State space models, filtering and environmental applications

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Section 1

Introductory Examples

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Basics of state space models and filtering

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Particle filters

Ensemble Kalman filters

Extensions of particle filters

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Numerical precipitation predictions

Numerical predictions of 3-h rainfall in northern Switzerland (50 \times 100 grid with spacing of 2.2 km), separated by 12 h.



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Statistical postprocessing

Numerical weather predictions (MWP) are deterministic with high spatial and temporal resolution. Statistical postprocessing is needed to

- Correct biases of deterministic forecasts,
- Quantify uncertainty of deterministic forecasts,
- Account correctly for spatial and temporal dependence,
- Obtain predictive distributions which are calibrated and sharp.

Sigrist et al. (2014) use NWP forecasts as explanatory variables in a statistical space-time model of precipitation. The Kalman filter is an essential tool for fitting this model and for making predictions.

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Effect of postprocessing

NWP forecast (left) and median of predictive distribution (right) for one day in the validation period.



Mean absolute error of 3 forecasts



Weather prediction and data assimilation

As noted in the previous example, weather prediction is based on deterministic physical models, formulated as systems of differential equations. Since these equations are sensitive to initial conditions, predictions become unreliable after a few days. One needs to readjust the initial conditions for the next integration cycle frequently, using observations about the state of the atmosphere. In contrast to the previous example, observations are used not only for postprocessing forecasts, but as an input to the next forecast cycle.

Methods which use observations iteratively to estimate the state of a system are called data assimilation. In engineering, the same problem is called filtering.

Lorenz models

Because any real data assimilation example from atmospheric physics or oceanography is extremely high-dimensional and complex, often simple toy models are used as testbeds. The two most famous are due to Lorenz in 1963 and 1996. The Lorenz 63 model is

$$\frac{d}{dt} \begin{pmatrix} X_t^1 \\ X_t^2 \\ X_t^3 \end{pmatrix} = \begin{pmatrix} 10(X_t^2 - X_t^1) \\ X_t^1(28 - X_t^3) - X_t^2 \\ X_t^1 X_t^2 - \frac{8}{3}X_t^3 \end{pmatrix}.$$

The Lorenz 96 model is

$$\frac{dX_t^k}{dt} = (X_t^{k+1} - X_t^{k-2})X_t^{k-1} - X_t^k + 8, \quad k = 1, \dots, 40, \ X^k \equiv X^{k+40}.$$

One can also consider stochastic differential equations by adding white noise on the right-hand side to take model deficiencies into account.

Data assimilation for the Lorenz 63 model

Still to come: A figure illustrating how the particle filter looses track.

The Lorenz 63 model

This model has a famous attractor of fractional dimension. Switches between the two parts seem to occur at random times.



Stochastic kinetic reaction networks

Systems biology studies reaction networks with *r* reactions and *p* species of molecules. The number of molecules at time *t* is $x_t = (x_t^1, ..., x_t^p)$. If reaction *i* occurs at time *t*, then x_t changes as follows:

$$x_t^j = x_{t-}^j - v_{ij} + u_{ij}$$
 $(j = 1, ..., p).$

This means that reaction *i* consumes v_{ij} molecules of type *j* and produces u_{ij} molecules of type *j*. Reaction *i* occurs at the rate

$$\mu_i(t) = \theta_i \prod_{j, \mathbf{v}_{ij} \ge 1} \begin{pmatrix} \mathbf{x}_t^j \\ \mathbf{v}_{ij} \end{pmatrix}$$

provided all $x_t^j \ge v_{ij}$. I.e. the rate of reaction *i* is proportional to the number of ways the required molecules can be selected.

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Stochastic kinetic reaction networks, ctd.

- (x_t) is a Markov process in continuous time:
 - At time *t*, the time until the next reaction is exponential with mean $1/(\mu_1(t) + ... + \mu_r(t))$.
 - ► If a reaction occurs, it is type *i* with probability $\mu_i(t)/(\mu_1(t) + ... + \mu_r(t))$.
 - The lack of memory of the exponential distribution means that the future depends only on the current state.

The goal is to estimate the rate constants θ_i and the trajectory of the process from noisy measurements of some components $x_{t_i}^j$ at discrete times t_i .

Because of the incomplete and noisy observations, this is an example of a state space model.

Rare event estimation in a queueing network

From Amrein & H.K., ACM Transactions, 21, 2011.

- Goal: Estimate probabilities of rare events in Markov models, e.g. overflow in 2 queues in series.
- Arrival at queue 1 according to a Poisson process with rate
 1. Service times independent exponential with means ρ_i.
- State $X_t = (X_t^1, X_t^2)$, where $X_t^i =$ number of customers in queue i at time *t*.
- Let $A = \{0, 0\}$, $B = \{(x^1, x^2); x^2 \ge L\}$ and τ_A , τ_B first times of (re)entering A, B. Estimate $\gamma = \mathbb{P}_{(0,0)}(\tau_B < \tau_A)$.
- If ρ₁ < 1, ρ₂ < 1, L ≫ 1, γ is small, and direct Monte Carlo fails.

A simulated example

Still to come.

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Importance splitting

Alternatives to direct Monte Carlo are importance sampling and importance splitting.

Importance splitting recursively simulates from $\mathcal{L}(X_{\tau_{B_k}} \mid \tau_{B_k} < \tau_A)$ and estimates $\mathbb{P}(\tau_{B_k} < \tau_A \mid \tau_{B_{k-1}} < \tau_A)$ where $A^c \supset B_1 \supset \ldots \supset B_n = B$.

It is a generalization of particle filtering where one wants to approximate conditional distributions of a latent Markov process given an increasing number of instantaneous noisy observations. More generally, one studies recursive simulation from a sequence of different target distributions, so-called sequential Monte Carlo.

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Illustration of results

100 replicates of an interval estimate of γ (on log-scale) for $\rho_1 = \frac{1}{2}$ and $\rho_2 = \frac{1}{2}, \frac{1}{3}, \frac{1}{5}$.



State space models

Let X_t be the state vector of a system at time t. It is typically not fully observed, but has a simple Markovian dynamics. Available are partial observations Y_i of X at certain time points t_i , corrupted by independent noise variables.

To simplify notation, assume $t_i = i$ and (X_t) is time-homogeneous with

$$X_t \mid X_{t-1} = x_{t-1} \sim \mathcal{K}(dx_t \mid x_{t-1}).$$

 (X_t) can be in continuous time and deterministic, i.e. given by a differential equation.

Observations Y_1, Y_2, \ldots are conditionally independent with

$$Y_t \mid (X_s) \sim g(y \mid x_t) dy.$$

Existence of a density is crucial, but it could be w.r. to a different reference measure.

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Section 2

Introductory Examples

Basics of state space models and filtering The state space model Filtering and smoothing distributions

Kalman filter and its applications

Particle filters

Ensemble Kalman filters

Extensions of particle filters

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Graphical representation of state space models

The dependence between the variables of a state space model can be represented as follows

$$\cdots \rightarrow X_{t-1} \rightarrow X_t \rightarrow X_{t+1} \rightarrow \cdots$$

 $\downarrow \qquad \downarrow \qquad \downarrow$
 $\cdots \qquad Y_{t-1} \qquad Y_t \qquad Y_{t+1} \qquad \cdots$

It implies various conditional independencies which we will use later.

An alternative representation of a general state space model is

$$X_t = F(X_{t-1}, U_t), \quad Y_t = H(X_t, V_t)$$

where (U_t) and (V_t) are two independent white noises and *F* and *H* are arbitrary functions.

Notation

 $X_{s:t}$ is shorthand for $(X_s, X_{s+1}, \ldots, X_t)$, similarly $y_{s:t}$.

Define conditional distribution of states in time interval [s, t] given observations up to time r:

 $\pi_{s:t|r}(dx_{s:t} \mid y_{1:r}) := \mathcal{L}(X_{s:t} \mid y_{1:r}).$

The *y*'s are fixed and thus are often omitted in $\pi_{s:t|r}$. To simplify, use $\pi_{t|r}$ for the marginal $\pi_{t:t|r}$. Call $\pi_{t|t-1}$ the prediction and $\pi_{t|t}$ the filter distribution. Distributions with s < r are called smoothing distributions.

By abuse of notation, we use P(P(. | .)) for other (conditional) distributions that will play a role, and p(.)(p(. | .)) for (conditional) densities. The arguments will indicate which variables are involved.

Filtering recursions

Prediction and filter distributions can be computed recursively:

• Propagation $\pi_{t-1|t-1} \longrightarrow \pi_{t|t-1}$ via conditioning on X_{t-1}

$$\pi_{t|t-1}(dx_t) = \int K(dx_t \mid x_{t-1}) \pi_{t-1|t-1}(dx_{t-1})$$

• Update $\pi_{t|t-1} \longrightarrow \pi_{t|t}$ via Bayes' formula:

$$\pi_{t|t}(dx_t) \propto \pi_{t|t-1}(dx_t) \times g(y_t \mid x_t).$$

We start the recursion with $\pi_{0|0}$ = the initial distribution of X_0 . The denominator in Bayes' formula is

$$\int g(y_t \mid x_t) \pi_{t|t-1}(dx_t) = p(y_t \mid y_{1:t-1})$$

Thus the joint density of $Y_{1:t}$ is a byproduct of the filter.

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Gibbs sampler for state space models

If $K(dx_t | x_{t-1}) = k(x_t | x_{t-1})dx_t$, full conditionals are

$$p(x_{s} \mid y_{1:t}, x_{0:s-1}, x_{s+1:t}) = p(x_{s} \mid y_{s}, x_{s-1}, x_{s+1}) \\ \propto g(y_{s} \mid x_{s})k(x_{s} \mid dx_{s})k(x_{s+1} \mid x_{s}).$$

If the model contains unknown parameters, also the density of parameters given $x_{0:t}$ and $y_{1:t}$ is available. Hence the Gibbs sampler can be used. But

- Convergence is in most cases slow: Full conditionals are too tight.
- Better to update $x_{0:t}$ jointly (or at least in big blocks).
- If observations become available sequentially, want to update the samples recursively.

Smoothing

Analogous recursions as for the filter hold for $\pi_{0:t|t}$ with an extension instead of a propagation step

$$\begin{array}{lll} \pi_{0:t|t-1}(dx_{0:t}) &= & \mathcal{K}(dx_t \mid x_{t-1})\pi_{0:t-1|t-1}(dx_{0:t-1}) \\ \pi_{0:t|t}(dx_{0:t}) &\propto & \pi_{0:t|t-1}(dx_{0:t}) \times g(y_t \mid x_t). \end{array}$$

Alternatively, we can use that given $y_{1:t}$ the state process is still a Markov chain, by rules for conditional independence. If $K(dx_t | x_{t-1}) = k(x_t | x_{t-1})dx_t$, then the backward transition of the conditional chain is a modification of the filter distribution:

$$\begin{array}{rcl} {\cal P}(dx_{s} \mid x_{s+1}, y_{1:t}) & = & {\cal P}(dx_{s} \mid x_{s+1}, y_{1:s}) \\ & \propto & k(x_{s+1} \mid x_{s}) \pi_{s|s}(dx_{s}). \end{array}$$

There are also expressions for the forward transition distributions which we will discuss later.

Implementations

- Although the recursions look simple, doing the integrations is in most cases difficult.
- Closed form solutions exist if X_t takes values in a finite set (Baum-Welch) or in the Gaussian linear case (Kalman) that we will discuss in the next section. These are all practically relevant cases.
- Analytical approximations exist (Extended Kalman filter, unscented Kalman filter), but they are limited in scope.
- Numerical approximations are difficult because X_t is often high-dimensional and π_{t|t} lives in different parts of the state space for different t's.
- Particle and Ensemble Kalman filters are recursive Monte Carlo approximations.

Gaussian linear state space models

A Gaussian linear state space model has the form

 $X_t = FX_{t-1} + U_t, \quad Y_t = HX_t + V_t$

where (U_t) and (V_t) are independent Gaussian white noises with mean 0 and covariance Q and R, respectively. In other words, we have a partially observed vector autoregression.

If also X_0 is Gaussian, then all conditional distributions $\pi_{s:t|r}$ are again Gaussian. The Kalman filter and smoother computes required conditional means and covariances recursively. It can be derived from the general recursions above, or directly from basic properties of orthogonal projections.

Section 3

Introductory Examples

Basics of state space models and filtering

Kalman filter and its applications

Gaussian linear state space models Kalman filter and smoother Fitting of stochastic advection-diffusion models

Particle filters

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Kalman filter

Denote mean and covariance of $\pi_{s|t}$ by $\mu_{s|t}$ and $P_{s|t}$). Then

$$\mu_{t|t-1} = F\mu_{t-1|t-1}, \quad P_{t|t-1} = FP_{t-1|t-1}F' + Q$$

and

$$\mu_{t|t} = \mu_{t|t-1} + K_t(y_t - H\mu_{t|t-1}),$$

$$P_{t|t} = (I - K_t H) P_{t|t-1} = (I - K_t H) P_{t|t}(I - K_t H)' + K_t R K_t'$$

where

$$K_t = K(H, P_{t|t-1}, R) = P_{t|t-1}H'(HP_{t|t-1}H' + R)^{-1}$$

= Cov $(X_t - \mu_{t|t-1}, Y_t - H\mu_{t|t-1})$ Cov $(Y_t - H\mu_{t|t-1})^{-1}$.

is the so-called Kalman gain.

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Kalman smoother

Because $P(dx_s | x_{s+1}, y_{1:t}) \propto k(x_{s+1} | x_s)\pi_{s|s}(dx_s)$, backward transitions are obtained as in the filter update

$$\begin{split} \mathbb{E}(X_{s} \mid X_{s+1}, y_{1:t}) &= \mathbb{E}(X_{s} \mid X_{s+1}, y_{1:s}) \\ &= \mu_{s|s} + K(F, P_{s|s}, Q)^{-1}(X_{s+1} - F\mu_{s|s}), \\ \text{Cov}\left(X_{s} \mid X_{s+1}, y_{1:t}\right) &= (I - K(F, P_{s|s}, Q))P_{s|s}. \end{split}$$

This is the basis of the forward-filtering backward simulation algorithm.

It is not difficult to derive also a backward recursion for the mean and covariance of $\pi_{s|t}$.

Back to the first example

I now show how the Kalman filter and smoother is used for the statistical postprocessing of numerical weather predictions (NWP).

Rainfall $y(t, \mathbf{s})$ is skewed and has an atom at zero, and thus we cannot use a Gaussian distribution. We assume it depends on a latent "potential rainfall" $w(t, \mathbf{s})$ which has a Gaussian distribution through

$$y(t, \mathbf{s}) = w(t, \mathbf{s})^{\lambda} \mathbf{1}_{\{w(t, \mathbf{s}) > 0\}}.$$

I will first discuss a class of Gaussian space-time models assuming Gaussian observations. At the end I show how we handle the non-linear dependence of y(t, s) on w(t, s).

Computational complexity

If $X_t \in \mathbb{R}^q$ and $Y_t \in \mathbb{R}^d$ and we consider *T* time steps:

- Forward filtering and likelihood evaluation has complexity $O(T(q^2 + d^3))$.
- Backward smoothing has additional complexity $O(Tq^3)$.
- If q and/or d are large, efficient computation becomes crucial.
- Savings are possible if P_{t-1|t} and P_{t|t} are sparse by using sparse Cholesky factorizations.
- We discuss next the use of Kalman filter and smoother for the introductory example of precipitation.

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Modeling potential rainfall

We assume that the potential rainfall w(t, s) is a Gaussian space-time random field

$$w(t, \mathbf{s}) = \beta_1 y_F(t, \mathbf{s})^{1/\tilde{\lambda}} + \beta_2 \mathbf{1}_{\{y_F(t, \mathbf{s}) = 0\}} + X(t, \mathbf{s}) + \nu(t, \mathbf{s}),$$

- $y_F(t, s)$ NWP forecast made at 0:00 UTC of the same day,
- $X(t, \mathbf{s})$ structured random effect (mean zero),
- $\nu(t, \mathbf{s})$ nugget effect,
- Unknown parameters: β_1 , β_2 , $\lambda > 0$, nugget variance σ_{ν}^2 , parameters for *X* (to be discussed later).

We use precipitation data from 32 stations that the NWP forecast does not use directly.

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Gaussian space-time fields

- X(t, s) = Gaussian process on [0,∞) × ℝ^d with mean zero and covariance function C_θ. How to choose C_θ ?
- General considerations:
 - Take different nature of time and space into account
 - Avoid separable models
 - Find compromise between simplicity and generality
 - Use parameters which have a clear interpretation
- Computational tractability:

Illustration of deterministic PDE

- Evaluation of likelihood has in general the complexity O((TN)³), T and N number of points in time and space.
- Adding noise to simple physics based models satisfies the requirements above. Our contribution: Solution of computational issues.

Stochastic advection-diffusion model

 Gaussian process defined through stochastic partial differential equation (SPDE)

$$\frac{\partial}{\partial t}X(t,\mathbf{s}) = -\boldsymbol{\mu}\cdot\nabla X(t,\mathbf{s}) + \nabla\cdot\boldsymbol{\Sigma}\nabla X(t,\mathbf{s}) - \kappa\,X(t,\mathbf{s}) + \boldsymbol{\epsilon}(t,\mathbf{s})$$

- $-\mu \cdot \nabla X(t, \mathbf{s})$ transport, μ drift vector ("advection")
- $\nabla \cdot \mathbf{\Sigma} \nabla X(t, \mathbf{s})$ anisotropic diffusion
- $-\kappa X(t, \mathbf{s})$ damping
- ϵ(t, s) temporally white noise and spatially colored forcing
 or source-sink ("convection")
- Continuous time and space model

Illustration of SPDE





Elementary solutions of the SPDE

Assume that for a fixed wavenumber $\textbf{\textit{k}} \in \mathbb{R}^2$

 $X(0, \mathbf{s}) = \alpha(0) \exp(i\mathbf{k}'\mathbf{s}), \quad \epsilon(t, \mathbf{s}) = \dot{W}(t) \exp(i\mathbf{k}'\mathbf{s})$

where \dot{W} is white noise (i.e. *W* is Brownian motion).

Then the solution is

$$X(t, \boldsymbol{s}) = \left(e^{-h(\boldsymbol{k})t}\alpha(0) + \int_0^t e^{-h(\boldsymbol{k})(t-u)}\dot{W}(u)du\right) \exp\left(i\boldsymbol{k}'\boldsymbol{s}\right)$$

where $h(\mathbf{k}) = i\mu'\mathbf{k} + \mathbf{k}'\mathbf{\Sigma}\mathbf{k} + \kappa$. Because h > 0, the solution forgets the initial amplitude $\alpha(0)$, and we have for large *t* a Gaussian amplitude which is stationary in time.

By linearity of the SPDE, sums of such solutions are again solutions.

Discretizing time and space

Assume we are interested in X at a finite set of locations and times. Truncating wavenumbers to a finite set, we have

$$\boldsymbol{X}(t_i) = \boldsymbol{\Phi} \boldsymbol{\alpha}(t_i), \quad \boldsymbol{\alpha}(t_i) = \boldsymbol{G} \boldsymbol{\alpha}(t_{i-1}) + \mathcal{N}(0, \boldsymbol{Q})$$

Here

- $\boldsymbol{\Phi}$ has elements $\exp(i\boldsymbol{k}_{\ell}'\boldsymbol{s}_{i})$.
- **G** and **Q** are diagonal.
- G depends on the time step t_i t_{i-1} and on the parameters μ, Σ and κ of the SPDE.
- ► Q depends on the time step t_i t_{i-1} and on the assumed spatial spectral density f_e of e(t, .). We use the Matérn density with smoothness parameter 1.
- The time discretization of α(t) is exact, the only approximation is the truncation of wavenumbers.

Stationary solutions of the SPDE

If $X(0, \mathbf{s})$ and $\epsilon(t, \mathbf{s})$ are stationary in space, by the Cramér representation

$$X(0, \mathbf{s}) = \int \exp{(i\mathbf{k}'\mathbf{s})} \alpha(0, d\mathbf{k}), \quad \epsilon(t, \mathbf{s}) = \int \exp{(i\mathbf{k}'\mathbf{s})} \dot{W}(t, d\mathbf{k}),$$

with $\mathbb{E}\left(\dot{W}(t, d\mathbf{k}) \overline{\dot{W}(u, d\mathbf{j})}\right) = \delta_{t,u} \delta_{\mathbf{k},\mathbf{j}} f_{\epsilon}(\mathbf{k}) d\mathbf{k}$, and similarly for $\alpha(0, d\mathbf{k})$.

Then the solution has the representation

$$\begin{aligned} X(t, \mathbf{s}) &= \int \exp{(i\mathbf{k}'\mathbf{s})}\alpha(t, d\mathbf{k}), \\ \alpha(t, d\mathbf{k}) &= e^{-h(\mathbf{k})t}\alpha(0, d\mathbf{k}) + \int_0^t e^{-h(\mathbf{k})(t-u)}\dot{W}(u, d\mathbf{k})du. \end{aligned}$$

Amplitudes develop independently for differen wavenumbers.

Fitting with linear Gaussian observations

Adding a nugget to \boldsymbol{X} , we obtain a Gaussian linear state space model with states $\alpha(t_i)$ and observations \boldsymbol{w}_i :

$$\alpha(t_i) = \boldsymbol{G}\alpha(t_{i-1}) + \mathcal{N}(0, \boldsymbol{Q}), \quad \boldsymbol{w}_i = \boldsymbol{\Phi}\alpha(t_i) + \mathcal{N}(0, \sigma_{\nu}^2 \boldsymbol{I}).$$

If we use sin and cos instead of the complex exponential, \boldsymbol{G} becomes 2 \times 2 block diagonal.

If also parameters are unknown, do simultaneous updates of parameters and states: Propose

$$(oldsymbol{lpha}^*, heta^*) \mid oldsymbol{lpha}, heta \sim oldsymbol{q}(heta^* \mid heta) oldsymbol{
ho}(oldsymbol{lpha}^* \mid oldsymbol{w}, heta^*) oldsymbol{d} heta^* oldsymbol{d}^*$$

and accept it with probability $\min(1, p(\theta^* | \boldsymbol{w}) / p(\theta | \boldsymbol{w}))$.

This is feasible as long the set of wavenumbers and the set of locations with observations are not too big. Sampling X on a finer grid can be done at the end.

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Fitting with linear Gaussian observations, ctd.

If **w** is on an $n \times n$ grid with spacing Δ , we can use wavenumbers on the grid with spacing $2\pi/(\Delta n)$. The Kalman filter and smoother decouple into 2-dimensional subproblems because Φ is then orthogonal:

$$oldsymbol{\Phi}'oldsymbol{w}_i = oldsymbol{lpha}(t_i) + \mathcal{N}(oldsymbol{0}, \sigma_
u^2oldsymbol{I})_i$$

and FFT makes evaluation of $\Phi' w_i$ fast.

If observations are available on only parts of the grid, we can impute the missing values in an iterative procedure.

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Section 4

Introductory Examples

Basics of state space models and filtering

Kalman filter and its applications

Particle filters

The basic particle filter Balanced sampling Auxiliary particle filter Likelihood estimation Importance splitting as a particle filter

Ensemble Kalman filters

Extensions of particle filters

Nonlinear and non-Gaussian observations

In the precipitation example, iterate between updates of $(\boldsymbol{w}, \lambda)$ given $\alpha, \boldsymbol{y}, \sigma_{\nu}^2$ and updates of (θ, α) given \boldsymbol{w} . For the former, propose first a new λ . If accepted, modify \boldsymbol{w} deterministically at sites where $\boldsymbol{y} > 0$. At other sites, sample \boldsymbol{w} from a truncated Gaussian distribution.

Similar procedure is possible for binary data in a probit model. Otherwise, need to use iterative methods with Gaussian approximations of $p(y(t, \mathbf{s}) | X(t, \mathbf{s}))$.

Particle Filter

Remember the filtering recursions:

- Propagation: $\pi_{t|t-1}(dx_t) = \int K(dx_t \mid x_{t-1})\pi_{t-1|t-1}(dx_{t-1})$.
- Update: $\pi_{t|t}(dx_t) \propto \pi_{t|t-1}(dx_t) \times g(y_t \mid x_t)$.

Recursive Monte Carlo implementation: If $(x_{t-1|t-1}^{j}; j = 1, ..., N)$ is an (approximate) sample from $\pi_{t-1|t-1}$, construct an (approximate) sample from $\pi_{t|t}$ by

- Propagation: Generate (x^j_{t|t-1}) by simulating the Markov process with initial conditions X_{t-1} = x^j_{t-1|t-1}, independently for different *j*'s.
- Update: Obtain $(x_{t|t}^{j})$ by resampling from $(x_{t|t-1}^{j})$ with probabilities proportional to $g(y_t | x_{t|t-1}^{j})$.

The $x_{t|t-1}^{j}$'s and $x_{t|t}^{j}$'s are called particles. They move in space and have offsprings. They interact via the number of offsprings.

Why resampling ?

 Update step can be split into a weighting and a resampling step. This gives a weighted approximation of

$$\pi_{t|t}(dx_t) \approx \sum_{j=1}^N w_t^j \Delta_{x_{t|t-1}^j}(dx_t), \quad w_t^j \propto g(y_t \mid x_{t|t-1}^j).$$

- Resampling introduces additional noise, but it avoids waisting simulation efforts in the next propagation step with unlikely starting values.
- Hence it is better to use the weighted approximation of \u03c0_{t|t} and to resample at the beginning of the propagation step.
- In particular, resampling introduces ties among x^j_{t|t}. These are broken in the next propagation step (called "rejuvenation") if state dynamics is stochastic.
- If some components of state change deterministically, can break ties by adding noise (compensated by shrinkage to the mean).

Several steps of the particle filter

Area of circles \propto weight. $Y_t = X_t^2$ plus noise, Blue=true (unobserved) X_t .



A single step of the particle filter

Left: Propagation (only few arrows shown). Right: Reweighting and resampling



Effective sample size (ESS)

If we have a weighted sample (X^j, w^j) of size *N* constructed by importance sampling, to how many unweighted sample values does this correspond ? Liu et al. gave the answer

$$\mathrm{ESS} = \left(\sum_{j=1}^{N} (w^j)^2\right)^{-1}.$$

This is correct in the two extreme cases $w^j = 1/N$ for all *j* and $w^j = 1$ for one *j*. The general case follows from an approximation of the variance of $\sum_i w^j \psi(X^j)$.

It has been suggested to resample only if ESS is below a certain threshold. Note however that in a recursive setting, ESS is presumably too optimistic since the propagation step does not fully compensate the loss of precision due to weighting and resampling at time t - 1.

Balanced sampling

For resampling, can use any scheme s.th. $x_{t|t-1}^{j}$ is chosen on average Nw_{t}^{j} times. A sampling scheme is called balanced if $x_{t|t-1}^{j}$ is chosen $\left[Nw_{t}^{j}\right]$ or $\left[Nw_{t}^{j}\right] + 1$ times.

There are many balanced sampling schemes. The simplest such scheme is

$$x_{t|t}^{j} = x_{t|t-1}^{l_{j}}, \quad l_{j} = F_{t}^{-1}((j-U)/N)$$

where *U* is uniform(0, 1) and F_t is the cdf of the discrete distribution on $\{1, 2, ..., N\}$ with weights w_t^j , see figure.

This is hard to analyze theoretically because no limit theory applies. So-called tree sampling is an alternative.

Auxiliary particle filter

Idea: Propagate particles with a transition $Q \neq K$ to bring them closer to y_t and adjust with suitable weights which are more equal

- Generate x^j_{t,*} ~ q(x_t | x^j_{t-1|t-1}, y_t)dx_t, independently for different *j*'s. (Change of notation since the x^j_{t,*} are no longer a sample from π_{t|t-1}).
- Resample the particles $x_{t,*}^j$ with probabilities

$$w_t^j \propto ilde w(x_{t-1|t-1}^j, x_{t,*}^j, y_t) = rac{g(y_t \mid x_{t,*}^j)k(x_{t,*}^j \mid x_{t-1|t-1}^j)}{q(x_{t,*}^j \mid x_{t-1|t-1}^j, y_t)}.$$

Illustration of balanced sampling



Auxiliary particle filter ctd.

- Weights w_t^j depend on $x_{t,*}^j$ and $x_{t-1|t-1}^j$.
- $((x_{t-1|t-1}^j, x_{t,*}^j), w_t^j)$ is a weighted sample from $\pi_{t-1:t|t}$.
- Variance of weights \tilde{w} is minimal for

$$q(x_t \mid x_{t-1}, y_t) = p(x_t \mid x_{t-1}, y_t) = \frac{g(y_t \mid x_t)k(x_t \mid x_{t-1})}{p(y_t \mid x_{t-1})}.$$

In this case, $w_t^j \propto p(y_t \mid x_{t-1|t-1}^j)$, and resampling before propagation increases diversity.

► Also in other cases one can add a resampling step with probabilities \(\tau\)(x^j_{t-1|t-1}, \(y\)t)\) before propagation. Then weights after propagation are

$$w_t^j \propto \tilde{w}(x_{t-1|t-1}^j, x_{t,*}^j, y_t) = \frac{g(y_t \mid x_{t,*}^j) k(x_{t,*}^j \mid x_{t-1|t-1}^j)}{\tau(x_{t-1|t-1}^j, y_t) q(x_{t,*}^j \mid x_{t-1|t-1}^j, y_t)}.$$

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Auxiliary particle filter, ctd.

Instead of sampling approximately from $\pi_{t-1:t|t}$, we can also look at sampling from the approximate filter density

$$\hat{\pi}_{t|t}(x_t) \propto \frac{1}{N} \sum_{j=1}^{N} g(y_t \mid x_t) k(x_t \mid x_{t-1|t-1}^j)$$

Importance sampling of $x_{t,*}$ alone leads to an $O(N^2)$ algorithm since computing a single weight involves a sum over N terms. Importance sampling of a mixture index I together with $x_{t,*}$ avoids this problem: If we use a proposal

$$\mathbb{P}(I=j) \propto \tau(x_{t-1|t-1}^{j}, y_{t}), \ p(x_{t,*} \mid I=j) = q(x_{t,*} \mid x_{t-1|t-1}^{j}, y_{t})$$

then the importance weights are indeed proportional to

$$\frac{g(y_t \mid x_{t,*}^j)k(x_{t,*}^j \mid x_{t-1|t-1}^j)}{\tau(x_{t-1|t-1}^j, y_t)q(x_{t,*}^j \mid x_{t-1|t-1}^j, y_t)}.$$

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Likelihood estimation

Remember that the the likelihood increment $p(y_t | y_{1:t-1})$ is the denominator in Bayes formula in the update step,

$$\int g(y_t \mid x_t) \pi_{t|t-1}(x_t) dx_t = \int g(y_t \mid x_t) \pi_{t-1:t|t-1}(x_t, x_{t-1}) dx_t dx_{t-1}.$$

Hence we have the estimate

$$\widehat{\rho}(y_t \mid y_{t-1}) = \frac{1}{N} \sum_{j=1}^{N} g(y_t \mid x_{t|t-1}^j)$$

which we compute when normalizing the weights.

If we use the auxiliary particle filter, we have similarly

$$\widehat{p}(y_t \mid y_{t-1}) = \frac{1}{N} \sum_{j=1}^{N} \widetilde{w}(x_{t-1|t-1}^j, x_{t,*}^j, y_t)$$

Implementing the auxiliary particle filter

The optimal choices $\tau(x_{t-1|t-1}^j, y_t) = p(y_t | x_{t-1|t-1}^j)$ and $q(x_t | x_{t-1}, y_t) = p(x_t | x_{t-1}, y_t)$ are usually not available explicitely, so need an approximation. Simplest choice by Gaussian approximation

$$\log k(x_t \mid x_{t-1}) + \log g(y_t \mid x_t) \\\approx c(x_{t-1}, y_t) + b(x_{t-1}, y_t)' x_t + \frac{1}{2} x_t' A(x_{t-1}, y_t) x_t,$$

obtained by using e.g. a Taylor approximation around the arg \max_{x_t} of $k(x_t \mid x_{t-1})g(y_t \mid x_t)$.

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Likelihood estimation is unbiased

In general the particle filter does not produce unbiased estimates of $\int \psi(x_t) \pi_{t|t}(dx_t)$. But we have

Lemma If x_0^j is sampled from the initial distribution of X_0 and if resampling is unbiased, then

$$\mathbb{E}(\prod_{t=1}^{n} \widehat{p}(y_t \mid y_{1:t-1})) = \prod_{t=1}^{n} p(y_t \mid y_{1:t-1}) = p(y_{1:n}),$$

but in general $\mathbb{E}(\widehat{\rho}(y_t \mid y_{1:t-1})) \neq \rho(y_t \mid y_{1:t-1}).$

This will be used in rare event estimation and in particle MCMC.

Properties of particle filter

- ► Under weak conditions, it is consistent (as N → ∞) and a CLT holds.
- Under strong conditions, can also show that consistency is uniform in *t* and variance in CLT is uniformly bounded in *t*.
- Respects constraints $\psi(x(t)) \equiv 0$.
- Weights degenerate quickly as dimensions grow (see Bickel et al. for a theoretical explanation). Particle filter can loose track easily.
- Auxiliary particle filter can bring substantial improvements, but not applicable if density k does not exist or is not available analytically.

Implementing importance splitting

- A particle with τ_{B_k} > τ_A gets weight 0 and is thus killed.
 Resampling inflates the number R_k of surviving particles at step k back to N. Impossible if R_k = 0, so then γ̂ := 0.
- ► Observations are not made available sequentially, so simulation effort can vary in each step. It is better to control precision by propagating particles until a fixed value ≥ 2 for R_k is achieved. Then N_{k-1} becomes random.
- Unbiasedness of $\widehat{\gamma}$ continues to hold if we use the UMVU estimator for the negative binomial

$$\widehat{\mathbb{P}}(\tau_{B_k} < \tau_A \mid \tau_{B_{k-1}} < \tau_A) = rac{R_k - 1}{N_{k-1} - 1}.$$

In our paper we also address the choices of n, B_k, R_k by a two-step procedure and construct confidence intervals for γ by running particle filters in parallel.

Importance splitting as a particle filter

- Let (X_t) be a Markov process on some set E, A, B ⊂ E A ∩ B = Ø. We want to estimate γ = P_{x0}(τ_B < τ_A) where τ_A and τ_B are first times of (re)entering A, B and x₀ ∉ B.
- Importance splitting chooses A^c ⊃ B₁ ⊃ ... B_n = B and simulates recursively from L(X_{τB_k} | τ_{B_k} < τ_A}.
- If we consider the state process X'_k = X_{τB_k} and assume that "observations" Y_k = 1_[τB_k<τ_A] all have the value 1, this becomes a filtering problem.
- Moreover

$$\gamma = \prod_{k=1}^n \mathbb{P}(\tau_{B_k} < \tau_A \mid \tau_{B_{k-1}} < \tau_A) = \prod_{k=1}^n \mathbb{P}(Y_k = 1 \mid Y_{1:k-1} \equiv 1).$$

Therefore estimation of the "likelihood" gives the desired estimate of γ , and it is unbiased !

Section 5

Introductory Examples

Basics of state space models and filtering

Kalman filter and its applications

Particle filters

Ensemble Kalman filters

Basic idea Square root and perturbed observation versions Fine tuning Bridging the two methods

Extensions of particle filters

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Basic idea of Ensemble Kalman filter

Assume Gaussian-linear observations

 $Y_t = HX_t + \mathcal{N}(0, R).$

Prediction step as for particle filter, but update as if $\pi_{t|t-1}$ is Gaussian with mean $\mu_{t|t-1}$ and covariance $P_{t|t-1}$.

Then $\pi_{t|t}$ is again Gaussian with mean

$$\mu_{t|t} = \mu_{t|t-1} + K(H, P_{t|t-1}, R)(y_t - H\mu_{t|t-1})$$

$$P_{t|t} = (I - K(H, P_{t|t-1}, R)H)P_{t|t-1}.$$

Ensemble Kalman filter estimates $\mu_{t|t-1}$ and $P_{t|t-1}$ and converts $(x_{t|t-1}^{j})$ into a sample with the correct first and second moment.

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Particle filter for banana-shaped prediction

A single observation $y = x_1 + \mathcal{N}(0, 0.5^2) \in \{-1, 1\}$. Banana-shaped prediction sample (black). Particle filter update (blue).



Square root and stochastic version

The Ensemble Kalman filter comes in two versions:

Stochastic: Perturbed observation EnKF

$$\mathbf{x}_{t|t}^{j} = \mathbf{x}_{t|t-1}^{j} + \widehat{K}_{t}(\mathbf{y}_{t} + \frac{\varepsilon_{t}^{j}}{t} - H\mathbf{x}_{t|t-1}^{j}), \quad \varepsilon_{t}^{j} \sim \mathcal{N}(\mathbf{0}, R)$$

where $\widehat{K}_t = K(H, \widehat{P}_{t|t-1}, R)$. This is the update for the mean with artificial noise added to the observation.

• Deterministic: Ensemble square root filter. It applies an affine linear transformation to $(x_{t|t-1}^{j})$.

Ensemble Kalman filter for banana-shaped prediction

A single observation $y = x_1 + \mathcal{N}(0, 0.5^2) \in \{-1, 1\}$. Banana-shaped prediction sample (black). Ensemble Kalman filter update (blue).



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Suppose that $\pi_{t|t-1}$ is not Gaussian. What can we say about the limiting distribution as $N \to \infty$? Because prediction sample is modified, the limit distribution is not Gaussian.

Some analysis is possible when $\pi_{t|t-1}$ is a Gaussian mixture. Limit distribution exists for both versions, but is different from the correct update according to Bayes formula, and depends on which version is used.

Typically (see next slide) the stochastic version is too close to a Gaussian distribution, the deterministic version often has peaks at wrong places.

Properties of the EnKF

Both versions:

- + With some tuning: Works well also for $N \ll \dim(y)$.
- Not consistent unless $\pi_{t|t-1}$ is Gaussian. Observation *y* influences only location, but not scale or shape of the updated ensemble. Does not respect constraints $\psi(x(t)) \equiv 0$.

Stochastic:

+ Robust in highly non-linear non-Gaussian systems.

Loss of skill due to additional Monte Carlo error in update.
 Deterministic:

- + No Monte Carlo error in update formula.
- Susceptible to outliers in the prediction ensemble that arise in highly non-linear systems.

Comparing the two versions: Examples

A single observation $y = x + \mathcal{N}(0, 1)$. The prediction distribution is a Gaussian mixture with *L* components and composite variance equal to one. True update distribution (black), EnSRF (blue), POEnKF (golden).



Fine tuning

Essentially, two different sources of error may impact filter performance:

- Systematic errors
 - Misspecifed dynamics
 - Misspecified observation mechanisms (representation errors, wrong error covariances)
 - Non-Gaussian prediction distributions
- Monte Carlo error due to limited ensemble size. Affects in particular estimation of covariance mat P_{t|t-1}.

Both sources of error typically lead to a too small spread of the updated ensemble, i.e., the uncertainty is underestimated.

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Covariance inflation

Basic idea: Artificially increase the spread of the prediction ensemble by a factor $\delta > 1$:

$$x_{t|t-1}^{j} \mapsto \tilde{x}_{t|t-1}^{j} = \overline{x}_{t|t-1} + \frac{\delta}{\delta} \cdot (x_{t|t-1}^{j} - \overline{x}_{t|t-1}). \quad (j = 1, \dots, N)$$

This leaves the prediction mean unchanged, but inflates the sample covariance: $\widehat{\text{Cov}}(\tilde{x}_{t|t-1}) = \delta^2 \cdot \widehat{\text{Cov}}(x_{t|t-1})$. It can be used for XEnKF and EnKPF.

Inflation factor δ may be spatially varying (a different factor for each coordinate). δ is typically chosen such that the filter is correctly calibrated, i.e., in the long run the observations should behave as if taken from the prediction or updated ensemble.

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Covariance tapering

Basic idea: Replace the sample covariance matrix $\hat{P}_{t|t-1}$ of the prediction ensemble by a regularized estimate that has a smaller Monte Carlo error.

If components of x correspond to spatial location, regularize by assuming that separated state variables should be uncorrelated (called tapering):

$$\hat{P}_{t|t-1}\mapsto C\circ\hat{P}_{t|t-1},$$

where C is a sparse correlation matrix and \circ denotes entry-wise multiplication.

Taper C is usually constructed via a compactly supported correlation function.

Localization

The components of x and y have a spatial interpretation in many applications. An observation y should have low impact on spatially distant state variables. Two paradigms:

- Covariance tapering ("covariance filtering, background") localization")
- Local updates ("observation localization, localization in grid space")

Bridging the two methods: Overview

- Many methods in the literature attempt to combine advantages of EnKF and PF.
- In our experience, these methods are difficult to implement: Either the description of the algorithm is not clear or it has many tuning parameters which are hard to choose or we could not reproduce the results.
- We have proposed two generalizations of PoEnKF, called XEnKF and EnKPF. Both rely on an exact update formula for Gaussian mixtures and have one tuning constant.
- XEnKF is based on clustering of the prediction particles. Tuning parameter is the number of clusters.
- In this talk, focus on the EnKPF.

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Updates of Gaussian mixtures

Explicit updates are possible if $\pi_{t|t-1}$ is a Gaussian mixture (GM). $\pi_{t|t}$ is again a Gaussian mixture:

$$\pi_{t|t-1} = \sum_{\ell=1}^{L} \alpha_{t|t-1}^{\ell} \mathcal{N}(\mu_{t|t-1}^{\ell}, P_{t|t-1}^{\ell}) \Rightarrow \pi_{t|t} = \sum_{\ell=1}^{L} \alpha_{t|t}^{\ell} \mathcal{N}(\mu_{t|t}^{\ell}, P_{t|t}^{\ell})$$

where $(\mu_{t|t-1}^\ell, P_{t|t-1}^\ell) o (\mu_{t|t}^\ell, P_{t|t}^\ell)$ as in the Kalman filter and

 $\alpha_{t|t}^{\ell} \propto \alpha_{t|t-1}^{\ell} \times \varphi(\mathbf{y}|\mathbf{H}\boldsymbol{\mu}_{t|t-1}^{\ell}, \mathbf{H}\mathbf{P}_{t|t-1}^{\ell}\mathbf{H}' + \mathbf{R})$

(φ denotes the Gaussian density). "Particle filter for the weights, Kalman filter for mean and variance".

EnKF as a Gaussian mixture

We can consider $x_{t|t}^j = x_{t|t-1}^j + \hat{K}(y + \varepsilon^j - Hx_{t|t-1}^j)$ (*j* = 1, 2, ..., *N*) as a balanced sample from the Gaussian mixture

$$\hat{\pi}_{t|t} = \frac{1}{N} \sum_{j=1}^{N} \mathcal{N}(x_p^j + \hat{K}(y - Hx_p^j), \hat{K}R\hat{K}').$$

Hence the EnKF in step 1 without sampling gives

$$\hat{\pi}_{t|t}^{\gamma} = \frac{1}{N} \sum_{j=1}^{N} \mathcal{N}(x_{p}^{j} + \mathcal{K}(\gamma \hat{P}_{t|t-1})(y - \mathcal{H}x_{p}^{j}), \frac{1}{\gamma} \mathcal{K}(\gamma \hat{P}_{t|t-1})\mathcal{R}\mathcal{K}(\gamma \hat{P}_{t|t-1}))').$$

Bayes formula converts $\hat{\pi}_{t|t}^{\gamma}$ in the second step into another Gaussian mixture from which we sample to get $(x_{t|t}^{j})$.

Bridging between EnKF and PF: EnKPF

We can apply Bayes formula in two steps (progressive correction)

$$\pi_{t|t}(dx) \propto \pi_{t|t-1}(dx) p(y|x)^{\gamma} p(y|x)^{1-\gamma} \quad (0 \leq \gamma \leq 1).$$

Step 1: From $\pi_{t|t-1}(dx)$ to $\pi_{t|t}^{\gamma}(dx) \propto \pi_{t|t-1}(dx)p(y|x)^{\gamma}$ by PoEnKF:

Simply replace *R* by R/γ or $\hat{P}_{t|t-1}$ by $\gamma \hat{P}_{t|t-1}$ in the Kalman gain.

Step 2: From $\pi_{t|t}^{\gamma}(dx)$ to $\pi_{t|t}(dx) \propto \pi_{t|t}^{\gamma}(dx)p(y|x)^{1-\gamma}$ by particle filter.

The essential trick is to do both steps analytically, and to sample only at the end. See next slide.

Choice of tuning parameter γ

The exponent γ trades off systematic and Monte Carlo errors:

- $\gamma = 0$ corresponds to PF, $\gamma = 1$ corresponds to EnKF, with continuous interpolation in between.
- A small γ means small bias but large Monte Carlo variance.
- > A large γ means large bias but small Monte Carlo variance.

Need to find an interesting compromise!

Ansatz: Choose γ to achieve a given sampling diversity, i.e., the mixture weights in step 2 should not be too different from uniform weights.

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Diversity as a function of γ

Left/right: Two different values of *y*. Two priors: Gaussian (top) and bimodal (bottom). State dimension: 10, 50, 250.



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Single update for bimodal prior II

As before, but with observation leading to a bimodal posterior.



Single update for bimodal prior I

Left: EnKF, Right: EnKPF, diversity \approx 40%. Dots: Prior sample, Dotted: Contours of underlying update density.



Section 6

Introductory Examples

Basics of state space models and filtering

Kalman filter and its applications

Particle filters

Ensemble Kalman filters

Extensions of particle filters Particle smoothing Parameter estimation General sequential Monte Carlo

Smoothing by filtering of paths

We can implement the recursion

$$\begin{array}{lll} \pi_{0:t|t-1}(dx_{0:t}) & = & \mathcal{K}(dx_t \mid x_{t-1})\pi_{0:t-1|t-1}(dx_{0:t-1}) \\ \pi_{0:t|t}(dx_{0:t}) & \propto & \pi_{0:t|t-1}(dx_{0:t}) \times g(y_t \mid x_t). \end{array}$$

by a particle filter that generates samples of paths $(x_{0:t|t}^{j})$: Attach the propagated particle to the current path

$$x_{0:t|t-1}^{j} = (x_{0:t-1|t-1}^{j}, x_{t|t-1}^{j})$$

and resample $(x_{0:t|t-1}^{j})$ with probabilities $\propto g(y_t \mid x_{t|t-1}^{j})$.

But then $(x_{s|t}^{j})$ degenerates quickly to a single value for any fixed *s* since this component is not rejuvenated. So this method is useless for uncertainty quantification, but it still can generate a reasonable path from $\pi_{0:t|t}$.

Smoothing by accept/reject

We can get rid of both problems by using the approximation

$$\pi_{s|s} \propto g(y_s \mid x_s) \sum_{j=1}^N k(x_s \mid x_{s-1|s-1}^j)$$

instead of the empirical distribution of $(x_{t|t}^j)$. This means that we sample $x_{s|t}^j$ given $x_{s+1|t}^j$ from the density which is proportional to

$$k(x_{s+1|t}^{j} | x_{s})g(y_{s} | x_{s}) \sum_{i=1}^{N} k(x_{s} | x_{s-1|s-1}^{i}).$$

Since we need only one draw from this distribution, we have to use the accept/reject method: Propose an index I_* and a value $x_{s,*}$ according to $\tau(i)q(x_s \mid x_{s|s}^i)$ and accept $x_{s,*}$ as $x_{s|t}^j$ with the appropriate probability.

Forward filtering, backward smoothing

We can combine the formula

$$\begin{array}{rcl} \mathsf{P}(dx_{s} \mid x_{s+1}, y_{1:t}) & = & \mathsf{P}(dx_{s} \mid x_{s+1}, y_{1:s}) \\ & \propto & k(x_{s+1} \mid x_{s}) \pi_{s|s}(dx_{s}). \end{array}$$

with the particle filter approximations of $\pi_{s|s}$ to generate recursively an approximate sample from $\pi_{0:t|t}$:

$$\mathbb{P}(x_{s|t}^{j} = x_{s|s}^{i} \mid x_{s+1|t}^{j}) = \frac{k(x_{s+1|t}^{j} \mid x_{s|s}^{i})}{\sum_{\ell=1}^{N} k(x_{s+1|t}^{j} \mid x_{s|s}^{\ell})}$$

If $\pi_{s|t}$ is much more concentrated than $\pi_{s|s}$, diversity is too low. Moreover, sampling from this discrete distribution by inversion has complexity $O(n^2)$.

Smoothing using the two filter formula

A different approach is based on the formula

$$\pi_{s|t} \propto \pi_{s|s-1} p(y_{s:t} \mid x_s).$$

The second factor is unknown, but it satisfies the recursion

$$p(y_{s+1:t} \mid x_s) = \int p(y_{s+1:t} \mid x_{s+1})k(x_{s+1} \mid x_s)dx_{s+1}$$

$$p(y_{s:t} \mid x_s) = g(y_s \mid x_s)p(y_{s+1:t} \mid x_s).$$

This has the same structure as the filter relations, except that $p(y_{s:r} | x_s)$ is not a probability density in x_s (hence there is no normalization in the update step).

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Approximating the 2-filter formula

We can turn this into a recursion for densities by setting

$$\tilde{\pi}_{s|s}(dx_s) = \frac{p(y_{s:t} \mid x_s)\gamma_s(x_s)}{\int p(y_{s:t} \mid x_s)\gamma_s(x_s)dx_s}$$

where γ_s is a density s.th. the denominator is finite.

Then we start with a sample $(\tilde{x}_{t|t}^j)$ from $\tilde{\pi}_{t|t} \propto g(y_t \mid x_t)\gamma_t$ and construct recursively approximate samples $(\tilde{x}_{s|s}^j)$ from $\tilde{\pi}_{s|s}$ by propagation, weighting and resampling.

In this way we obtain an approximation of $\pi_{\textit{s}|\textit{t}}$ with density proportional to

$$\sum_{i=1}^{N} k(x_{s} \mid x_{s-1|s-1}^{i}) g(y_{s} \mid x_{s}) \sum_{j=1}^{N} \frac{k(\tilde{x}_{s+1|s+1}^{j} \mid x_{s})}{\gamma_{s+1}(\tilde{x}_{s+1|s+1}^{j})}$$

Parameters as components of the state

We discuss next the estimation of parameters θ in the state transition *K* and/or the observation density *g*.

The easiest method is to include θ as a deterministic component of the state:

 $\mathcal{K}(d\theta_t, dx_t \mid \theta_{t-1}, x_{t-1}) = \Delta_{\theta_{t-1}}(d\theta_t) \mathcal{K}(dx_t \mid x_{t-1}, \theta_{t-1})$

where Δ is a point mass. The particle filter degenerates quickly because there is no rejuvenation of the θ -component.

One can avoid this by adding noise to $\theta_{t|t-1}^{j}$, possibly combined with shrinkage to the mean. But we need to let the variance of the noise go to zero in order that $(\theta_{t|t}^{j})$ approximates $p(\theta \mid y_{1:t})$.

An O(N) smoothing algorithm

The density on the previous slide is a mixture of N^2 components. Like in the auxiliary particle filter we generate a weighted sample by sampling first the two indices of the mixture and then x_s from an approximation to the chosen component and finally by weighting.

If we try to use an approximately optimal choice of the two indices, we end up again by an $O(N^2)$ algorithm. However, we obtain an O(N) algorithm if we choose the two indices independently. The price we pay is a potential loss of effective sample size.

Particle MCMC

In the linear Gaussian case, we used simultaneous updates of parameters and states: We proposed

 $(x_{0:t}^*, \theta^*) \mid (x_{0:t}, \theta) \sim q(\theta^* \mid \theta) d\theta^* P(dx_{0:t} \mid y_{1:t}, \theta^*)$

and accept it with probability $\min(1, p(\theta^* \mid y_{1:t}) / p(\theta \mid y_{1:t}))$.

This cannot be used in general because we cannot draw true realizations from $P(dx_{0:t} | y_{1:t}, \theta)$, and for the acceptance probability we need $p(y_{1:t} | \theta)$ which we cannot compute exactly.

Andrieu et al. JRSS B (2010) showed that using particle approximations at both instances still gives a consistent algorithm.

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Sampling from moving targets

In many applications outside from state space models, one also wants to sample not from one distribution π , but from a sequence of related distributions π_t , t = 0, 1, ..., n.

In particular, if the target π is difficult to sample, one can start with a simpler distribution π_0 and construct an approximating sequence such that $\pi_n = \pi$. Examples are importance splitting as discussed in the Introduction, or simulated tempering where

$$\pi_t(dx) \propto \left(\frac{d\pi}{d\pi_0}(x)\right)^{\beta_t} \pi_0(dx) \quad (0 = \beta_0 < \beta_1 < \ldots < \beta_n = 1).$$

To simplify the notation, assume that all π_t have densities which are known up to a normalizing constant.

Resampling pairs of particles

- We cannot compute the density of $(x_{t,*}^j)$, but the density of $(x_{t-1}^j, x_{t,*}^j)$ is known: $\pi_{t-1}(x_{t-1})q_t(x_t \mid x_{t-1})$.
- Importance resampling can convert (x^j_{t-1}, x^j_{t,*}) to a sample from a distribution with second marginal π_t.
- Such distributions have the density $\pi_t(x_t)r_{t-1}(x_{t-1} \mid x_t)$ where r_{t-1} is an arbitrary transition density.
- By marginalization, we can choose any "backward transition" r_{t-1} and resample (x^j_{t,*}) with probabilities

$$w_t^j \propto \tilde{w}_t(x_{t-1}^j, x_{t,*}^j) = rac{\pi_t(x_{t,*}^j)r_{t-1}(x_{t-1}^j \mid x_{t,*}^j)}{\pi_{t-1}(x_{t-1}^j)q_t(x_{t,*}^j \mid x_{t-1}^j)}.$$

Sequential sampling

Generalizing the particle filter, we want to approximate π_t by a sequence of particles (x_t^j) which evolve by propagation and resampling.

If in the propagation step $x_{t,*}^j \sim q_t(x_t \mid x_{t-1}^j) dx_t$ (independently for different *j*'s), then resampling must be with probabilities

$$w_t^j = \propto \tilde{w}_t(x_{t,*}^j) = \frac{\pi_t(x_{t,*}^j)}{\int \pi_{t-1}(x_{t-1})q_t(x_{t,*}^j \mid x_{t-1})dx_{t-1}}$$

But typically, the integral in the denominator cannot be computed analytically. One exception is when q_t leaves π_{t-1} invariant, e.g. a Metropolis-Hastings transition. But then we move from π_{t-1} to π_t only by importance resampling, propagation just does some rejuvenation by breaking ties from the resampling step.

Choice of the transitions

We are free to choose forward and backward transitions q_t and r_{t-1} . For given q_t , the variance of \tilde{w}_t is minimal if

$$r_{t-1}(x_{t-1} \mid x_t) = \frac{\pi_{t-1}(x_{t-1})q_t(x_t \mid x_{t-1})}{\int \pi_{t-1}(x_{t-1})q_t(x_t \mid x_{t-1})dx_{t-1}},$$

bringing us back to the problem at the start. Still, one can approximate the optimal choice by something which does not involve an integral.

E.g. approximating $\log \pi_{t-1}(x_{t-1}) + \log q_t(x_t \mid x_{t-1})$ by a quadratic expression in x_{t-1} , will lead to a Gaussian backward density.

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The case of the particle filter

For the particle filter $\pi_t = \pi_{t|t}$ for which we do not have an explicit expression up to a normalizing constant. But if we choose the pair target density $\pi_t(x_t)r_{t-1}(x_{t-1} \mid x_t)$ to be

$$\pi_{t-1:t|t}(x_t, x_{t-1}) = \frac{\pi_{t-1}(x_{t-1})k(x_t \mid x_{t-1})g(y_t \mid x_t)}{p(y_t \mid y_{1:t-1})},$$

the weights are explicit up to a normalizing constant

$$ilde{w}_t(x_{t-1}, x_t) \propto rac{k(x_t \mid x_{t-1})g(y_t \mid x_t)}{q(x_t \mid x_{t-1}, y_t)}.$$

As stated earlier, the optimal choice of *q* is $p(x_t | x_{t-1}, y_t)$.