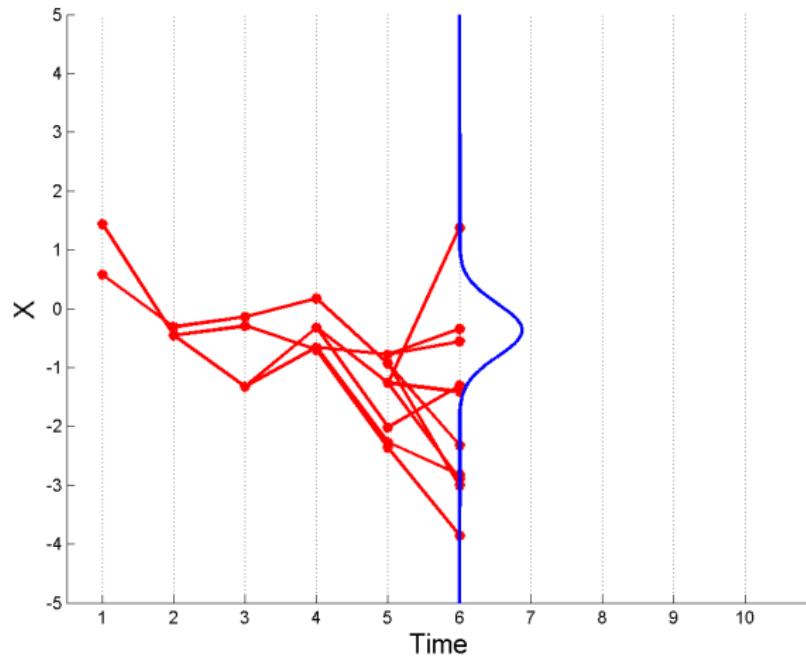
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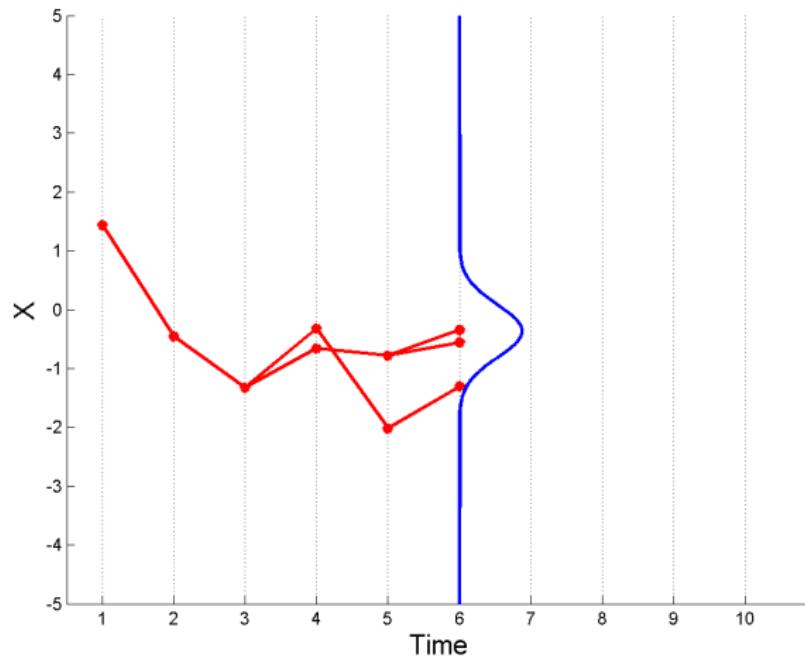
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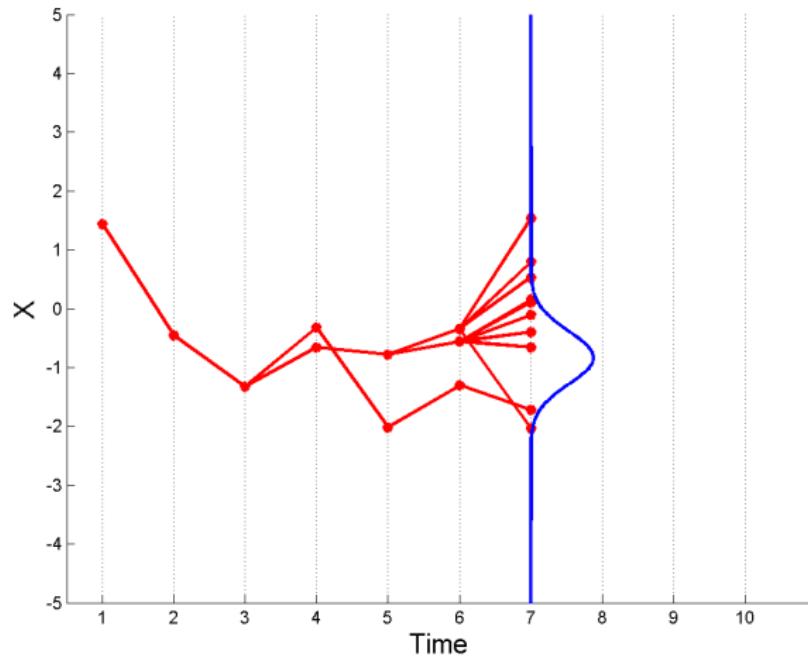
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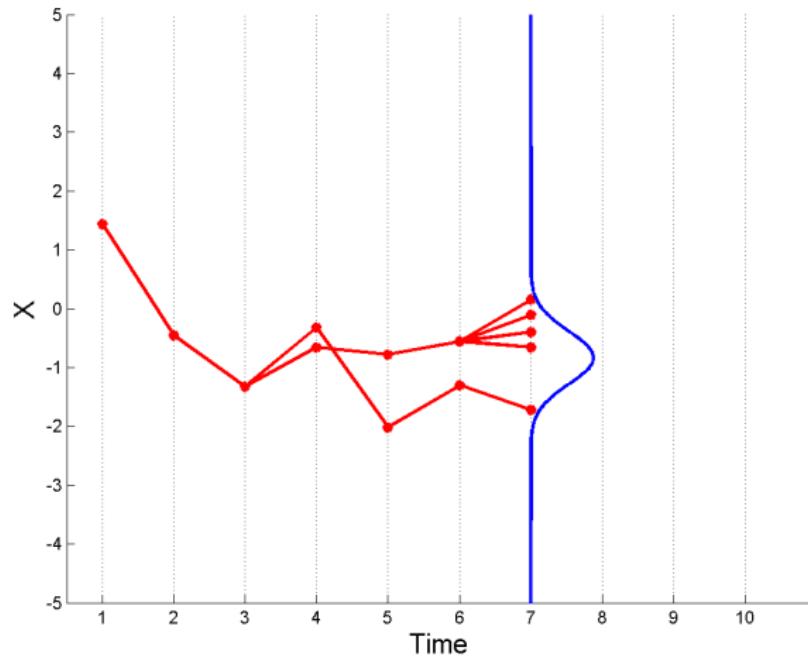
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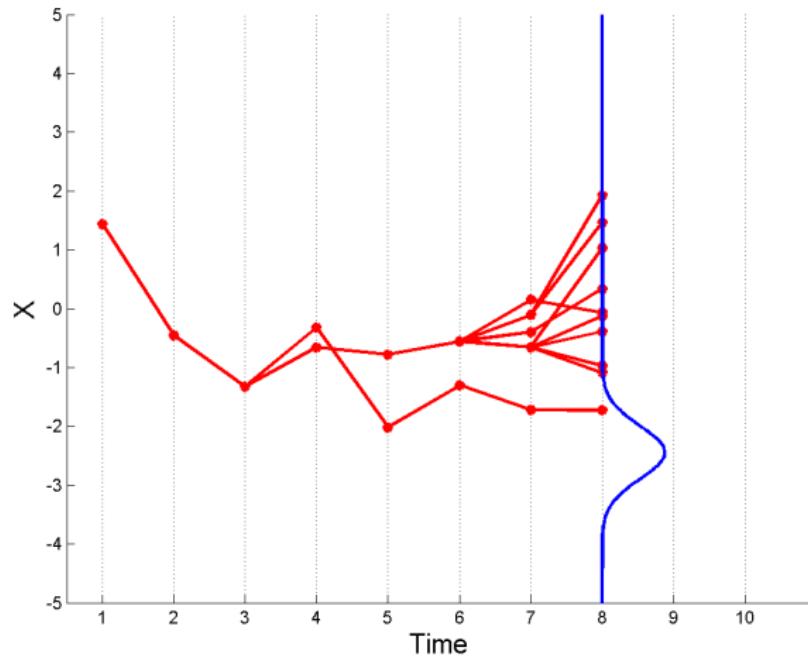
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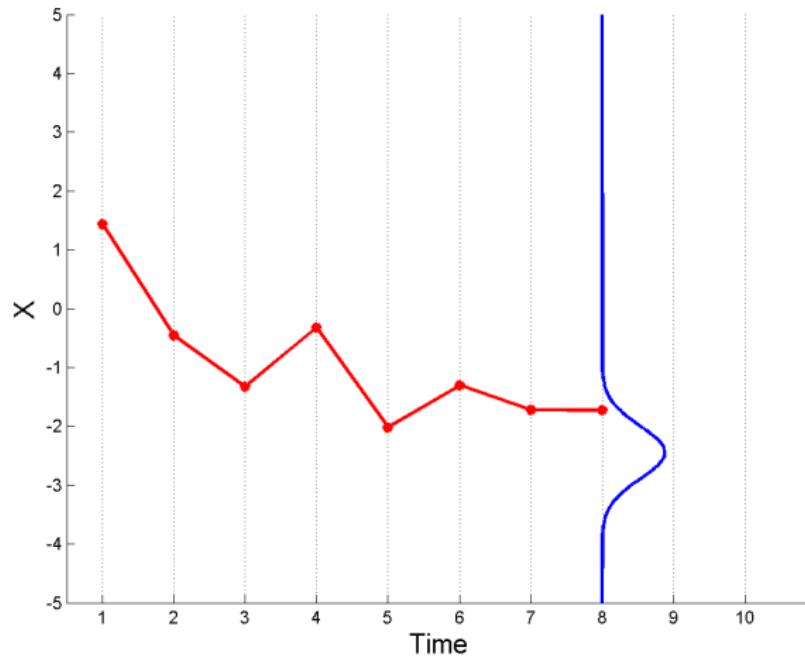
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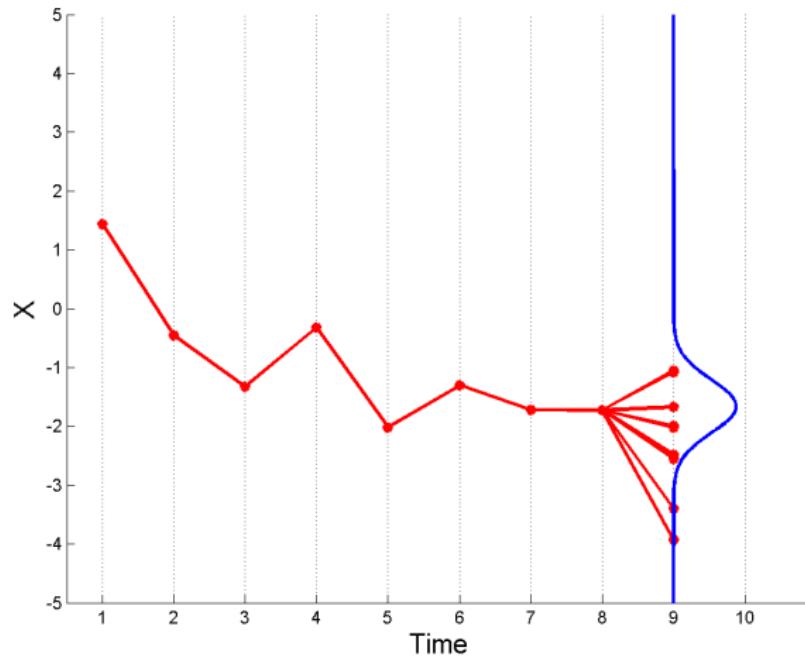
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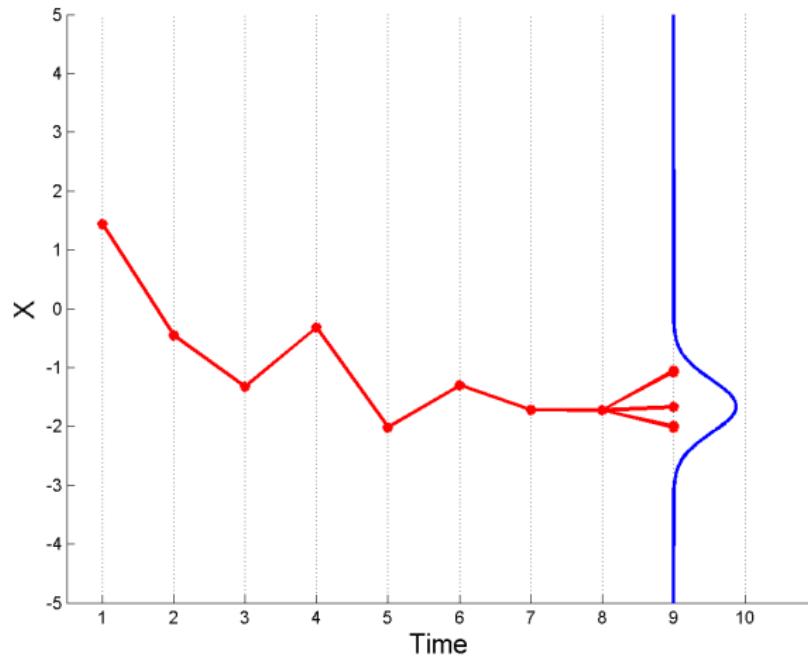
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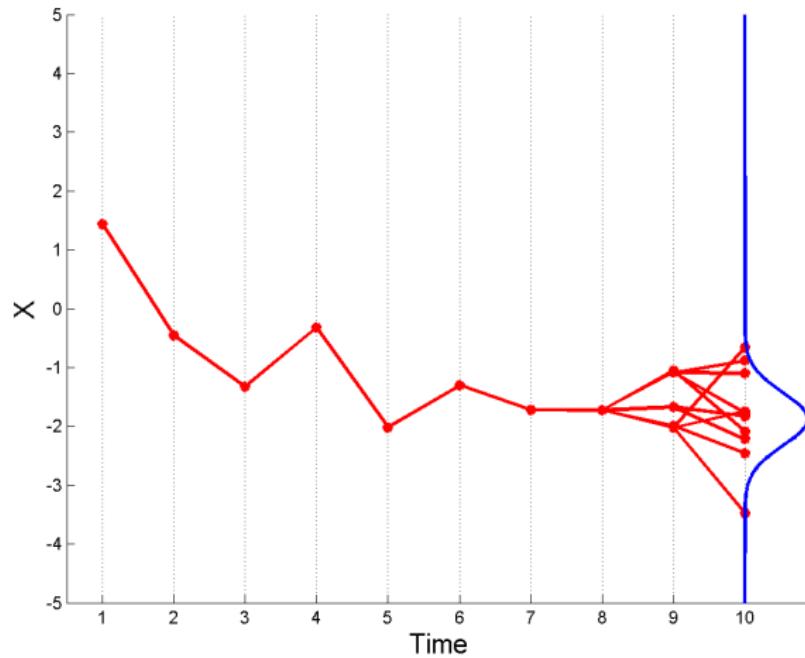
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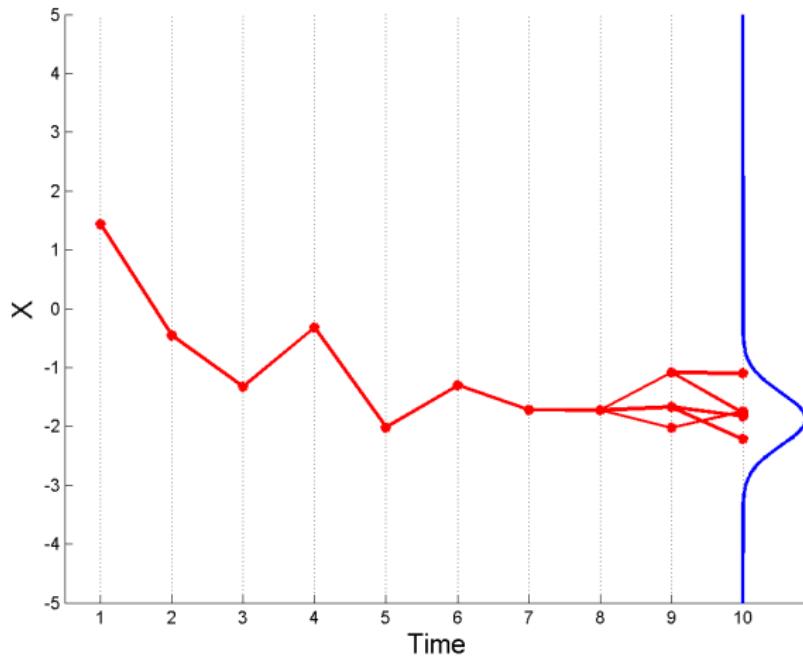
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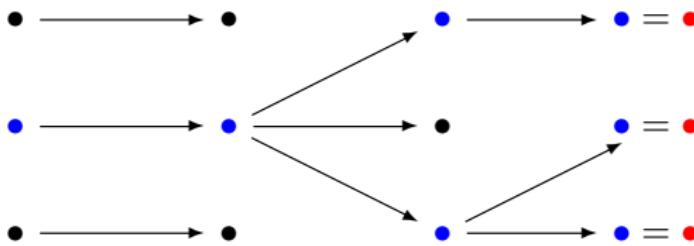
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Graphical illustration : $\eta_n^N := \frac{1}{N} \sum_{1 \leq i \leq N} \delta_{\xi_i} \simeq \eta_n$



Two more particle estimates



- Ancestral lines = Almost i.i.d. sampled paths w.r.t. \mathbb{Q}_n .

$$(\xi_{0,n}^i, \xi_{1,n}^i, \dots, \xi_{n,n}^i) := \text{*i*-th ancestral line } \text{*i*-th current individual} = \xi_n^i$$

$$\frac{1}{N} \sum_{1 \leq i \leq N} \delta_{(\xi_{0,n}^i, \xi_{1,n}^i, \dots, \xi_{n,n}^i)} \xrightarrow{N \rightarrow \infty} \mathbb{Q}_n$$

- Unbiased particle free energy functions

$$\mathcal{Z}_n^{\textcolor{red}{N}} = \prod_{0 \leq p < n} \eta_p^{\textcolor{red}{N}}(G_p) \xrightarrow{N \rightarrow \infty} \mathcal{Z}_n = \prod_{0 \leq p < n} \eta_p(G_p)$$

... and the last particle measure

$$\mathbb{Q}_n^N(d(x_0, \dots, x_n)) := \eta_n^N(dx_n) \mathbb{M}_{n, \eta_{n-1}^N}(x_n, dx_{n-1}) \dots \mathbb{M}_{1, \eta_0^N}(x_1, dx_0)$$

with the random particle matrices:

$$\mathbb{M}_{n+1, \eta_n^N}(x_{n+1}, dx_n) \propto \eta_n^N(dx_n) H_{n+1}(x_n, x_{n+1})$$

Example: Normalized additive functionals

$$\mathbf{f}_n(x_0, \dots, x_n) = \frac{1}{n+1} \sum_{0 \leq p \leq n} f_p(x_p)$$



$$\mathbb{Q}_n^N(\mathbf{f}_n) := \frac{1}{n+1} \sum_{0 \leq p \leq n} \underbrace{\eta_n^N \mathbb{M}_{n, \eta_{n-1}^N} \dots \mathbb{M}_{p+1, \eta_p^N}(f_p)}_{\text{matrix operations}}$$

Island models

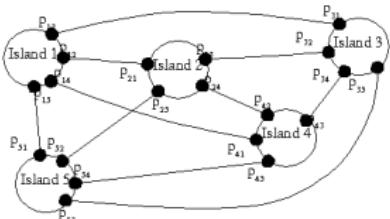


Fig. 3.4 Schematic of a genetic algorithm using island migration

Reminder : the unbiased property

$$\begin{aligned}\mathbb{E} \left(f_n(\mathbf{X}_n) \prod_{0 \leq p < n} G_p(\mathbf{X}_p) \right) &= \mathbb{E} \left(\eta_n^N(f_n) \prod_{0 \leq p < n} \eta_p^N(G_p) \right) \\ &= \mathbb{E} \left(F_n(\mathcal{X}_n) \prod_{0 \leq p < n} \mathcal{G}_p(\mathcal{X}_p) \right)\end{aligned}$$

with the Island evolution Markov chain model

$$\mathcal{X}_n := \eta_n^N \quad \text{and} \quad \mathcal{G}_n(\mathcal{X}_n) = \eta_n^N(\mathbf{G}_n) = \mathcal{X}_n(\mathbf{G}_n)$$

\Rightarrow particle model with $(\mathcal{X}_n, \mathcal{G}_n(\mathcal{X}_n)) = \text{Interacting Island particle model}$

Introduction

Feynman-Kac models

Interacting particle systems

Concentration inequalities

Current population models

Particle free energy

Genealogical tree models

Backward particle models

Some ideas on how it works

Current population models

Constants (c_1, c_2) related to (bias, variance), c universal constant
Test funct. $\|f_n\| \leq 1, \forall (x \geq 0, n \geq 0, N \geq 1)$.

- ▶ The probability of the event

$$[\eta_n^N - \eta_n](f) \leq \frac{c_1}{N} (1 + x + \sqrt{x}) + \frac{c_2}{\sqrt{N}} \sqrt{x}$$

is greater than $1 - e^{-x}$.

- ▶ $x = (x_i)_{1 \leq i \leq d} \rightsquigarrow (-\infty, x] = \prod_{i=1}^d (-\infty, x_i]$ cells in $E_n = \mathbb{R}^d$.

$$F_n(x) = \eta_n(1_{(-\infty, x]}) \quad \text{and} \quad F_n^N(x) = \eta_n^N(1_{(-\infty, x]})$$

The probability of the following event

$$\sqrt{N} \|F_n^N - F_n\| \leq c \sqrt{d(x+1)}$$

is greater than $1 - e^{-x}$.

Particle free energy models

Constants (c_1, c_2) related to (bias, variance), c universal constant
 $\forall (x \geq 0, n \geq 0, N \geq 1)$

- Unbiased property

$$\mathbb{E} \left(\eta_n^N(f_n) \prod_{0 \leq p < n} \eta_p^N(G_p) \right) = \mathbb{E} \left(f_n(X_n) \prod_{0 \leq p < n} G_p(X_p) \right)$$

- For any $\epsilon \in \{+1, -1\}$, the probability of the event

$$\frac{\epsilon}{n} \log \frac{\mathcal{Z}_n^N}{\mathcal{Z}_n} \leq \frac{c_1}{N} (1 + x + \sqrt{x}) + \frac{c_2}{\sqrt{N}} \sqrt{x}$$

is greater than $1 - e^{-x}$.

note $(0 \leq \epsilon \leq 1 \Rightarrow (1 - e^{-\epsilon}) \vee (e^\epsilon - 1) \leq 2\epsilon)$

$$e^{-\epsilon} \leq \frac{z^N}{z} \leq e^\epsilon \Rightarrow \left| \frac{z^N}{z} - 1 \right| \leq 2\epsilon$$

Genealogical tree models := η_n^N (in path space)

Constants (c_1, c_2) related to (bias, variance), c universal constant
 \mathbf{f}_n test function $\|\mathbf{f}_n\| \leq 1$, $\forall (x \geq 0, n \geq 0, N \geq 1)$.

- ▶ The probability of the event

$$[\eta_n^N - \mathbb{Q}_n](f) \leq c_1 \frac{n+1}{N} (1 + x + \sqrt{x}) + c_2 \sqrt{\frac{(n+1)}{N}} \sqrt{x}$$

is greater than $1 - e^{-x}$.

- ▶ \mathcal{F}_n = indicator fct. \mathbf{f}_n of cells in $\mathbf{E}_n = (\mathbb{R}^{d_0} \times \dots \times \mathbb{R}^{d_n})$
The probability of the following event

$$\sup_{\mathbf{f}_n \in \mathcal{F}_n} |\eta_n^N(\mathbf{f}_n) - \mathbb{Q}_n(\mathbf{f}_n)| \leq c (n+1) \sqrt{\frac{\sum_{0 \leq p \leq n} d_p}{N} (x+1)}$$

is greater than $1 - e^{-x}$.

Backward particle models

Constants (c_1, c_2) related to (bias, variance), c universal constant.

\mathbf{f}_n normalized additive functional with $\|f_p\| \leq 1$, $\forall (x \geq 0, n \geq 0, N \geq 1)$

- ▶ The probability of the event

$$[\mathbb{Q}_n^N - \mathbb{Q}_n](\bar{\mathbf{f}}_n) \leq c_1 \frac{1}{N} (1 + (x + \sqrt{x})) + c_2 \sqrt{\frac{x}{N(n+1)}}$$

- ▶ $\mathbf{f}_{a,n}$ normalized additive functional w.r.t. $f_p = 1_{(-\infty, a]}$, $a \in \mathbb{R}^d = E_n$

.

The probability of the following event

$$\sup_{a \in \mathbb{R}^d} |\mathbb{Q}_n^N(\mathbf{f}_{a,n}) - \mathbb{Q}_n(\mathbf{f}_{a,n})| \leq c \sqrt{\frac{d}{N}(x+1)}$$

is greater than $1 - e^{-x}$.

Introduction

Feynman-Kac models

Interacting particle systems

Concentration inequalities

Some ideas on how it works

Some ideas on how it works

Feynman-Kac evolution equations

$$\eta_n \xrightarrow{\text{Correction-Selection}} \widehat{\eta}_n = \Psi_{G_n}(\eta_n) \xrightarrow{\text{Prediction-Mutation}} \eta_{n+1} = \widehat{\eta}_n M_{n+1}$$

with the nonlinear mass transport equation

$$\Psi_{G_n}(\eta_n)(dy) := \frac{1}{\mu(G_n)} G_n(y) \eta_n(dy) = \int \eta_n(dx) S_{n,\eta_n}(x, dy)$$

with

$$S_{n,\eta_n}(x, .) := G_n(x) \delta_x + (1 - G_n(x)) \Psi_{G_n}(\eta_n)$$

↓

$$\eta_{n+1} = \eta_n (S_{n,\eta_n} M_{n+1}) := \eta_n K_{n+1,\eta_n} = \text{Law}(\overline{X}_n)$$

McKean Markov chain model \overline{X}_n = Perfect sampler

η_n = Law of \overline{X}_n = **nonlinear Markov chain**

$$\mathbb{P}(\overline{X}_{n+1} \in dx \mid \overline{X}_n) = K_{n+1, \eta_n}(\overline{X}_n, dx)$$

Mean field particle model = Stochastic population dynamic model

$\rightsquigarrow \xi_n = (\xi_n^1, \dots, \xi_n^N) \in E_n^N$ **such that**

$$\eta_n^N := \frac{1}{N} \sum_{1 \leq i \leq N} \delta_{\xi_n^i} \simeq_{N \uparrow \infty} \eta_n$$

\rightsquigarrow Natural approximated transitions

$$\xi_n^i \rightsquigarrow \xi_{n+1}^i \sim K_{n+1, \eta_n^N}(\xi_n^i, dx)$$

Some key advantages

- ▶ Mean field models = Stochastic linearization/perturbation technique

$$\eta_n^N = \eta_{n-1}^N K_{n,\eta_{n-1}^N} + \frac{1}{\sqrt{N}} W_n^N$$

with $W_n^N \simeq W_n$ independent centered Gaussian fields .

- ▶ $\eta_n = \eta_{n-1} K_{n,\eta_{n-1}}$ stable \Rightarrow Non propagation of local sampling errors

\implies Uniform control w.r.t. the time horizon

- ▶ "No burning, no need to study the stability of MCMC models".
- ▶ Stochastic adaptive grid approximation
- ▶ Nonlinear system \rightsquigarrow positive - beneficial interactions.
- ▶ Simple and natural sampling algorithm.
- ▶ Local conditional iid samples \oplus Stability of nonlinear sg
 \rightsquigarrow New concentration and empirical process theory