

# Sequential Monte Carlo methods

## Lecture 15 – General Sequential Monte Carlo

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**Aim:** Show how SMC can be used for a much wider class of problems than inference in state-space models.

## Outline:

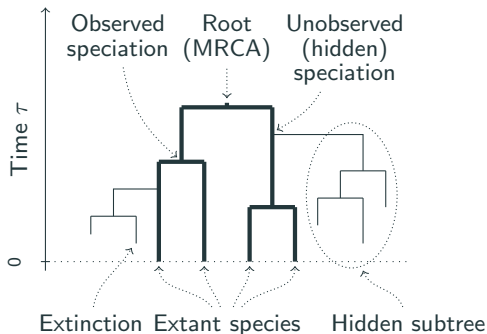
1. Examples of probabilistic models
2. General SMC formulation
3. Locally optimal proposals

## Examples of probabilistic models

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# Phylogenetic trees

A phylogenetic (evolutionary) tree shows the inferred evolutionary relationships among various species based upon similarities and differences in their physical or genetic characteristics.



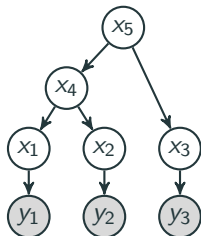
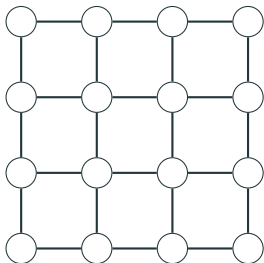
$$p(\mathcal{T}_r) = \begin{cases} 2 \prod_{c \in \text{Ch}(r)} p(\mathcal{T}_c) & \text{if } r \text{ is the root node,} \\ 2\lambda e^{-(\lambda+\mu)\Delta_r} \prod_{c \in \text{Ch}(r)} p(\mathcal{T}_c) & \text{if } r \text{ is a speciation,} \\ \mu e^{-(\lambda+\mu)\Delta_r} & \text{if } r \text{ is an extinction,} \\ e^{-(\lambda+\mu)\Delta_r} & \text{if } r \text{ is an extant species,} \end{cases}$$

(More on Lectures 17-18.)

# Probabilistic graphical models

A **probabilistic graphical model** (PGM) is a probabilistic model where a graph  $\mathcal{G} = (\mathcal{V}, \mathcal{E})$  represents the conditional independency structure between random variables,

1. a set of **vertices**  $\mathcal{V}$  (nodes) represents the random variables
2. a set of **edges**  $\mathcal{E}$  containing elements  $(i, j) \in \mathcal{E}$  connecting a pair of nodes  $(i, j) \in \mathcal{V} \times \mathcal{V}$



# Gaussian process state-space model

The Gaussian process (GP) is a **non-parametric** and **probabilistic** model for nonlinear functions.

**Non-parametric** means that it does not rely on any particular parametric functional form to be postulated.

$$\begin{aligned} X_t &= f(X_{t-1}) + V_t, & \text{s.t. } f(X) &\sim \mathcal{GP}(0, \kappa_{\eta, f}(x, x')), \\ Y_t &= g(X_t) + E_t, & \text{s.t. } g(X) &\sim \mathcal{GP}(0, \kappa_{\eta, g}(x, x')). \end{aligned}$$

The model functions  $f$  and  $g$  are assumed to be realizations from Gaussian process priors and  $V_t \sim \mathcal{N}(0, Q)$ ,  $E_t \sim \mathcal{N}(0, R)$ .

**Task:** Compute the posterior  $p(f, g, Q, R, \eta, x_{0:T} | y_{1:T})$ .

# Dirichlet process mixtures models

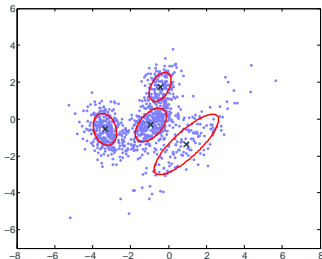
The **Dirichlet process** is a Bayesian nonparametric model, which can be used to construct mixture models with an **unknown** and **possibly infinite** number of components.

**Generative process:**

$$p(x_{k+1} = j \mid x_{1:k}) = \begin{cases} \frac{n_{k,j}}{k+\alpha} & \text{for } j = 1, \dots, J_k, \\ \frac{\alpha}{k+\alpha} & \text{for } j = J_k + 1, \end{cases},$$

$$\theta_k \sim F(\theta), \quad k = 1, 2, \dots,$$

$$p(y_k \mid x_k, \{\theta_k\}_{k \geq 1}) = G(y_k \mid \theta_{x_k}),$$



# General SMC formulation

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# Model specification

SMC can be used to approximate a **sequence** of probability distributions on a sequence of probability spaces of **increasing dimension**.

Let  $\{\pi_k(\mathbf{x}_{1:k})\}_{k \geq 1}$  be an arbitrary sequence of target distributions

$$\pi_k(\mathbf{x}_{1:k}) = \frac{\tilde{\pi}_k(\mathbf{x}_{1:k})}{Z_k}$$

- The domain of  $x_k$  is  $\mathcal{X}_k$ , and  $\mathcal{X}_{1:k} = \mathcal{X}_k \times \mathcal{X}_{1:k-1}$  for all  $k$ .
- $\tilde{\pi}_k(\mathbf{x}_{1:k})$  can be evaluated pointwise.
- The normalizing constant  $Z_k$  may be unknown.

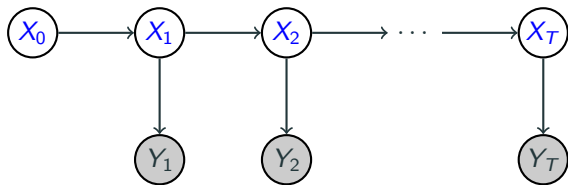
Common tasks:

1. Approximate the normalization constant  $Z_k$ .
2. Approximate  $\pi_k(\mathbf{x}_k)$  and compute  $\int \phi(\mathbf{x}_k) \pi_k(\mathbf{x}_k) d\mathbf{x}_k$ .

## ex) State space model

The sequence of target distributions  $\{\pi_k(\mathbf{x}_{0:k})\}_{k=1}^n$  can be constructed in **many** different ways.

The most basic construction arises from **chain-structured graphs**, such as the state space model (SSM).



$$\underbrace{p(\mathbf{x}_{0:t} | \mathbf{y}_{1:t})}_{\pi_t(\mathbf{x}_{0:t})} = \frac{\underbrace{p(\mathbf{x}_{0:t}, \mathbf{y}_{1:t})}_{\tilde{\pi}_t(\mathbf{x}_{0:t})}}{\underbrace{p(\mathbf{y}_{1:t})}_{Z_t = \int \tilde{\pi}(\mathbf{x}_{0:t}) d\mathbf{x}_{0:t}}}$$

Sequential Monte Carlo approximates

$$\pi_k(x_{0:k}) \approx \sum_{i=1}^N w_k^i \delta_{x_{0:k}^i}(x_{0:k}).$$

The weighted particle populations  $\{x_{0:k}^i, w_k^i\}_{i=1}^N$  are generated sequentially for  $k = 1, 2, \dots$

# General Sequential Monte Carlo



Assume that we have obtained  $\{x_{0:k-1}^i, w_{k-1}^i\}_{i=1}^N$

**Resampling:** Sample  $a_k^i$  with  $\mathbb{P}(a_k^i = j) = \nu_{k-1}^j, j = 1, \dots, N$ .

**Propagation:**  $x_k^i \sim q_k(x_k | x_{1:k-1}^{a_k^i})$  and  $x_{0:k}^i = (x_{0:k-1}^{a_k^i}, x_k^i)$ .

**Weighting:**  $w_k^i \propto \frac{w_{k-1}^{a_k^i}}{\nu_{k-1}^{a_k^i}} \frac{\tilde{\pi}_k(x_{0:k}^i)}{\tilde{\pi}_{k-1}(x_{0:k-1}^{a_k^i}) q_k(x_k^i | x_{0:k-1}^{a_k^i})}$ .

The result is a new weighted set of particles  $\{x_{0:k}^i, w_k^i\}_{i=1}^N$ .

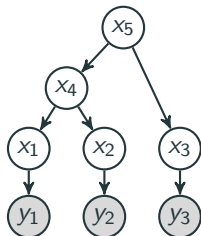
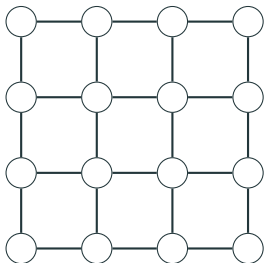
# SMC for probabilistic graphical models

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## Recall – Probabilistic graphical models

A **probabilistic graphical model** (PGM) is a probabilistic model where a graph  $\mathcal{G} = (\mathcal{V}, \mathcal{E})$  represents the conditional independency structure between random variables,

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## Key idea

SMC methods are used to approximate a **sequence of probability distributions** on a sequence of spaces of increasing dimension.

### Key idea:

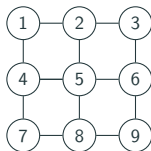
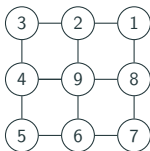
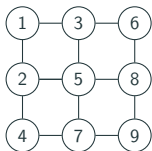
1. Introduce a **sequential decomposition** of the PGM.
2. Each **subgraph** induces an intermediate target dist.
3. Apply SMC to the sequence of intermediate target dist.

Using an artificial sequence of intermediate target distributions for an SMC method is a powerful (quite possibly underutilized) idea.

**Key question:** Exactly how do we define  $\{\tilde{\pi}_k(x_{1:k})\}_{k \geq 1}$ ?

## ex) Illustrating possible graph decomposition

Using a 2D lattice model from statistical physics,  $\mathbf{x} \in (-\pi, \pi]$ .



$$p(\mathbf{x}_{\mathcal{V}}) \propto e^{-\beta H(\mathbf{x}_{\mathcal{V}})}, \quad H(\mathbf{x}_{\mathcal{V}}) = - \sum_{(i,j) \in \mathcal{E}} J_{ij} \cos(x_i - x_j),$$

The **intermediate sequence** of target distributions can be chosen

$$\tilde{\pi}_k(\mathbf{x}_{\mathcal{L}_k}) \propto \tilde{\pi}_{k-1}(\mathbf{x}_{\mathcal{L}_{k-1}}) e^{\kappa(\mathbf{x}_{\mathcal{L}_{k-1}}) \cos(x_k - \mu(\mathbf{x}_{\mathcal{L}_{k-1}}))}.$$

$\mathcal{L}_k$  – index to the nodes in the  $k^{\text{th}}$  intermediate target  $\tilde{\pi}_k(\mathbf{x}_{\mathcal{L}_k})$ .



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## Algorithm SMC for graphical models

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### 1. Initialize ( $k = 1$ ):

- (a) Draw  $x_{\mathcal{L}_1}^i \sim q_1(\cdot)$ .
- (b) Set  $w_1^i = W_1(x_{\mathcal{L}_1}^i)$ .

### 2. For $k = 2$ to $K$ do:

- (a) **Resampling:** Draw  $a_k^i$ ,  $\mathbb{P}(a_k^i = j) = \tilde{w}_{k-1}^j / \sum_l \tilde{w}_{k-1}^l$ .
- (b) **Propagation:** Draw  $\xi_k^i \sim q_k(\cdot | x_{\mathcal{L}_{k-1}}^{a_k^i})$ , set  $x_{\mathcal{L}_k}^i = x_{\mathcal{L}_{k-1}}^{a_k^i} \cup \xi_k^i$ .
- (c) **Weighting:** Set  $\tilde{w}_k^i = W_k(x_{\mathcal{L}_k}^i)$ .

### 3. End

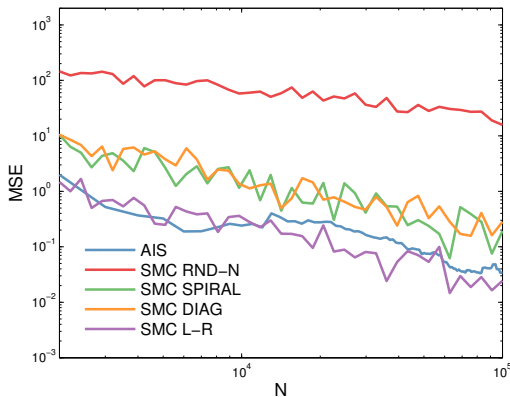
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$\mathcal{L}_k$  – index to the nodes in the  $k^{\text{th}}$  intermediate target  $\tilde{\pi}_k(x_{\mathcal{L}_k})$ .

$\xi_k^i$  – nodes added at step  $k$ .

Also provides an unbiased estimate of the **normalizing constant!**

## ex) Classical XY-model



This model is borrowed from



John M. Kosterlitz and David J. Thouless. **Ordering, metastability and phase transitions in two-dimensional systems.** *J. of Physics C: Solid State Physics*, 6(7):1181–1203, 1973.

# How to select the targets?

We want to sample from  $p(x_{\mathcal{Y}}) \Rightarrow$  **Require:**  $\pi_K(x_{\mathcal{L}_K}) = p(x_{\mathcal{Y}})$ , but...

... all **intermediate targets**  $\{\pi_k(x_{\mathcal{L}_k})\}_{k=1}^{K-1}$  are design choices!

**Idea:**

$$\tilde{\pi}_k^{\psi}(x_{\mathcal{L}_k}) = \underbrace{\tilde{\pi}_k(x_{\mathcal{L}_k})}_{\text{From graph decomp.}} \overbrace{\psi_k(x_{\mathcal{L}_k})}^{\text{Twisting funct.}}$$

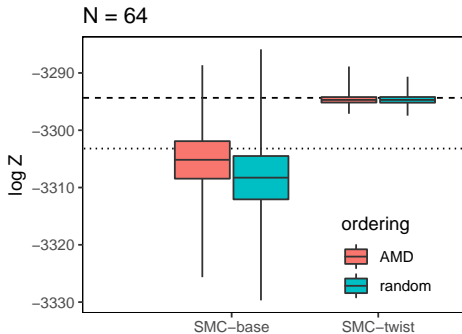
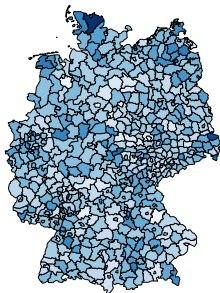
- In theory, possible to select  $\psi_k$  to get exact samples from  $p(x_{\mathcal{Y}})$  at iteration  $K$ .
- Use **deterministic inference method** to approximate optimal  $\psi_k$

SMC works as a **post-correction** of the **biases** associated with the deterministic inference method of choice.

## ex) CAR-Binomial

Model:

- Precision:  $Q_{tt} = 0.1 \times \{|\text{Ne}(t)| + 1\}$  and  $Q_{tt'} = -0.1$  if  $t \sim t'$
- $y_t \sim \text{Binomial}(10, \text{logit}^{-1}(x_t))$
- Spatial structure  $\sim$  regions in Germany,  $T = 544$



## What about stability?

**Recall:** for a state-space model we need exponential forgetting for the particle filter to be stable.

### **The same is true in the general case!**

If there are **strong** and **long-ranging** dependencies among the variables  $X_{1:k}$  under the distribution  $\pi_k$ , then the asymptotic variance of SMC may be exponential in  $k$ .

However,

- In many applications we *do* have fast enough forgetting (though, it can be difficult to verify theoretically)
- Even if this is not the case, SMC can give good results for moderate values of  $k$

- Bootstrap particle filter invented around 1992–1993
- Auxiliary particle filter, 1999
- Convergence theory: many results in the early 2000 but still an active research area
- SMC Samplers, 2006 (similar ideas going back to at least 2002)
- Particle Markov chain Monte Carlo, around 2010
- SMC for PPL, graphical models, etc. 2010–present