#### BACKWARD FILTERING FORWARD GUIDING

#### Frank van der Meulen (Vrije Universiteit Amsterdam)

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Algorithms and Computationally Intensive Inference Seminar (ACIIS) Warwick, December 2 (2022)

General problem setting

Conditioning, Doob's h-transform and the Backward Information Filter

Guided process

Discrete case

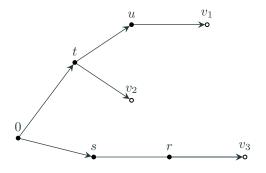
Numerical illustration

Continuous time transitions

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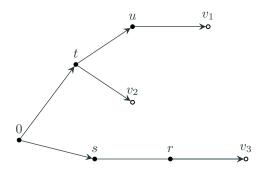
Wrap-up / conclusions

Consider a directed *Markovian* tree:



• latent vertices, o leaf/observation-vertices.

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To each edge corresponds a Markov kernel  $\kappa_{t}(x_{pa(t)}, dx_{t})$ .

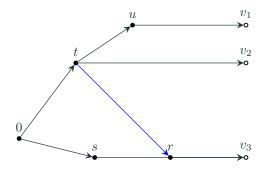
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#### We aim for

- 1. sampling values at ●, conditional on values at o;
- 2. estimating parameters in kernels;
- 3. not just on a tree, but on a general Directed Acyclic Graph (DAG).

# **Example of a DAG**



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- population of *n* individuals;
- each individual is either Susceptible, Infected or Recovered;
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- If  $x_i = \mathbf{I}$ , then it transitions to  $\mathbf{R}$  with intensity  $\mu$ .
- If  $x_i = \mathbf{R}$ , it transitions to  $\mathbf{S}$  with intensity  $\nu$ .

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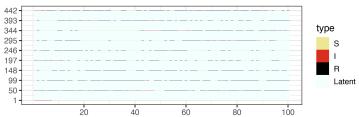
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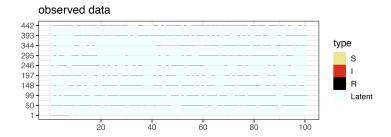
The transition matrix for individual i at time t, given "full state" x:

$$\kappa_i(t,x) = \begin{bmatrix} \psi(\lambda N_i(t,x)) & 1 - \psi(\lambda N_i(t,x)) & 0\\ 0 & \psi(\mu) & 1 - \psi(\mu)\\ 1 - \psi(\nu) & 0 & \psi(\nu) \end{bmatrix},$$

where  $\psi(u) = \exp(-\tau u)$ 

#### observed data





Observe state of each individual at times  $t_0 < t_1 < \cdots t_n$ .

#### Goals:

- identify most probable latent states (partial observations...);
- estimate rate parameters  $\lambda$ ,  $\mu$  and  $\nu$ .

 $\triangle$  Dimension of state-space is  $3^n$ .

Variation: observe

$$V_{t_i} \sim \text{Bin}(I(X_{t_i}), \rho),$$

with  $I(X_{t_i})$  the number of infected individuals at time  $t_i$ .

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Ju, Heng, Jacob - Sequential Monte Carlo algorithms for agent-based models of disease transmission

#### **Example 2: Wright-Fisher diffusion on a tree**

Diffusion approximation to Wright-Fisher model (with mutation) for  ${\cal N}$  diploid individuals.

Consider a directed tree where along each branch

$$dX_t = (\beta_1(1 - X_t) + \beta_2 X_t) dt + \sqrt{X_t(1 - X_t)} dW_t.$$

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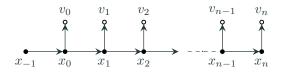
Wish to

- reconstruct  $X_t$  along edges;
- estimate parameters  $(\beta_1, \beta_2)$ .

STOLTZ ET AL. – Bayesian inference of species trees using diffusion models

General problem setting

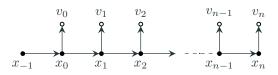
### State-space models / hidden Markov models



Well-known filtering, smoothing algorithms dating back to 1960-1970.

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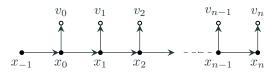


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- Finite state space: Baum-Welch, Viterbi, forward-backward algorithm.
- Linear Gaussian models: Kalman filter, Rauch-Tung-Striebel smoother.
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More recently much work on SMC (twisted particle samplers, controlled SMC).

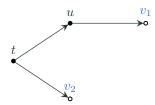
# Conditioning, Doob's

Information Filter

h-transform and the Backward

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- $V_t$ : all leaf descendants of vertex t.
- $V_t = \{v_1, v_2\}.$



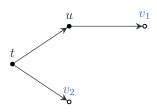
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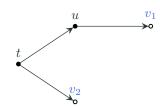
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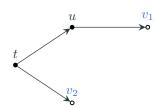


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 $\Lambda$  Within the subtree,  $h_t(x_t)$  is the likelihood of  $x_t$ .

• *Doob's* h-transform: Transformation of each  $\kappa_s$  with  $h_s$  to  $\kappa_s^*$ :

$$\kappa_{s}^{\star}(x, dy) = \frac{\kappa_{s}(x, dy)h_s(y)}{\int \kappa_{s}(x, dy)h_s(y)}, \quad s \in \mathcal{S}.$$

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- Recursive computation of  $h_s$  in a backward pass: (Backward Information Filter):
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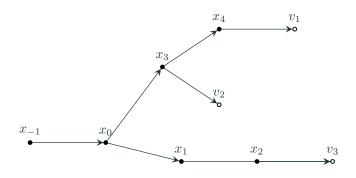
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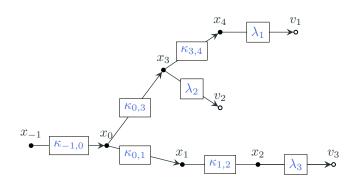
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  - Only in very specific models tractable.
- $\bigwedge$  On a DAG conditioning changes the dependency structure. There are no conditional kernels  $\kappa_{\Rightarrow s}^{\star}$  from pa(s) to s.

#### **Backward Information Filter**



# Make kernels explicit



## **Example:** finite state space

• Suppose  $x_t \in \{\textcircled{1}, \textcircled{2}, \textcircled{3}\}$  and  $v_t \in \{\textcircled{1}, \textcircled{2}, \textcircled{3}\}$ . Idea: in observations we cannot distinguish 1 and 2.

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- Finite state space ⇒ Markov kernels can be identified with matrices

$$\lambda_i = \begin{bmatrix} 1 & 0 \\ 1 & 0 \\ 0 & 1 \end{bmatrix} \qquad \kappa_{s,t} = \begin{bmatrix} 1 - \theta & \theta & 0 \\ 0.25 & 0.5 & 0.25 \\ 0.4 & 0.3 & 0.3 \end{bmatrix},$$

for  $i \in \{1, 2, 3\}$ ,  $s \in \{0, 1, 3\}$  and  $t \in \operatorname{ch}(s)$ .

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• Prior on initial state: set  $x_{-1} = 0$  and

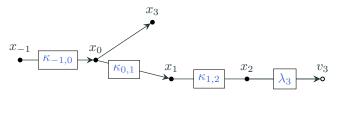
$$\kappa_{-1,0} = [\pi_1, \ \pi_2, \ \pi_3] =: \boldsymbol{\pi}.$$

• BIF: efficient way to compute  $x \mapsto h_t(x)$ .

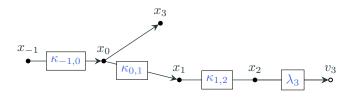
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- Initialise from observations: for t = 1, 2, 3

$$h_t^{\text{obs}} := \begin{bmatrix} 1 \\ 0 \end{bmatrix} \mathbf{1} \{ v_t = (1,2) \} + \begin{bmatrix} 0 \\ 1 \end{bmatrix} \mathbf{1} \{ v_t = (3) \}.$$

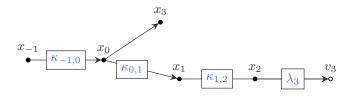


$$h_2 = \lambda_3 h_3^{\text{obs}}$$
  $h_1 = \kappa_{1,2} h_2.$ 



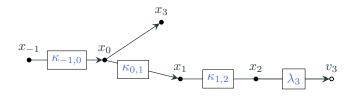
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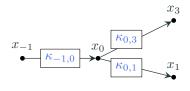
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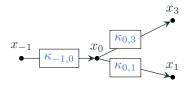
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$$= \sum_{x_2} \underbrace{p(v_3 \mid \cancel{x_1}, x_2)}_{h_2(x_2)} p(x_2 \mid x_1).$$



Get

$$h_{0 \Rightarrow 3} = \kappa_{0,3} h_3 \quad \text{and} \quad h_{0 \Rightarrow 1} = \kappa_{0,1} h_1$$

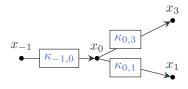


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$$h_0(x) = h_{0 \to 1}(x) h_{0 \to 3}(x).$$



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- ⚠ This is all tractable because
  - 1. the DAG is a directed tree;
  - 2. the state space is finite.

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Then in BIF, with  $h_{1,3}(y)=h_{0\rightarrow 1}(y_1)h_{0\rightarrow 3}(y_2)$ , we get

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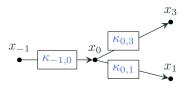
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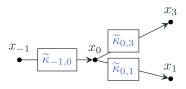
Natural to convert DAG to string diagram...

Key idea: replace  $h_{s o t}$  by  $g_{s o t}$  that makes BIF tractable.

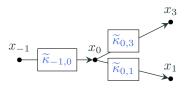
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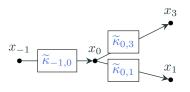
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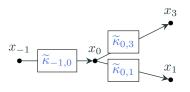
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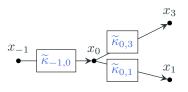
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Guided process Discrete case

Let the maps  $x\mapsto g_{s o t}(x)$  be specified for each edge (s,t) and define

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Practical way to choose  $g_{s o t}$ : replace kernel  $\kappa_{s o t}$  by approximation  $\widetilde{\kappa}_{s o t}$ .

**Define** the guided process  $X^\circ$  as the process starting in  $X_0^\circ = x_0$  and from the roots onwards evolving on the DAG  $\mathcal G$  according to transition kernel

$$\kappa_{\mathrm{pa}(s) \to s}^{\circ}(x_{\mathrm{pa}(s)}; \mathrm{d}y) = \frac{g_s(y) \kappa_{\mathrm{pa}(s) \to s}(x_{\mathrm{pa}(s)}; \mathrm{d}y)}{\int g_s(y) \kappa_{\mathrm{pa}(s) \to s}(x_{\mathrm{pa}(s)}; \mathrm{d}y)}, \qquad s \in \mathcal{S}.$$

#### Use of guided process

Let S denote the set of non-leaf vertices.

**Theorem.** Assume kernels towards leaf-nodes admit densities  $h_{\mathrm{pa}(v) \rightarrow v}$ . Then

$$h_0(x_0) = g_0(x_0) \mathbb{E} \left[ \prod_{s \in \mathcal{S}} w_{\mathrm{pa}(s) \to s}(X_{\mathrm{pa}(s)}^{\circ}) \prod_{v \in \mathcal{V}} \frac{h_{\mathrm{pa}(v) \to v}(X_{\mathrm{pa}(v)}^{\circ})}{g_{\mathrm{pa}(v) \to v}(X_{\mathrm{pa}(v)}^{\circ})} \right]$$

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Computationally, this implies a bidirectional scheme:

- 1. Backward pass for Filtering;
- 2. Forward pass for Guiding.

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- 1 In general BIF is intractable.
- Resolve by backward filtering with simpler kernels and forward simulating the corresponding guided process.
- This results in weighted samples from the conditioned process.

# Application: interacting particle process

Forward transitions:

$$\kappa_i(t,x) = \begin{bmatrix} \psi\left(\lambda N_i(t,x)\right) & 1 - \psi\left(\lambda N_i(t,x)\right) & 0\\ 0 & \psi(\mu) & 1 - \psi(\mu)\\ 1 - \psi(\nu) & 0 & \psi(\nu) \end{bmatrix},$$

where

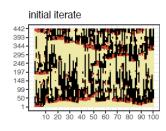
$$N_i(x) = \{\text{number of infected neighbours of individual } i \text{ in state } x\}$$

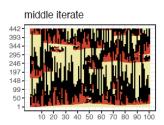
and 
$$\psi(u) = \exp(-\tau u)$$
.

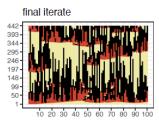
Auxiliary kernel for backward filtering:

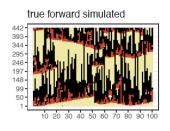
$$\widetilde{\kappa}_i = \begin{bmatrix} \psi(\widetilde{\lambda}_i(t)) & 1 - \psi(\widetilde{\lambda}_i(t)) & 0 \\ 0 & \psi(\mu) & 1 - \psi(\mu) \\ 1 - \psi(\nu) & 0 & \psi(\nu) \end{bmatrix}.$$

# **Application: interacting particle process**

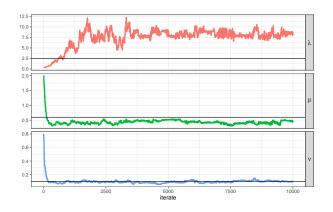








# **Application: interacting particle process**



Rethinking the discrete-time case:

• Edge

$$x_S \qquad x_T \\ \bullet \qquad \rightarrow \bullet$$

Suppose  $x\mapsto h(T,x)$  is given; wish to find  $x\mapsto h(S,x).$ 

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$$(\mathcal{A}h)(S,x) := \mathbb{E}[h(T,X_T) - h(S,X_S) \mid X_S = x]$$
$$= \int h(T,y)\kappa_{S\to T}(x, dy) - h(S,x).$$

•  $\bigwedge$  Obtain  $x \mapsto h(S, x)$  by solving (Ah)(S, x) = 0.

Define the infinitesimal generator of the space-time process  $(t,X_t)$ : for S < s < s + h < T

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⚠ Solving Kolmogorov backward equation is usually intractable.

# Defining the guided process via its inf.generator

• Backward filter with  $\widetilde{\mathcal{L}}$  instead of  $\mathcal{L}$ , such that solving  $(\widetilde{\mathcal{L}}g)(s,x)+\frac{\partial}{\partial s}g(s,x)=0$  becomes tractable.

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• Correct for "wrong" h by weight

$$\exp\left(\int_{t_i}^{t_{i+1}} \frac{(\mathcal{L}-\widetilde{\mathcal{L}})g}{g}(u,X_u^{\circ}) \,\mathrm{d}u\right).$$

• Care is needed in choice of  $\widetilde{\mathcal{L}}$ : matching conditions.

# How to solve the Kolmogorov Backward Equation?

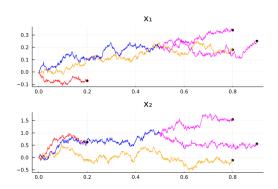
For  $s \in (S, T]$ ,

$$(\widetilde{\mathcal{L}}g)(s,x) + \frac{\partial}{\partial s}g(s,x) = 0, \qquad g(T,\cdot) = g_T(\cdot).$$

#### Examples/strategies:

- 1. If  $\widetilde{\mathcal{L}}$  is the infinitesimal generator of a linear diffusion process, then  $\log g(t,x)=c(t)+F(t)'x+x'H(t)x$  with ODE-system for (H(t),F(t),c(t)).
- 2. Ansatz  $g(t,x) = \sum_j c(t) \psi_j(t)$ . Derive ODE for c(t).

# **Example: branching diffusion**

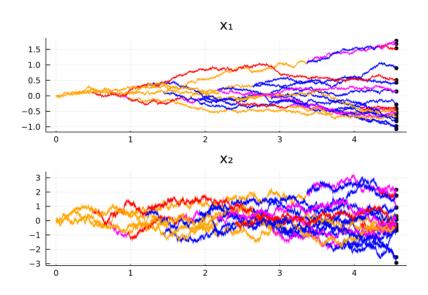


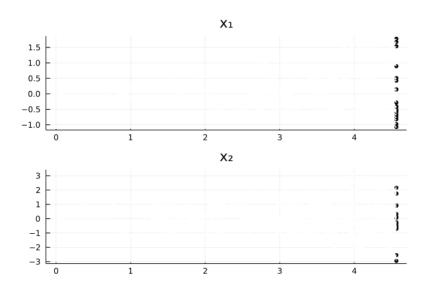
#### SDE on a tree where on each branch

$$\mathrm{d}X_t = \tanh. \left( \begin{bmatrix} -\theta_1 & \theta_1 \\ \theta_2 & -\theta_2 \end{bmatrix} X_t \right) \, \mathrm{d}t + \begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix} \, \mathrm{d}W_t.$$

Guided process Numerical illustration

33





On each branch

$$dX_t = \tanh \cdot \left( \begin{bmatrix} -\theta_1 & \theta_1 \\ \theta_2 & -\theta_2 \end{bmatrix} X_t \right) dt + \begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix} dW_t.$$

35

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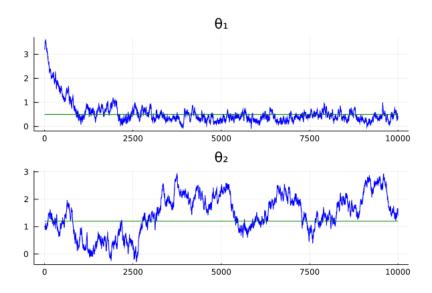
- Backward filter a linear process (essentially  $\widetilde{\kappa}$ )
- Write  $X^\circ$  as pushforward of  $(x_0,\xi,Z)$ , with  $\xi=(\theta_1,\theta_2,\sigma_1,\sigma_2)$
- $\bullet \ \ \mathsf{MCMC} \ \ \mathsf{on} \ \ (\xi,Z)$

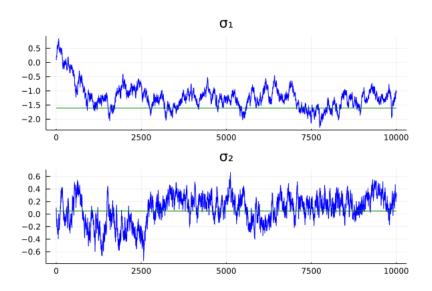
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Implementation in MitosisStochasticDiffEq.jl by Frank Schäfer (MIT).





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Ongoing: SPDEs, SDEs on manifolds, chemical reaction networks. Open postdoc position at VU Amsterdam.

Continuous-discrete smoothing of diffusions
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- Inference in Hidden Markov Models, CAPPÉ, MOULINES AND RYDÉN

Good source on filtering, smoothing, parameter estimation in HMM.