

Some stoch. models and methods in risk analysis

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SOME HYPER-REFS:

- ▶ Feynman-Kac formulae, Genealogical & Interacting Particle Systems with appl., Springer (2004)
- ▶ Sequential Monte Carlo Samplers JRSS B. (2006). (joint work with A. Doucet & A. Jasra)
- ▶ A Backward Particle Interpretation of Feynman-Kac Formulae M2AN (2010). (joint work with A. Doucet & S.S. Singh)
- ▶ On the concentration of interacting processes. Foundations & Trends in Machine Learning [170p.] (2012). (joint work with Peng Hu & Li Ming Wu) [+ Refs]
- ▶ More ref. on the website : Feynman-Kac particle models & their applications domains [+ Links]

Introduction

Feynman-Kac models

7 rare event models/problems

Exponential concentration analysis

Introduction

Some basic notation

Importance sampling

Acceptance-rejection samplers

Feynman-Kac models

7 rare event models/problems

Exponential concentration analysis

Basic notation $\mu \in \mathcal{P}(E)$ **proba. meas.**, $f \in \mathcal{B}(E)$ **bounded funct. on E .**

- ▶ $Q(x_1, dx_2)$ **integral operator** $x_1 \in E_1 \rightsquigarrow x_2 \in E_2$

$$Q(f)(x_1) = \int Q(x_1, dx_2) f(x_2) \quad \text{and} \quad \mu(f) = \int \mu(dx_2) f(x_2)$$

$$[\mu Q](dx_2) = \int \mu(dx_1) Q(x_1, dx_2) \quad (\implies [\mu Q](f) = \mu[Q(f)])$$

- ▶ **Boltzmann-Gibbs transformation** [$G \geq 0$ and μ s.t. $\mu(G) > 0$]

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

(Bayes' rule/conditioning/restriction formula)

Importance sampling and optimal twisted measures

$\mathbb{P}(X \in A) = \mathbb{P}_X(A) = 10^{-10} \rightsquigarrow$ Find \mathbb{P}_Y t.q. $\mathbb{P}_Y(A) = \mathbb{P}(Y \in A) \simeq 1$

\rightsquigarrow **Crude Monte Carlo sampling** Y^i i.i.d. \mathbb{P}_Y

$$\mathbb{P}_Y \left(\frac{d\mathbb{P}_X}{d\mathbb{P}_Y} 1_A \right) = \mathbb{P}_X(A) \simeq \mathbb{P}_X^N(A) := \frac{1}{N} \sum_{1 \leq i \leq N} \frac{d\mathbb{P}_X}{d\mathbb{P}_Y}(Y^i) 1_A(Y^i)$$

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Optimal twisted measure = Conditional distribution

$$\text{Variance} = 0 \iff \mathbb{P}_Y = \text{Law}(X \mid X \in A) = \psi_{1_A}(\mathbb{P}_X)$$



Perfect or MCMC samplers = acceptance-rejection techniques

BUT

(Very often) with very small acceptance rates

Conditional distributions and Feynman-Kac models

Example : Markov chain models X_n restricted to subsets A_n

$$\mathbf{X} = (X_0, \dots, X_n) \in \mathbf{A} = (A_0 \times \dots \times A_n)$$

The conditional distributions

$$\text{Law}(\mathbf{X} \mid \mathbf{X} \in \mathbf{A}) = \text{Law}((X_0, \dots, X_n) \mid X_p \in A_p, p < n) = \mathbb{Q}_n$$

and

$$\text{Proba}(X_p \in A_p, p < n) = \mathcal{Z}_n$$

Conditional distributions and Feynman-Kac models

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$$\text{Proba}(X_p \in A_p, p < n) = \mathcal{Z}_n$$

are given by the Feynman-Kac measures

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

with

$$\mathbb{P}_n = \text{Law}(X_0, \dots, X_n) \quad \& \quad G_p = 1_{A_p}, \quad p < n$$

Introduction

Feynman-Kac models

Nonlinear evolution equation

Interacting particle samplers

Particle estimates

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Exponential concentration analysis

Feynman-Kac models (general $G_n(X_n)$ & $X_n \in E_n$)

Flow of n -marginals

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

$$\Downarrow (\gamma_n(1) = \mathcal{Z}_n)$$

Nonlinear evolution equation :

$$\begin{aligned} \eta_{n+1} &= \Psi_{G_n}(\eta_n) M_{n+1} \\ \mathcal{Z}_{n+1} &= \eta_n(G_n) \times \mathcal{Z}_n \end{aligned}$$

with the Markov transitions

$$M_{n+1}(x_n, dx_{n+1}) = \mathbb{P}(X_{n+1} \in dx_{n+1} \mid X_n = x_n)$$

Note : $[X_n = (X'_0, \dots, X'_n) \text{ \& } G_n(X_n) = G'(X'_n)] \implies \eta_n = \mathbb{Q}'_n$

Interacting particle samplers

Nonlinear evolution equation :

$$\begin{aligned}\eta_{n+1} &= \Psi_{G_n}(\eta_n) M_{n+1} \\ \mathcal{Z}_{n+1} &= \eta_n(G_n) \times \mathcal{Z}_n\end{aligned}$$

↪ **(Sequential) Interacting Particle simulation technique**

G_n -acceptance-rejection with recycling \oplus M_n -propositions

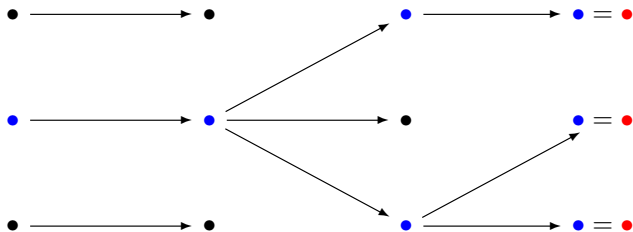
\Updownarrow (Genetic type branching/Interacting jump particle model)

$$\xi_n = (\xi_n^i)_{1 \leq i \leq N} \xrightarrow{G_n\text{-selection}} \widehat{\xi}_n = (\widehat{\xi}_n^i)_{1 \leq i \leq N} \xrightarrow{M_n\text{-mutation}} \xi_{n+1}$$

Key observations :

- ▶ $X_n = (X'_0, \dots, X'_n)$, $G_n(X_n) = G'(X'_n) \Rightarrow$ **Genealogical tree model**
- ▶ Law(An ancestral line | acceptance/selection $p = 0, \dots, n) = \mathbb{Q}_n$

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

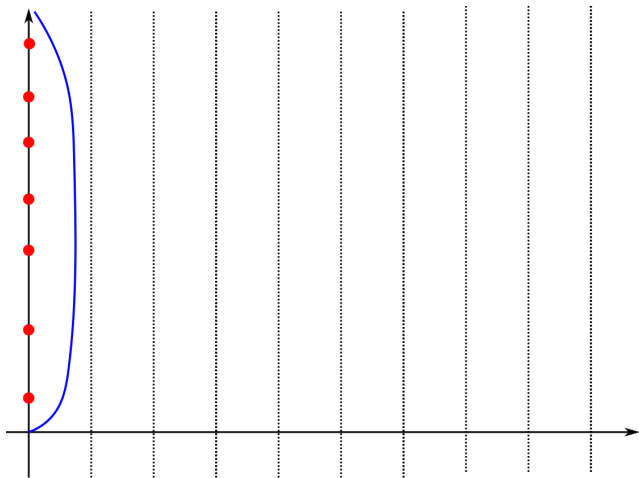


Some particle estimates

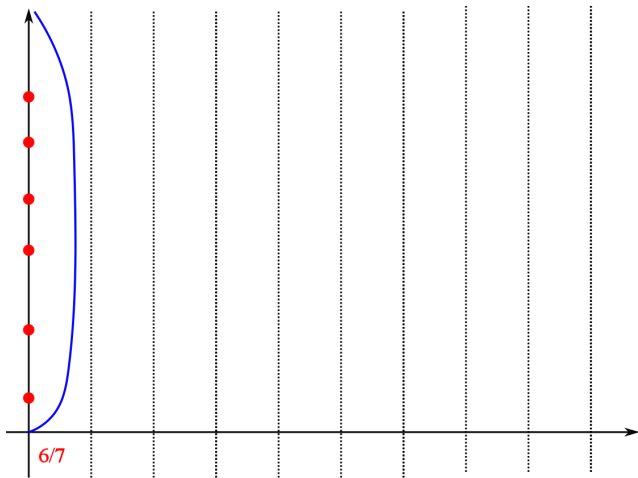
- ▶ Individuals ξ_n^i "almost" iid with law $\eta_n \simeq \eta_n^N = \frac{1}{N} \sum_{1 \leq i \leq N} \delta_{\xi_n^i}$
- ▶ Ancestral lines "almost" iid with law $\mathbb{Q}_n \simeq \frac{1}{N} \sum_{1 \leq i \leq N} \delta_{\text{Line}_n(i)}$
- ▶ **Unbiased** Particle normalizing constants

$$\mathcal{Z}_{n+1} = \prod_{0 \leq p \leq n} \eta_p(G_p) \simeq_{N \uparrow \infty} \mathcal{Z}_{n+1}^N = \prod_{0 \leq p \leq n} \eta_p^N(G_p) \quad (\text{or } \prod \text{ success proportions})$$

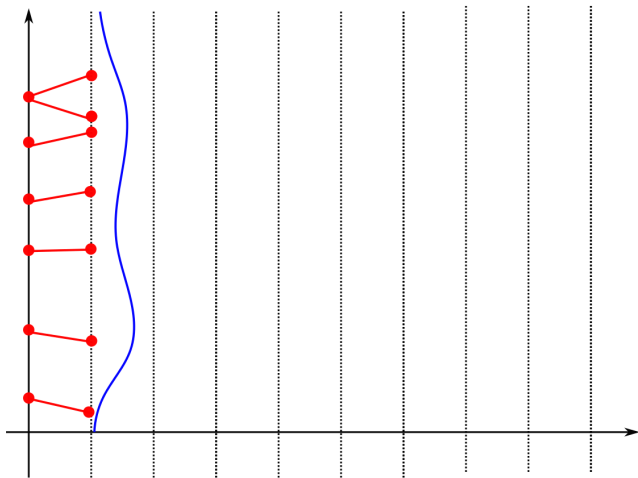
Graphical illustration : $\eta_n \simeq \eta_n^N := \frac{1}{N} \sum_{1 \leq i \leq N} \delta_{\xi_n^i}$



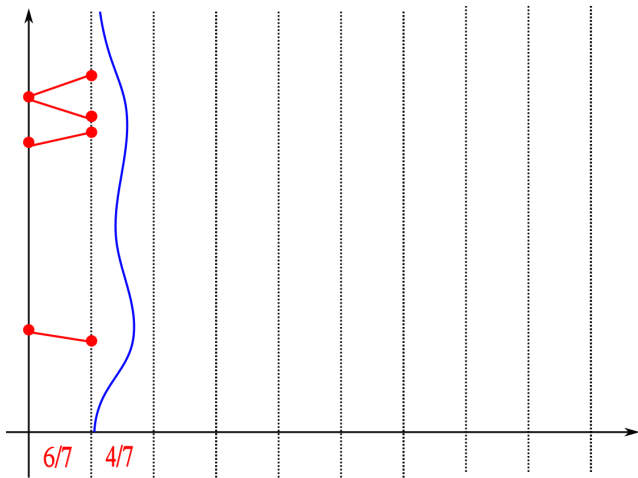
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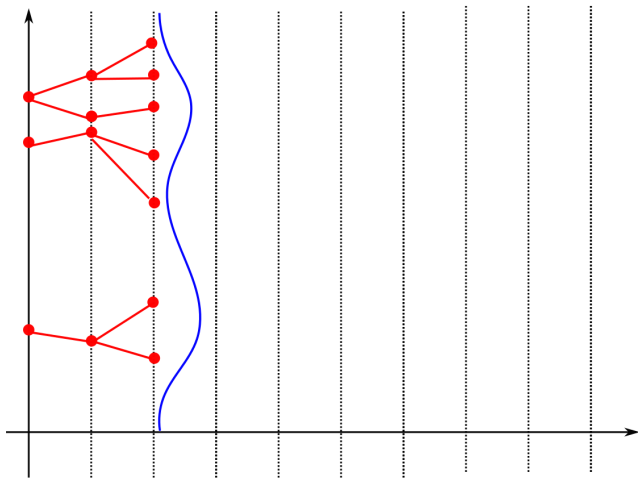
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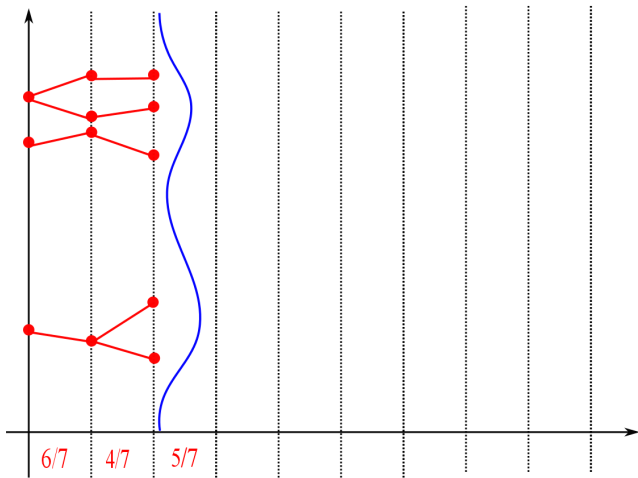
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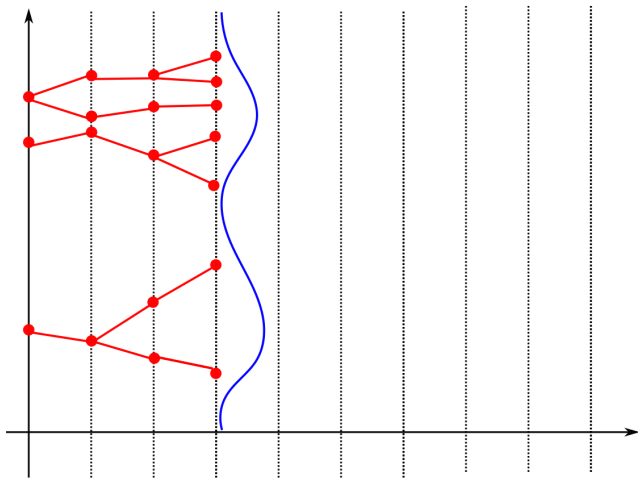
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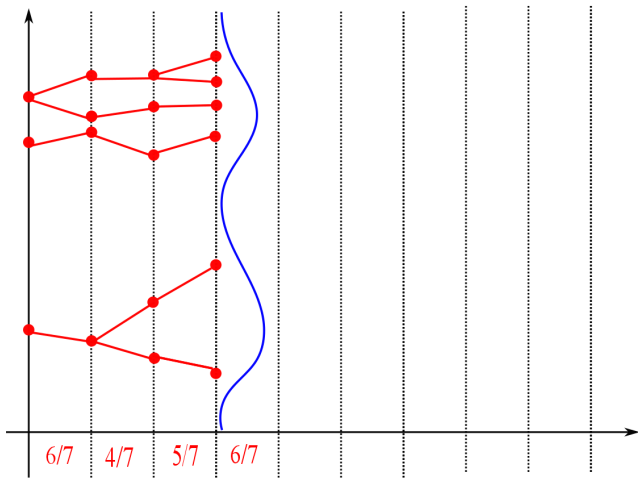
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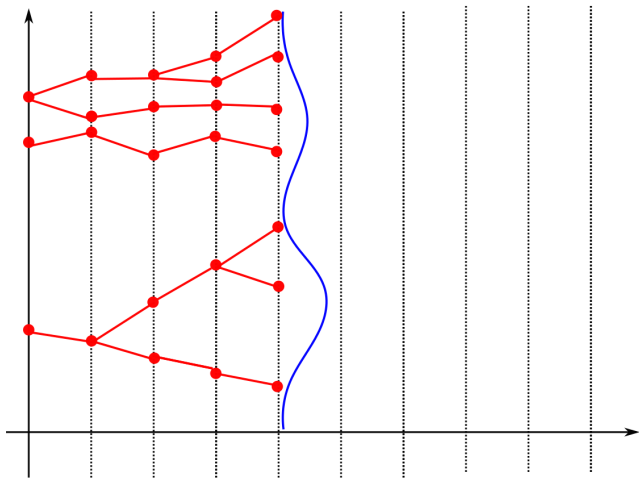
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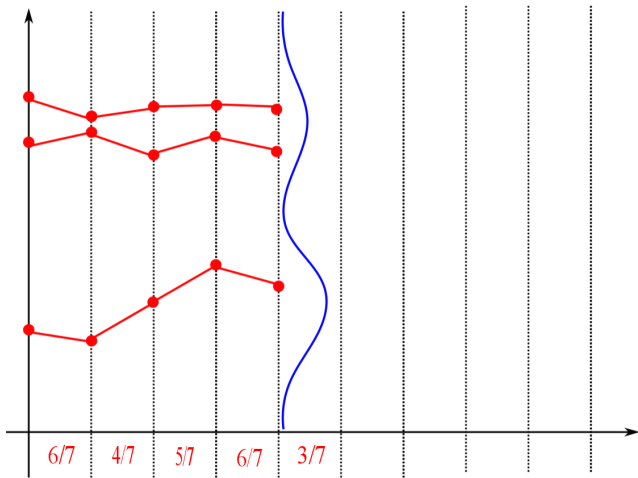
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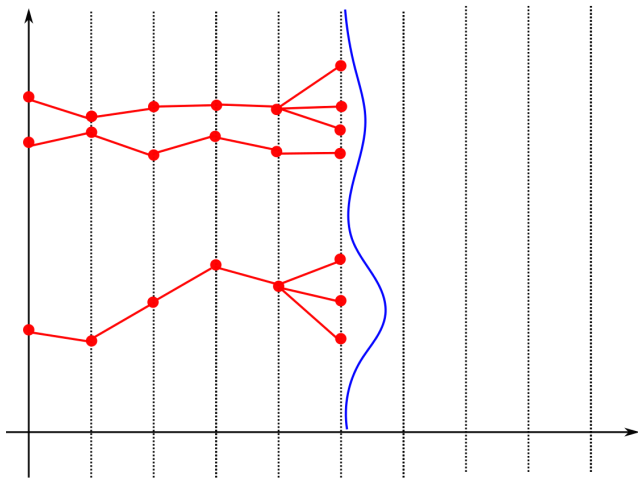
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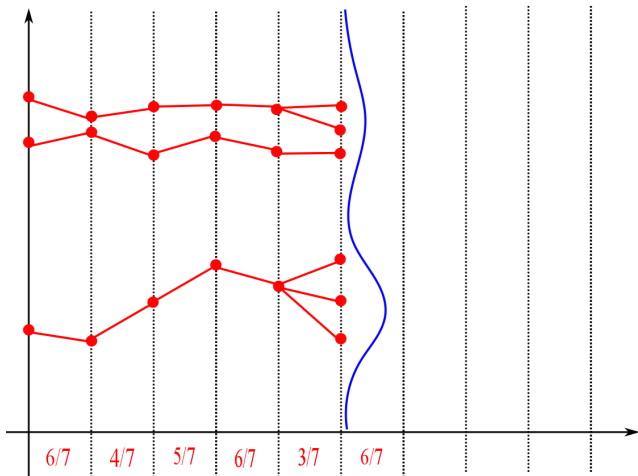
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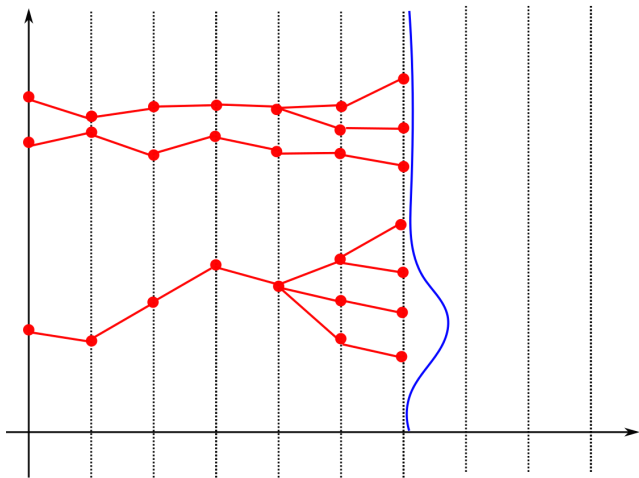
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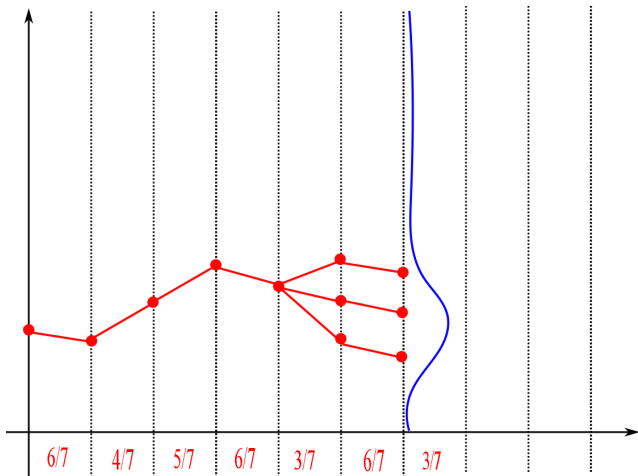
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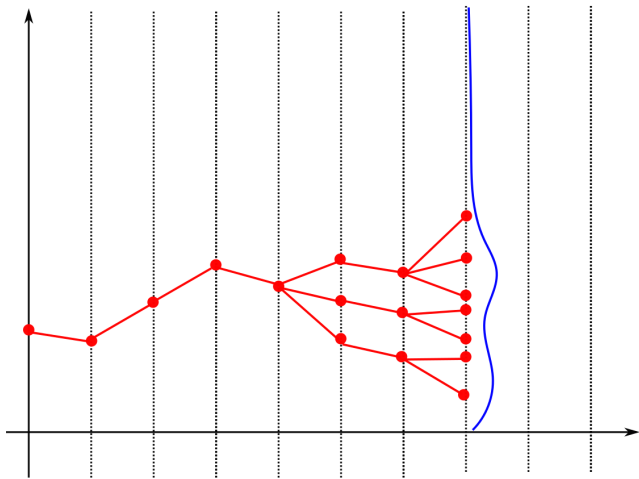
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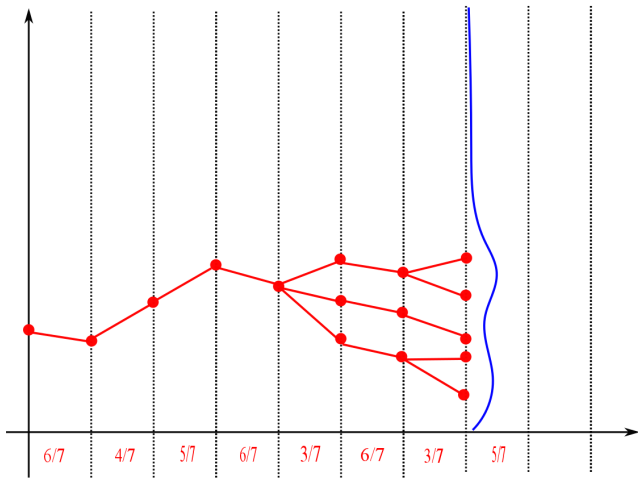
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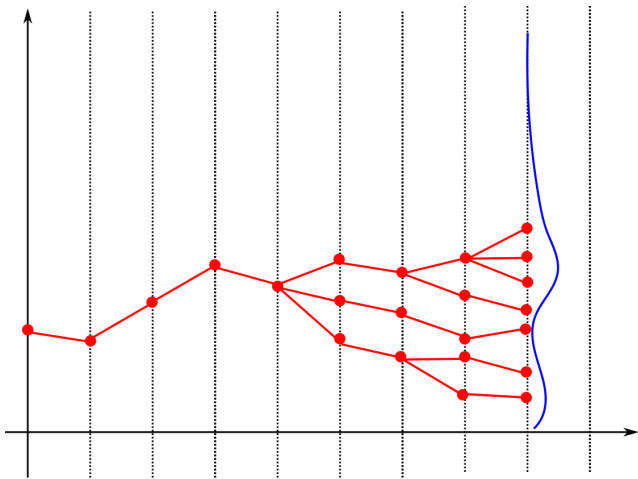
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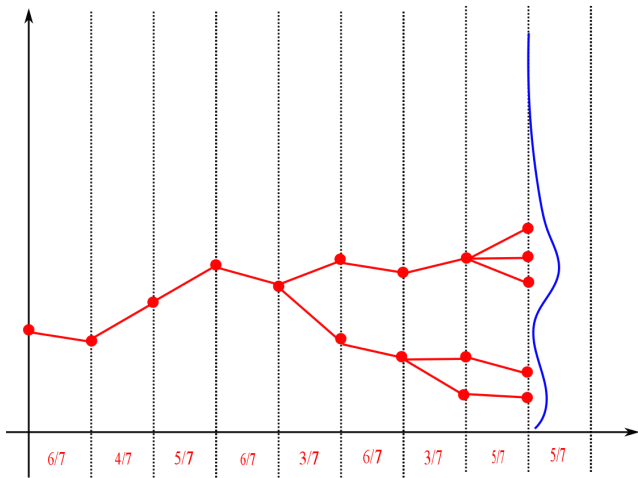
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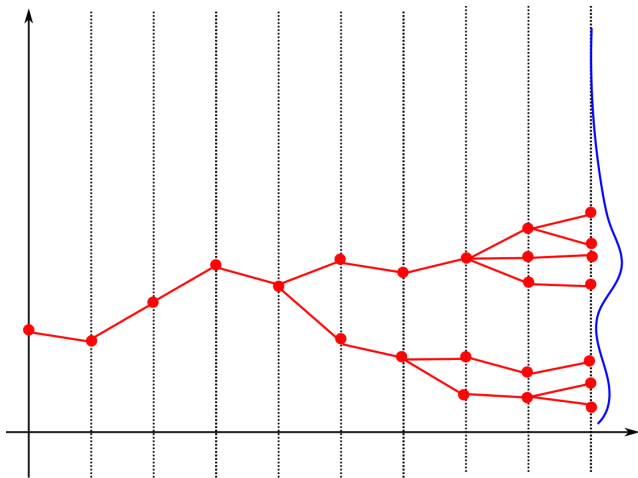
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How to use the full ancestral tree model ?

$$G_{n-1}(x_{n-1})M_n(x_{n-1}, dx_n) \stackrel{\text{hyp}}{=} H_n(x_{n-1}, x_n) \nu_n(dx_n)$$

⇒ **Backward Markov model :**

$$\begin{aligned} \mathbb{Q}_n(d(x_0, \dots, x_n)) &= \eta_n(dx_n) \underbrace{\mathbb{M}_{n, \eta_{n-1}}(x_n, dx_{n-1})}_{\propto \eta_{n-1}(dx_{n-1}) H_n(x_{n-1}, x_n)} \dots \mathbb{M}_{1, \eta_0}(x_1, dx_0) \end{aligned}$$

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Particle approximation

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

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

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

Introduction

Feynman-Kac models

7 rare event models/problems

Self avoiding walks

Level crossing probabilities

Particle absorption models

Quasi-invariant measures

Doob h -processes

Boltzmann-Gibbs measures

Restriction models

Exponential concentration analysis

Self avoiding walks in \mathbb{Z}^d

Feynman-Kac model with

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



Conditional distributions

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

and

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

Level crossing probabilities (1)

$$\mathbb{P}(V_n(X_n) \geq a) \quad \text{or} \quad \mathbb{P}(X \text{ hits } A_n \text{ before } B)$$

- ▶ Level crossing at a fixed given time

$$\begin{aligned} \mathbb{P}(V_n(X_n) \geq a) &= \mathbb{E} \left(f_n(\mathbf{X}_n) e^{V_n(X_n)} \right) \\ &= \mathbb{E} \left(\mathbf{f}_n(\mathbf{X}_n) \prod_{0 \leq p < n} G_p(\mathbf{X}_p) \right) \end{aligned}$$

with

- ▶ The Markov chain on transition space

$$\mathbf{X}_n = (X_n, X_{n+1}) \quad \text{and} \quad G_n(\mathbf{X}_n) = \exp[V_{n+1}(X_{n+1}) - V_n(X_n)]$$

- ▶ The test functions

$$f_n(\mathbf{X}_n) = 1_{V_n(X_n) \geq a} e^{-V_n(X_n)}$$

Level crossing probabilities (2)

- **Excursion level crossing** $A_n \downarrow$, with B non critical recurrent subset.

$$\mathbb{P}(X \text{ hits } A_n \text{ before } B) = \mathbb{E} \left(\prod_{0 \leq p \leq n} 1_{A_p}(X_{T_p}) \right)$$

$$T_n := \inf \{p \geq T_{n-1} : X_p \in (A_n \cup B)\}$$

Feynman-Kac model

$$\mathbb{E} \left(\prod_{0 \leq p \leq n} 1_{A_p}(X_{T_p}) \right) = \mathbb{E} \left(\prod_{0 \leq p < n} G_p(\mathbf{X}_p) \right)$$

with

$$\mathbf{X}_n = (X_t)_{t \in [T_n, T_{n+1}]} \quad \& \quad G_n(\mathbf{X}_n) = 1_{A_{n+1}}(X_{T_{n+1}})$$

Absorption models (Ground state energies, extinction probabilities, trapping models)

- ▶ Sub-Markov semigroups

$$Q_n(x, dy) = G_{n-1}(x) M_n(x, dy) \rightsquigarrow E_n^c = E_n \cup \{c\}$$

- ▶ Absorbed Markov chain

$$X_n^c \in E_n^c \xrightarrow{\text{absorption} \sim (1-G_n)} \widehat{X}_n^c \xrightarrow{\text{exploration} \sim M_n} X_{n+1}^c$$

↓

$$\mathbb{Q}_n = \text{Law}((X_0^c, \dots, X_n^c) \mid T^{\text{absorption}} \geq n)$$

and

$$\mathcal{Z}_n = \text{Proba}(T^{\text{absorption}} \geq n)$$

Homogeneous models $(G_n, M_n) = (G, M)$

- ▶ Reversibility condition : $\mu(dx)M(x, dy) = \mu(dy)M(y, dx)$

$$\text{Proba} (T^{\text{absorption}} \geq n) \simeq \lambda^n$$

with $\lambda =$ top eigenvalue of

$$Q(x, dy) = G(x) M(x, dy)$$

- ▶ Frobenius theorem $\rightsquigarrow \exists! h$ s.t. $Q(h) = \lambda h$

- ▶ Quasi-invariant/Yaglom measure/F-K sg fixed point :

$$\mathbb{P}(X_n^c \in dx \mid T^{\text{absorption}} > n) \xrightarrow{n \uparrow \infty} \frac{1}{\mu(h)} h(x) \mu(dx)$$

- ▶ Doob h -process X^h :

$$M^h(x, dy) = \frac{1}{\lambda} h^{-1}(x) Q(x, dy) h(y) = \frac{Q(x, dy) h(y)}{Q(h)(x)} = \frac{M(x, dy) h(y)}{M(h)(x)}$$

Homogeneous models $(G_n, M_n) = (G, M)$

$$\mathbb{Q}_n(d(x_0, \dots, x_n)) \propto \mathbb{P}((X_0^h, \dots, X_n^h) \in d(x_0, \dots, x_n)) h^{-1}(x_n)$$

- ▶ *Invariant measure $\mu_h = \mu_h M^h$ & normalized additive functionals*

$$\bar{F}_n(x_0, \dots, x_n) = \frac{1}{n+1} \sum_{0 \leq p \leq n} f(x_p) \implies \mathbb{Q}_n(\bar{F}_n) \simeq_n \mu_h(f)$$

- ▶ *If $G = G^\theta$ depends on some $\theta \in \mathbb{R} \rightsquigarrow f := \frac{\partial}{\partial \theta} \log G^\theta$*

$$\frac{\partial}{\partial \theta} \log \lambda^\theta \simeq_n \frac{1}{n+1} \frac{\partial}{\partial \theta} \log \mathcal{Z}_{n+1}^\theta = \mathbb{Q}_n(\bar{F}_n)$$

NB : Similar expression when M^θ depends on some $\theta \in \mathbb{R}$.

Boltzmann-Gibbs measures (Stochastic optimization, free energies, bayesian inference)

$$\eta_n(dx) := \frac{1}{\mathcal{Z}_n} e^{-\beta_n V(x)} \lambda(dx) \quad \text{with } \beta_n \uparrow$$

- ▶ For any MCMC transition M_n with target η_n

$$\eta_n = \eta_n M_n$$

- ▶ Updating of the temperature parameter

$$\eta_{n+1} = \Psi_{G_n}(\eta_n) \quad \text{with } G_n = e^{-(\beta_{n+1}-\beta_n)V}$$

Proof : $e^{-\beta_{n+1}V} = e^{-(\beta_{n+1}-\beta_n)V} \times e^{-\beta_n V}$

Consequence :

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

and ($\beta_0 = 0$)

$$\lambda(e^{-\beta_n V}) = \mathcal{Z}_n = \prod_{0 \leq p < n} \eta_p(G_p)$$

Restriction models (Uncertainty propagations, tail probabilities, black box input/output)

$$\eta_n(dx) := \frac{1}{\mathcal{Z}_n} \mathbf{1}_{A_n}(x) \lambda(dx) \quad \text{with} \quad A_n \downarrow$$

- ▶ For any MCMC transition M_n with target η_n

$$\eta_n = \eta_n M_n$$

- ▶ Updating of the subset

$$\eta_{n+1} = \Psi_{G_n}(\eta_n) \quad \text{with} \quad G_n = \mathbf{1}_{A_{n+1}}$$

Proof : $\mathbf{1}_{A_{n+1}} = \mathbf{1}_{A_{n+1}} \times \mathbf{1}_{A_n}$

Consequence :

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

and $(\lambda(A_0) = 1)$

$$\lambda(A_n) = \mathcal{Z}_n = \prod_{0 \leq p < n} \eta_p(G_p)$$

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- A brief review on genetic style algorithms

- Current population models

- Particle free energy

- Genealogical tree models

- Backward particle models

Equivalent Algorithms

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

Equivalent Algorithms

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Sampling Algorithms	Transition proposals	Accept-reject-recycle

More "natural" botanical names:

bootstrapping, spawning, cloning, pruning, replenish, multi-level splitting, subset branching, enrichment, go with the winner, ...

Equivalent Algorithms

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

More "natural" botanical names:

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1950 \leq Heuristic style algo. \leq 1996 \leq Particle Feynman-Kac models

Convergence analysis \supset CLT, LDP, \mathbb{L}_p -estimates, Empirical processes convergence, Moderate deviations, Berry-Essen, Propagations of chaos, Exact weak expansions,

Concentration analysis = Exponential deviation proba. estimates

Current population models

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

$\forall (x \geq 0, n \geq 0, N \geq 1)$

- ▶ Unbias prop. (\rightsquigarrow Island models [cf. DM, Dubarry, Moulines CS2])

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

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

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

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 $:= \eta_n^N$ (in path space)

Constants (c_1, c_2) related to (bias, variance), c universal constant \perp time
 \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

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

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

Backward particle models

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

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

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

- ▶ $\mathbf{f}_{a,n}$ 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 $1 - e^{-x}$.