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Genealogical tree models and Nonlinear estimation

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\hookrightarrow *Feynman-Kac Formulae, Genealogical and Interacting Particle Systems with Applications, Springer NY. Series: Probability and Applications, (2004)*

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Evolutionary type models

Simple Genetic Branching Algo.	<i>Mutation</i> <i>Selection/Branching</i>
Metropolis-Hastings Algo.	<i>Proposal</i> <i>Acceptance/Rejection</i>
Sequential Monte Carlo methods	<i>Sampling</i> <i>Resampling (SIR)</i>
Filtering/Smoothing	<i>Prediction</i> <i>Updating/Correction</i>
Particle \in Absorbing Medium	<i>Evolution</i> <i>Killing/Creation/Anhiling</i>

Other Botanical Names: multi-level splitting (Khan-Harris 51), prune enrichment (Rosenbluth 1955), switching algo. (Magill 65), matrix reconfiguration (Hetherington 84), restart (Villen-Altamirano 91), particle filters (Rigal-Salut-DM 92), SIR filters (Gordon-Salmon-Smith 93, Kitagawa 96), go-with-the-winner (Vazirani-Aldous 94), ensemble Kalman-filters (Evensen 1994), quantum Monte Carlo methods (Melik-Nightingale 1999), spawning filters (Fisher-Maybeck 2002), SIR Pilot Exploration Resampling (Liu-Zhang 2002),...

\iff Particle Interpretations of Feynman-Kac models

Since R. Feynman's PhD. on path integrals 1942

Physics \longleftrightarrow Biology \longleftrightarrow Engineering Sciences \longleftrightarrow Probability/Statistics

- **Physics :**

- $FKS \in$ nonlinear integro-diff. éq. (\sim generalized Boltzmann models).
- Spectral analysis of Schrödinger operators and large matrices with nonnegative entries.
(particle evolutions in disordered/absorbing media)
- Multiplicative Dirichlet problems with boundary conditions.
- Microscopic and macroscopic interacting particle interpretations.

- **Biology:**

- Self-avoiding walks, macromolecular polymerizations.
- Branching and genetic population models.
- Coalescent and Genealogical evolutions.

- **Rare events analysis:**
 - Multisplitting and branching particle models (Restart).
 - Importance sampling and twisted probability measures.
 - Genealogical tree based simulation methods.
- **Advanced Signal processing:**
 - Optimal filtering/smoothing/regulation, open loop optimal control.
 - Interacting Kalman-Bucy filters.
 - Stochastic and adaptative grid approximation-models
- **Statistics/Probability:**
 - Restricted Markov chains (w.r.t terminal values, visiting regions,...)
 - Analysis of Boltzmann-Gibbs type distributions (simulation, partition functions,...).
 - Random search evolutionary algorithms, interacting Metropolis/simulated annealing algo.

Simple Genetic evolution/simulation models —→ only 2 ingredients!!

(Discrete time parameter $n \in \mathbb{N} = \{0, 1, 2, \dots\}$, state spaces E_n ($\in \{\mathbb{Z}^d, \mathbb{R}^d, \underbrace{\mathbb{R}^d \times \dots \times \mathbb{R}^d}_{(n+1)-times}, \dots\}$))

- *Mutation/exploration/prediction/proposal :*
→ Markov transitions $M_n(x_{n-1}, dx_n)$ from E_{n-1} into E_n .
- *Selection/absorption/updating/acceptance :*
→ Potential functions G_n from E_n into $[0, 1]$.

A Genetic Evolution Model \Rightarrow Markov chain $\xi_n = (\xi_n^1, \dots, \xi_n^N) \in E_n^N = \underbrace{E_n \times \dots \times E_n}_{N-times}$

$$\xi_n \in E_n^N \xrightarrow{\text{selection}} \widehat{\xi}_n \in E_n^N \xrightarrow{\text{mutation}} \xi_{n+1} \in E_{n+1}^N$$

- **Selection transition** ($\exists \neq$ types \rightarrow Ex.: accept/reject)

$$\xi_n^i \rightsquigarrow \widehat{\xi}_n^i = \xi_n^i \quad \text{with proba. } G_n(\xi_n^i) \quad \boxed{\text{[Acceptance]}}$$

Otherwise we select a better fitted individual in the current configuration

$$\widehat{\xi}_n^i = \xi_n^j \quad \text{with proba. } G_n(\xi_n^j) / \sum_{k=1}^N G_n(\xi_n^k) \quad \boxed{\text{[Rejection + Selection]}}$$

- **Mutation transition**

$$\widehat{\xi}_n^i \rightsquigarrow \xi_{n+1}^i \sim M_{n+1}(\widehat{\xi}_n^i, \bullet)$$

A Genealogical tree model

Important observation [Historical process]

$$X'_n \in E'_n \quad \text{Markov chain}$$



$$X_n = (X'_0, \dots, X'_n) \in E_n = (E'_0 \times \dots \times E'_n) \quad \text{Markov chain} \in \text{path spaces}$$

→ *Markov transitions* $M_n(x_{n-1}, dx_n)$ [elementary extensions]

$$X_{n+1} = ((X'_0, \dots, X'_n), X'_{n+1}) = (X_n, X'_{n+1})$$

Genetic Evolution Model on Path Spaces=Genealogical tree model

$$X_n = (X'_0, \dots, X'_n) \quad \text{Markov transitions} \quad M_n \quad \text{and} \quad G_n(X_n) = G'_n(X'_n)$$

↓

Genetic path-valued particle Model

$$\left\{ \begin{array}{lcl} \xi_n^i & = & (\xi_{0,n}^i, \xi_{1,n}^i, \dots, \xi_{n,n}^i) \\ \widehat{\xi}_n^i & = & (\widehat{\xi}_{0,n}^i, \widehat{\xi}_{1,n}^i, \dots, \widehat{\xi}_{n,n}^i) \in E_n = (E'_0 \times \dots \times E'_n) \end{array} \right.$$

- Path acceptance/(rejection+selection).
- Path mutation = path elementary extensions.

Occupation/Empirical measures ($\forall f_n$ test function on E_n)

$$\eta_n^N(f_n) = \frac{1}{N} \sum_{i=1}^N f_n(\xi_n^i) = \frac{1}{N} \sum_{i=1}^N f_n \underbrace{(\xi_{0,n}^i, \xi_{1,n}^i, \dots, \xi_{n,n}^i)}_{i\text{-th ancestral lines}}$$

\downarrow
Unbias-particle measures & Unnormalized Feynman-Kac measures :

$$\gamma_n^N(f_n) = \eta_n^N(f_n) \times \prod_{0 \leq p < n} \eta_p^N(G_p) \xrightarrow{N \rightarrow \infty} \gamma_n(f_n) = \mathbb{E}(f_n(X_n) \prod_{0 \leq p < n} G_p(X_p))$$

Notes:

- $f_n = 1 \Rightarrow \gamma_n^N(1) = \prod_{0 \leq p < n} \eta_p^N(G_p) \xrightarrow{N \rightarrow \infty} \gamma_n(1) = \mathbb{E}(\prod_{0 \leq p < n} G_p(X_p))$
- *Path-space models*

$$[X_n = (X'_0, \dots, X'_n) \text{ and } G_n(X_n) = G'_n(X'_n)] \Rightarrow \gamma_n(f_n) = \mathbb{E}(f_n(X'_0, \dots, X'_n) \prod_{0 \leq p < n} G'_p(X'_p))$$

\Rightarrow Occupation measure & Normalized Feynman-Kac measures:

$$\eta_n^N(f_n) = \frac{1}{N} \sum_{i=1}^N f_n(\xi_n^i) = \gamma_n^N(f_n)/\gamma_n^N(1) \xrightarrow{N \rightarrow \infty} \eta_n(f_n) = \gamma_n(f_n)/\gamma_n(1)$$

Path-space models

$$[X_n = (X'_0, \dots, X'_n) \text{ and } G_n(X_n) = G'_n(X'_n)]$$

\Downarrow

$$\eta_n(f_n) = \frac{\mathbb{E}(f_n(X'_0, \dots, X'_n) \prod_{0 \leq p < n} G'_p(X'_p))}{\mathbb{E}(\prod_{0 \leq p < n} G'_p(X'_p))}$$

Note:

$$\gamma_n(f_n) = \eta_n(f_n) \times \prod_{0 \leq p < n} \eta_p(G_p) \quad (\leftarrow \gamma_n^N(f_n) = \eta_n^N(f_n) \times \prod_{0 \leq p < n} \eta_p^N(G_p))$$

Complete asymptotic theory $(n, N) \rightarrow \infty$ (weak+strong LLN, CLT, LDP,...)

→ *Feynman-Kac Formulae, Genealogical and Interacting Particle Systems with Applications, Springer NY. Series: Probability and Applications (2004)*

Some examples:

- Weak convergence [$p \geq 1$ + \mathcal{F}_n not too large + regular mutations]

$$\sup_{n \geq 0} \mathbb{E} \left(\sup_{f_n \in \mathcal{F}_n} |\eta_n^N(f_n) - \eta_n(f_n)|^p \right)^{1/p} \leq c(p)/\sqrt{N}$$

Ex.:

$$E_n = \mathbb{R}, \quad \mathcal{F}_n = \{\mathbf{1}_{]-\infty, x]}; \quad x \in \mathbb{R}\} \Rightarrow \sup_{n \geq 0} \mathbb{E} \left(\sup_{x \in \mathbb{R}} |\eta_n^N(\mathbf{1}_{]-\infty, x]}) - \eta_n(\mathbf{1}_{]-\infty, x}]|^p \right)^{1/p} \leq c(p)/\sqrt{N}$$

- Propagation-of-chaos estimates [$q \leq N$ finite block size, regular mutations]

$$\text{Law}(\xi_n^1, \dots, \xi_n^q) \simeq \eta_n^{\otimes q} + \frac{1}{N} \mathcal{M}_n^{(q)} \quad \text{with} \quad \mathcal{M}_n^{(q)} \quad \text{signed meas. s.t. } \sup_{n \geq 0} \sup_{\|F\| \leq 1} |\mathcal{M}_n^{(q)}(F)| \leq c q^2$$

Particle physics: Markov $X_n \in \mathbf{Absorbing\ medium}$ $G(x) = e^{-V(x)} \in [0, 1]$

$$X_n^c \in E^c = E \cup \{c\} \xrightarrow{\text{absorption}} \widehat{X}_n^c \xrightarrow{\text{exploration}} X_{n+1}^c$$

Absorption/killing: $\longrightarrow \widehat{X}_n^c = X_n^c$, with proba $G(X_n^c)$; otherwise the particle is killed and $\widehat{X}_n^c = c$.

\Downarrow

$$A = \{x : G(x) = 0\} \longrightarrow \text{Hard obstacles}$$

$$T = \inf \{n \geq 0 ; \widehat{X}_n^c = c\} \longrightarrow \text{Absorption time } X_{T+n}^c = \widehat{X}_{T+n}^c = c$$

$\implies \underline{\text{Feynman-Kac models}} (G, X_n) : \gamma_n = \text{Law}(X_n^c ; T \geq n) \quad \text{and} \quad \gamma_n(1) = \text{Proba}(T \geq n)$

\Downarrow

$$\eta_n = \text{Law}(X_n^c \mid T \geq n) = \text{Law}((X_0'^c, \dots, X_n'^c) \mid T \geq n)$$

Biology: Macromolecules and Directed Polymers

- *Self avoiding walks* $X'_n \in \mathbb{Z}^d$

$$X_n = (X'_0, \dots, X'_n) \quad \text{and} \quad G_n(X_n) = 1_{\notin \{X'_0, \dots, X'_{n-1}\}}(X'_n)$$

$$\gamma_n(1) = \text{Proba}(\forall 0 \leq p \neq q \leq n, \ X'_p \neq X'_q) \quad \text{and} \quad \eta_n = \text{Law}(X'_0, \dots, X'_n \mid \forall 0 \leq p \neq q \leq n, \ X'_p \neq X'_q)$$

- *Edwards' model*

$$X_n = (X'_0, \dots, X'_n) \quad \text{and} \quad G_n(X_n) = \exp \left\{ -\beta \sum_{0 \leq p < n} 1_{X'_p}(X'_n) \right\}$$

Statistics: Sequential MCMC and Feynman-Kac-Metropolis models

Metropolis potential [π target measure] + [(K, L) pair Markov transitions]

$$G(y_1, y_2) = \frac{\pi(dy_2)L(y_2, dy_1)}{\pi(dy_1)K(y_1, dy_2)}$$

Ex. π Gibbs measure:

$$\pi(dy) \propto e^{-V(y)} \lambda(dy) \Rightarrow G(y_1, y_2) = e^{(V(y_1) - V(y_2))} \frac{\lambda(dy_2)L(y_2, dy_1)}{\lambda(dy_1)K(y_1, dy_2)}$$

Note: $(K = L \text{ } \lambda\text{-reversible}) \text{ or } (\lambda K = \lambda \text{ and } L(y_2, dy_1) = \lambda(dy_1) \frac{dK(y_1, \cdot)}{d\lambda}(y_2))$

$$\Downarrow$$
$$G(y_1, y_2) = \exp(V(y_1) - V(y_2))$$

Notation $\mathbb{E}_\nu^M(\cdot)$ = Expectation w.r.t. Markov [transition M , initial condition ν]

Theorem: (Time reversal formula), [A. Doucet, P.DM; (Séminaire Probab. 2003)]

$$\mathbb{E}_\pi^L(f_n(Y_n, Y_{n-1}, \dots, Y_0) | Y_n = y) = \frac{\mathbb{E}_y^K(f_n(Y_0, Y_1, \dots, Y_n) \{ \prod_{0 \leq p < n} G(Y_p, Y_{p+1}) \})}{\mathbb{E}_y^K(\{ \prod_{0 \leq p < n} G(Y_p, Y_{p+1}) \})}$$

In addition :

⊕ FK-Metropolis n -marginal: $\lim_{n \rightarrow \infty} \eta_n = \pi$ (cv. decays $\perp \pi$)

⊕ Nonhomogeneous models: (π_n, L_n, K_n)

$\pi_n(dy) \propto e^{-\beta_n V(y)} \lambda(dy)$, cooling schedule $\beta_n \uparrow \infty$, mutation s.t. $\pi_n = \pi_n K_n$, and $\text{Law}(X_0) = \pi_0$

↓

$$G_n(y_1, y_2) = \exp [-(\beta_{n+1} - \beta_n)V(y_1)] \implies \eta_n = \pi_n$$

Rare events analysis

- **Importance sampling and Twisted Feynman-Kac measures**

$$\mathbb{P}(V_n(X_n) \geq a) = \mathbb{E}(\mathbf{1}_{V_n(X_n) \geq a} e^{-\beta_n V_n(X_n)} e^{+\beta_n V_n(X_n)})$$

↓

Importance potentials/measures:

$$G_n(X_n, X_{n+1}) = e^{\beta_n(V_{n+1}(X_{n+1}) - V_n(X_n))} \implies \mathbb{P}(V_n(X_n) \geq a) = \gamma_n(\mathbf{1}_{V_n \geq a} e^{-\beta_n V_n})$$

In addition:

$$\mathbb{E}(f_n(X_n) \mid V_n(X_n) \geq a) = \eta_n(f_n \mathbf{1}_{V_n \geq a} e^{-\beta_n V_n}) / \eta_n(\mathbf{1}_{V_n \geq a} e^{-\beta_n V_n})$$

⊕ Path-space models ⇒ weighted genealogies

$$X_n = (X'_0, \dots, X'_n) \text{ and } V_n(X_n) = V'_n(X'^n)$$

↓

$$\mathbb{E}(f_n(X'_0, \dots, X'_n) \mid V'_n(X'_n) \geq a) = \eta_n(f_n \mathbf{1}_{V_n \geq a} e^{-\beta_n V_n}) / \eta_n(\mathbf{1}_{V_n \geq a} e^{-\beta_n V_n})$$

- **Multi-splitting Feynman-Kac models** (\neq importance sampling)

$(E = A \cup A^c)$, Y_n Markov, $Y_0 \in A_0 (\subset A) \rightsquigarrow A^c = (B \cup C)$, C = absorbing set/hard obstacle

Multi-level decomposition $B = B_m \subset \dots \subset B_1 \subset B_0 \quad (A_0 = B_1 - B_0, \ B_0 \cap C = \emptyset)$

\Downarrow

$$\mathbb{P}(Y_n \text{ hits } B \text{ before } C) = \mathbb{E} \left(\prod_{1 \leq p \leq m} G_p(X_p) \right)$$

Inter-level excursions : $T_n = \inf \{p \geq T_{n-1} : Y_p \in B_n \cup C\}$

$$X_n = (Y_p ; T_{n-1} \leq p \leq T_n) \in \text{Excursion space} \quad G_n(X_n) = 1_{B_n}(Y_{T_n})$$

\Downarrow

FK interpretation

$$\mathbb{E}(f(Y_0, \dots, Y_{T_m}) 1_{B_m}(X_{T_m})) = \mathbb{E}(f(X_0, \dots, X_m) \prod_{1 \leq p \leq m} G_p(X_p))$$

Advanced signal processing → filtering/hidden Markov chains/Bayesian methodology

Signal process

$X_n = \text{Markov chain} \in E_n$

Observation/Sensor eq. $Y_n = H_n(X_n, V_n) \in F_n$ with $\mathbb{P}(H_n(x_n, V_n) \in dy_n) = g_n(x_n, y_n) \lambda_n(dy_n)$

Example: $Y_n = h_n(X_n) + V_n \in F_n = \mathbb{R}$, with Gaussian noise $V_n = \mathcal{N}(0, 1)$

⇓

$$\mathbb{P}(h_n(x_n) + V_n \in dy_n) = (2\pi)^{-1/2} e^{-\frac{1}{2}(y_n - h_n(x_n))^2} \quad dy_n = \underbrace{\exp [h_n(x_n)y_n - h_n^2(x_n)/2]}_{g_n(x_n, y_n)} \underbrace{\mathcal{N}(0, 1)(dy_n)}_{\lambda_n(dy_n)}$$

Prediction/filtering/smoothing → **Feynman-Kac representation** $G_n(x_n) = g_n(x_n, y_n)$

$$\eta_n = \text{Law}(X_n \mid Y_0 = y_0, \dots, Y_{n-1} = y_{n-1}) = \text{Law}(X'_0, \dots, X'_n \mid Y_0 = y_0, \dots, Y_{n-1} = y_{n-1})$$

Partially linear/Gaussian models

$$X_n^1 = \text{Markov } \in E_n + \begin{cases} X_n^2 &= A_n(X_n^1) X_{n-1}^2 + a_n(X_n^1) + B_n(X_n^1) W_n \in \mathbb{R}^d \\ Y_n &= C_n(X_n^1) X_n^2 + c_n(X_n^1) + D_n(X_n^1) V_n \in \mathbb{R}^{d'} \end{cases}$$

Given a realization $X^1 = x \rightarrow \text{Kalman-Bucy optimal one step predictor}$

$$\widehat{X}_{x,n+1}^2 = \mathbb{E}(X_{n+1}^2 \mid Y_0, \dots, Y_n, X^1 = x) \quad \text{and} \quad P_{x,n+1}^- = \mathbb{E}([X_{n+1}^2 - \widehat{X}_{x,n+1}^2][X_{n+1}^2 - \widehat{X}_{x,n+1}^2]')$$

↓

Quenched Kalman-Bucy recursion: $(\widehat{X}_{x,n+1}^2, P_{x,n+1}^-) = \mathcal{B}_{n+1}[(x_n, x_{n+1}), (\widehat{X}_{x,n}^2, P_{x,n}^-)]$

Feynman-Kac representation: $\eta_n \sim (\mathbf{X}_n, \mathbf{G}_n)$ s.t.

$$\begin{aligned}\mathbf{X}_n &= (X_n^1, (\widehat{X}_{X^1,n+1}^2, P_{X^1,n+1}^-)) \text{ Markov chain } \in \mathbf{E}_n = (E_n \times \mathbb{R}^d \times \mathbb{R}^{d \times d}) \\ \mathbf{G}_n(x, m, P) &= \frac{d\mathcal{N}(C_n(x) m + c_n(x), C_n(x) P C_n(x)' + D_n(x) R_n^v D_n(x)')}{d\mathcal{N}(0, D_n(x) R_n^v D_n(x)')} (y_n)\end{aligned}$$

$$\Downarrow \quad [\text{virtual sensor} : \quad Y_n = \{C_n(X_n^1) \widehat{X}_{X^1,n}^2 + c_n(X_n^1)\} + \widehat{V}_{X^1,n}]$$

$$\begin{aligned}F_n(x, m, P) = f_n(x) &\implies \eta_n(F_n) = \mathbb{E}(f_n(X_n^1) \mid Y_0, \dots, Y_{n-1}) \\ F_n(x, m, P) = \mathcal{N}(m, P)(f_n) &\implies \eta_n(F_n) = \mathbb{E}(f_n(X_n^2) \mid Y_0, \dots, Y_{n-1})\end{aligned}$$

Note: \rightsquigarrow Interacting Kalman-Bucy filters and for path-space models we have

$$X_n^1 = (X_0^1 ', \dots, X_n^1 ') \rightsquigarrow \text{Law}((X_0^1 ', \dots, X_n^1 ') \mid Y_0, \dots, Y_{n-1})$$