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Gaussian filtering based maximum likelihood and maximum a posteriori estimation in discrete-time state-space models

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1 Introduction

(Murphy 2002)

SSMs vs Box-Jenkins

Role of static parameters

Importance of estimating static parameters

Overview of different approaches

2 Background

2.1 State space models

State space models (SSMs) provide a unified probabilistic methodology for modeling sequential data (Ljung et al. 1994; Durbin et al. 2012; Cappé et al. 2005; Barber et al. 2011). Sequential data arise in numerous applications, typically in the form of time-series measurements. However it is not necessary for the sequence index to have a temporal meaning. In probabilistic terms a time-series can be described by a *stochastic process* $\mathbf{z} = \{\mathbf{z}_k : k \in K\}$, where \mathbf{z}_k is a random variable and $K \subset \mathbb{R}$ for continuous time or $K \subset \mathbb{N}$ for discrete time sequences. In this thesis we will only be concerned with discrete time processes and the sample space of \mathbf{z}_k will be \mathbb{R}^d . The shorthand $\mathbf{z}_{1:k}$ will be used to mean the subset $\{\mathbf{z}_1, \dots, \mathbf{z}_k\}$. We will also denote by \mathbf{Z} the $d \times T$ matrix, that has all T values of the process \mathbf{z} as columns.

examples

A fundamental question in probabilistic models for sequential data is how to model the dependence between variables. It is infeasible to assume that every random variable in the process depends on all the others. Thus it is common to assume a *Markov chain*, where the distribution of the process at the current timestep depends only on the distribution in the previous timestep. A further assumption in SSMs is that the process of interest, the dynamic process \mathbf{x} , is not directly observed but only through another stochastic process, the *measurement process* \mathbf{y} . Since \mathbf{x} is not observed, SSMs belong to the class of *latent variable models*. Sometimes, as in Cappé et al. (2005), SSMs are called *hidden Markov models* (HMM) but usually this implies that the sample space of \mathbf{x} is discrete. Another assumption is that the values of the measurement process are conditionally independent given the latent Markov process. An intuitive way to present conditional independence properties between random variables is a *Bayes network* presented by a directed acyclic graph (DAG) (Pearl 1988; Bishop 2006). A Bayes network presentation of a discrete-time SSM is given in figure 1.

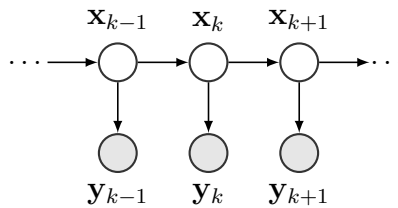


Figure 1: SSM as a graphical model presented with a directed acyclic graph

{fig:ssm_graph}

The value $\mathbf{x}_k \in \mathcal{X} \subset \mathbb{R}^{d_x}$ of the dynamic process at time k is called the *state* at time k . For the measurements we define $\mathbf{y}_k \in \mathcal{Y} \subset \mathbb{R}^{d_y}$. Taking into account the Markov property

Explain state

$$p(\mathbf{x}_k | \mathbf{x}_{1:k-1}) = p(\mathbf{x}_k | \mathbf{x}_{k-1}) \quad (1)$$

of the dynamic process and the conditional independence property

$$p(\mathbf{y}_k | \mathbf{x}_{1:k}, \mathbf{y}_{1:k-1}) = p(\mathbf{y}_k | \mathbf{x}_k) \quad (2)$$

of the measurement process, the joint distribution of states and measurements factorises as

$$p(\mathbf{X}, \mathbf{Y} | \boldsymbol{\theta}) = p(\mathbf{x}_0 | \boldsymbol{\theta}) \prod_{k=1}^T p(\mathbf{x}_k | \mathbf{x}_{k-1}, \boldsymbol{\theta}) p(\mathbf{y}_k | \mathbf{x}_k, \boldsymbol{\theta}). \quad (3) \quad \text{\texttt{\{eq:complete_}}}$$

Thus in order to describe a SSM one needs to specify three probability distributions:

Prior distribution $p(\mathbf{x}_0 | \boldsymbol{\theta})$ is the distribution assumed for the state prior to observing any measurements. The sensitivity of the posterior distributions to the prior depends on the amount of data (the more data the less sensitivity).

Dynamic model $p(\mathbf{x}_k | \mathbf{x}_{k-1}, \boldsymbol{\theta})$ dictates the time evolution of the states

Measurement model $p(\mathbf{y}_k | \mathbf{x}_k, \boldsymbol{\theta})$ models how the observations depend on the state and the statistics of the noise

In this thesis it is assumed that the parametric form of these distributions is known for example by physical modeling (Ljung et al. 1994). However the distributions are dependent on the vector parameter $\boldsymbol{\theta} \in \Theta \subset \mathbb{R}^{d_\theta}$ whose value (or distribution) we would like to learn from the measurements.

Traditionally SSMs are specified as a pair of equations specifying the dynamic and measurement models. The most general presentation of discrete-time SSMs is

$$\mathbf{x}_k = \mathbf{f}_{\boldsymbol{\theta},k}(\mathbf{x}_{k-1}, \mathbf{q}_{k-1}) \quad (4a)$$

$$\mathbf{y}_k = \mathbf{h}_{\boldsymbol{\theta},k}(\mathbf{x}_k, \mathbf{r}_k). \quad (4b)$$

Here the stochasticity is separated into the noise processes \mathbf{q} and \mathbf{r} which are usually assumed to be zero mean, white and independent of each other. We will restrict ourselves to the case of zero mean, white and additive Gaussian noise and in our case the mappings $\mathbf{f}_{\boldsymbol{\theta},k}$ and $\mathbf{h}_{\boldsymbol{\theta},k}$ and the noise processes \mathbf{q} and \mathbf{r} will be stationary (i.e. independent of k). Thus the SSMs considered in this thesis are of the form

$$\mathbf{x}_k = \mathbf{f}_{\boldsymbol{\theta}}(\mathbf{x}_{k-1}) + \mathbf{q}_{k-1} \quad \mathbf{q}_{k-1} \sim \mathcal{N}(\mathbf{0}, \mathbf{Q}) \quad (5a) \quad \text{\texttt{\{eq:ssm_gener\}}}$$

$$\mathbf{y}_k = \mathbf{h}_{\boldsymbol{\theta}}(\mathbf{x}_k) + \mathbf{r}_k \quad \mathbf{r}_k \sim \mathcal{N}(\mathbf{0}, \mathbf{R}) \quad (5b) \quad \text{\texttt{\{eq:Q\}}}$$

$$\mathbf{x}_0 \sim \mathcal{N}(\boldsymbol{\mu}_0, \boldsymbol{\Sigma}_0) \quad , \quad (5c) \quad \text{\texttt{\{eq:R\}}}$$

We will assume an implicit dependence of the distributional parameters \mathbf{Q} , \mathbf{R} , $\boldsymbol{\mu}_0$ and $\boldsymbol{\Sigma}_0$ on $\boldsymbol{\theta}$. Regarding the Gaussian distribution, suppose $\mathbf{x} \in \mathbb{R}^m$ is normally distributed with mean \mathbf{m} and covariance matrix \mathbf{P} . We will then use the notation

$$\mathbf{x} \sim \mathcal{N}(\mathbf{m}, \mathbf{P}) \quad (6)$$

$$\Leftrightarrow p(\mathbf{x} | \mathbf{m}, \mathbf{P}) = \mathcal{N}(\mathbf{x} ; \mathbf{m}, \mathbf{P}) \quad (7)$$

$$= (2\pi)^{-m/2} |\mathbf{P}|^{-1/2} e^{-1/2(\mathbf{x}-\mathbf{m})^T \mathbf{P}^{-1}(\mathbf{x}-\mathbf{m})} \quad (8) \quad \text{\texttt{\{eq:prior\}}}$$

Now clearly the mappings $\mathbf{f}_{\boldsymbol{\theta}} : \mathcal{X} \rightarrow \mathcal{X}$ and $\mathbf{h}_{\boldsymbol{\theta}} : \mathcal{X} \rightarrow \mathcal{Y}$ specify the means of the dynamic and the measurement models:

$$p(\mathbf{x}_k | \mathbf{x}_{k-1}, \boldsymbol{\theta}) = \mathcal{N}(\mathbf{f}_{\boldsymbol{\theta}}(\mathbf{x}_{k-1}), \mathbf{Q}) \quad (9a) \quad \text{\texttt{\{eq:ssm_distr\}}}$$

$$p(\mathbf{y}_k | \mathbf{x}_k, \boldsymbol{\theta}) = \mathcal{N}(\mathbf{h}_{\boldsymbol{\theta}}(\mathbf{x}_k), \mathbf{R}) \quad (9b) \quad \text{\texttt{\{eq:dynamics_}}}$$

\text{\texttt{\{eq:measurement\}}}

Example: 1D random walk

The simplest example is a one dimensional random-walk observed in Gaussian noise. We will assume $p(\mathbf{x}_0) = \mathcal{N}(0, P_0)$. In an alternative (but equivalent) notation the dynamics model is now

$$p(x_k | x_{k-1}) = x_{k-1} + q_{k-1}, \quad (10)$$

where $q_{k-1} \sim \mathcal{N}(0, Q)$ and the measurement model is

$$p(y_k | x_k) = x_k + r_k, \quad (11)$$

where $r_k \sim \mathcal{N}(0, R)$. A simulation from the model is presented in figure 2.

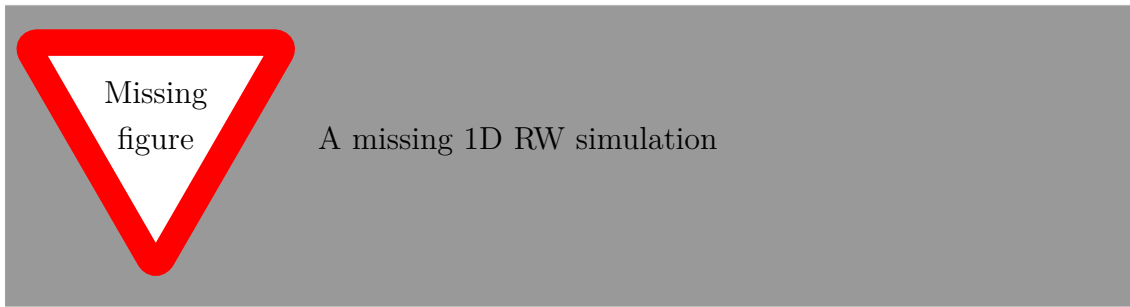


Figure 2: Simulation from the 1D RW model

{fig:rw1d}

2.2 Bayesian optimal filtering and smoothing

Inference can be defined as answering questions of interest with a probability distribution (Barber et al. 2011). In case of SSMS there are many questions of interest, but most commonly one would like to know the *marginal posterior distribution* of the states. State inference can be divided into subcategories based on the temporal relationship between the state and the observations (Särkkä 2006):

Predictive distribution $p(\mathbf{x}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta})$ is the predicted distribution of the state in the next timestep (or more generally at timestep $k + h$, where $h > 0$) given the previous measurements

Filtering distribution $p(\mathbf{x}_k | \mathbf{y}_{1:k}, \boldsymbol{\theta})$ is the marginal posterior distribution of any state \mathbf{x}_k given the measurements up to and including \mathbf{y}_k

Smoothing distribution $p(\mathbf{x}_k | \mathbf{y}_{1:T}, \boldsymbol{\theta})$ is the marginal posterior distribution of any state \mathbf{x}_k given the measurements up to and including \mathbf{y}_T where $k < T$

Predictive distribution

Let us now derive a recursive formulation for computing the filtering distribution at time k . Let $p(\mathbf{x}_{k-1} | \mathbf{y}_{1:k-1})$ be the filtering distribution of the previous step. Then

$$\begin{aligned} p(\mathbf{x}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) &= \int p(\mathbf{x}_k, \mathbf{x}_{k-1} | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) d\mathbf{x}_{k-1} \\ &= \int p(\mathbf{x}_k | \mathbf{x}_{k-1}) p(\mathbf{x}_{k-1} | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) d\mathbf{x}_{k-1}, \end{aligned} \quad (12) \quad \text{\texttt{eq:pred_bayes}}$$

which is known as the *Chapman-Kolmogorov equation* (Särkkä 2006). In this thesis the predictive distributions will be Gaussian or approximated with a Gaussian

$$p(\mathbf{x}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) = \mathcal{N}(\mathbf{x}_k; \mathbf{m}_{k|k-1}, \mathbf{P}_{k|k-1}) \quad (13)$$

Filtering distribution

Incorporating the newest measurement can be achieved with the Bayes' rule (see for example Gelman et al. 2004)

$$\begin{aligned} \underbrace{p(\mathbf{x}_k | \mathbf{y}_{1:k}, \boldsymbol{\theta})}_{\text{posterior}} &= \frac{\overbrace{p(\mathbf{y}_k | \mathbf{x}_k, \boldsymbol{\theta})}^{\text{likelihood}} \overbrace{p(\mathbf{x}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta})}^{\text{prior}}}{\underbrace{p(\mathbf{y}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta})}_{\text{marginal likelihood}}} \\ &= \frac{p(\mathbf{y}_k | \mathbf{x}_k) p(\mathbf{x}_k | \mathbf{y}_{1:k-1})}{\int p(\mathbf{y}_k | \mathbf{x}_k) p(\mathbf{x}_k | \mathbf{y}_{1:k-1}) d\mathbf{x}_k} \end{aligned} \quad (14)$$

check marginal
likelihood
wording

which is called the measurement update equation. In this thesis the filtering distributions will be Gaussian or approximated with a Gaussian

$$p(\mathbf{x}_k | \mathbf{y}_{1:k}, \boldsymbol{\theta}) = \mathcal{N}(\mathbf{x}_k; \mathbf{m}_{k|k}, \mathbf{P}_{k|k}) \quad (15)$$

Smoothing distribution

The smoothing distributions can also be computed recursively by assuming that the filtering distributions and the smoothing distribution $p(\mathbf{x}_{k+1} | \mathbf{y}_{1:T})$ of the “previous” step are available. Since

$$\begin{aligned} p(\mathbf{x}_k | \mathbf{x}_{k+1}, \mathbf{y}_{1:T}, \boldsymbol{\theta}) &= p(\mathbf{x}_k | \mathbf{x}_{k+1}, \mathbf{y}_{1:k}, \boldsymbol{\theta}) \\ &= \frac{p(\mathbf{x}_k, \mathbf{x}_{k+1} | \mathbf{y}_{1:k}, \boldsymbol{\theta})}{p(\mathbf{x}_{k+1} | \mathbf{y}_{1:k}, \boldsymbol{\theta})} \\ &= \frac{p(\mathbf{x}_{k+1} | \mathbf{x}_k, \boldsymbol{\theta}) p(\mathbf{x}_k | \mathbf{y}_{1:k}, \boldsymbol{\theta})}{p(\mathbf{x}_{k+1} | \mathbf{y}_{1:k}, \boldsymbol{\theta})} \end{aligned}$$

we get

$$p(\mathbf{x}_k | \mathbf{y}_{1:T}, \boldsymbol{\theta}) = p(\mathbf{x}_k | \mathbf{y}_{1:k}, \boldsymbol{\theta}) \int \left[\frac{p(\mathbf{x}_{k+1} | \mathbf{x}_k, \boldsymbol{\theta}) p(\mathbf{x}_{k+1} | \mathbf{y}_{1:T}, \boldsymbol{\theta})}{p(\mathbf{x}_{k+1} | \mathbf{y}_{1:k}, \boldsymbol{\theta})} \right] d\mathbf{x}_{k+1}, \quad (16)$$

where $p(\mathbf{x}_{k+1} | \mathbf{y}_{1:k})$ can be computed by equation (12). In this thesis the smoothing distributions will be Gaussian or approximated with a Gaussian

$$p(\mathbf{x}_k | \mathbf{y}_{1:T}) = \mathcal{N}(\mathbf{x}_k; \mathbf{m}_{k|T}, \mathbf{P}_{k|T}) \quad (17)$$

Marginal likelihood

An important quantity concerning parameter estimation is the marginal likelihood $p(\mathbf{Y} | \boldsymbol{\theta})$. If we're able to compute the distributions

$$p(\mathbf{y}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) = \int p(\mathbf{y}_k | \mathbf{x}_k, \boldsymbol{\theta}) p(\mathbf{x}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) d\mathbf{x}_k \quad (18)$$

then by repeatedly applying the definition of conditional probability we find that the marginal likelihood can be computed from

$$p(\mathbf{Y} | \boldsymbol{\theta}) = p(\mathbf{y}_1 | \boldsymbol{\theta}) \prod_{k=2}^T p(\mathbf{y}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) \quad (19) \quad \text{\texttt{eq:lh_factor}}$$

3 State estimation

3.1 Linear-Gaussian State Space Models

Since linear mappings can be described by matrices, stationary linear-Gaussian SSMs are described by the subset of SSMs of the form (5) where

$$\mathbf{f}_{\boldsymbol{\theta}}(\mathbf{x}_{k-1}) = \mathbf{A}\mathbf{x}_{k-1} \quad (20)$$

$$\mathbf{h}_{\boldsymbol{\theta}}(\mathbf{x}_k) = \mathbf{H}\mathbf{x}_k \quad (21)$$

3.1.1 Kalman filter

The recursions are as follows (Kalman et al. 1960; Jazwinski 2007):

Predict:

$$\mathbf{m}_{k|k-1} = \mathbf{A}\mathbf{m}_{k-1|k-1} \quad (22a)$$

$$\mathbf{P}_{k|k-1} = \mathbf{A}\mathbf{P}_{k-1|k-1}\mathbf{A}^T + \mathbf{Q} \quad (22b)$$

Update:

$$\mathbf{v}_k = \mathbf{y}_k - \mathbf{H}\mathbf{m}_{k|k-1} \quad (22c)$$

$$\mathbf{S}_k = \mathbf{H}\mathbf{P}_{k|k-1}\mathbf{H}^T + \mathbf{R} \quad (22d)$$

$$\mathbf{K}_k = \mathbf{P}_{k|k-1}\mathbf{H}^T\mathbf{S}_k^{-1} \quad (22e)$$

$$\mathbf{m}_{k|k} = \mathbf{m}_{k|k-1} + \mathbf{K}_k\mathbf{v}_k \quad (22f)$$

$$\mathbf{P}_{k|k} = \mathbf{P}_{k|k-1} - \mathbf{K}_k\mathbf{S}_k\mathbf{K}_k^T \quad (22g)$$

Explain something about linear-Gaussian SSMs

eq:Kalman filter
the Kalman filter

This includes the sufficient statistics for the T joint distributions

$$p(\mathbf{x}_k, \mathbf{y}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) = N \left(\begin{bmatrix} \mathbf{x}_k \\ \mathbf{y}_k \end{bmatrix} ; \begin{bmatrix} \mathbf{m}_{k|k-1} \\ \mathbf{H}\mathbf{m}_{k|k-1} \end{bmatrix}, \begin{bmatrix} \mathbf{P}_{k|k-1} & \mathbf{P}_{k|k-1}\mathbf{H}^T \\ \mathbf{H}\mathbf{P}_{k|k-1}^T & \mathbf{S}_k \end{bmatrix} \right) \quad (23) \quad \{\text{eq:joint_per}\}$$

3.1.2 Rauch-Tung-Striebel Smoother

The standard Rauch-Tung-Striebel (RTS) smoother gives the statistics $\mathbf{m}_{k|T}$ and $\mathbf{P}_{k|T}$ (Jazwinski 2007; Rauch et al. 1965).

$$\mathbf{J}_k = \mathbf{P}_{k|k} \mathbf{A}^T \mathbf{P}_{k|k+1}^{-1} \quad (24a)$$

$$\mathbf{m}_{k|T} = \mathbf{m}_{k|k} + \mathbf{J}_k (\mathbf{m}_{k+1|T} - \mathbf{m}_{k+1|k}) \quad (24b)$$

$$\mathbf{P}_{k|T} = \mathbf{P}_{k|k} + \mathbf{J}_k (\mathbf{P}_{k+1|T} - \mathbf{P}_{k+1|k}) \mathbf{J}_k^T \quad (24c)$$

$$p(\mathbf{x}_k, \mathbf{x}_{k-1} | \mathbf{Y}, \boldsymbol{\theta}) = N \left(\begin{bmatrix} \mathbf{x}_k \\ \mathbf{x}_{k-1} \end{bmatrix} ; \begin{bmatrix} \mathbf{m}_{k|T} \\ \mathbf{m}_{k-1|T} \end{bmatrix}, \begin{bmatrix} \mathbf{P}_{k|T} & \mathbf{P}_{k|T} \mathbf{J}_k^T \\ \mathbf{J}_k \mathbf{P}_{k|T} & \mathbf{P}_{k-1|T} \end{bmatrix} \right) \quad (25)$$

In (Elliott et al. 1999) a new kind of filter is presented that can compute (??) with only forward recursions.

Elaborate
on the RTS
smoother

3.2 Nonlinear-Gaussian SSMs

In the nonlinear case at least one of the mappings $\mathbf{f}_{\boldsymbol{\theta}}$ and $\mathbf{h}_{\boldsymbol{\theta}}$ in (5) is nonlinear. Unfortunately in this case computing the filtering distributions in closed form becomes intractable and one has to resort to approximations. Finding good approximations is however extremely valuable since a great many physically modeled dynamical phenomena could be formulated as nonlinear-Gaussian SSM.

The types of approximate filtering (and smoothing) solutions can be put into two categories: particle filtering (or sequential Monte Carlo), which is asymptotically exact but computationally demanding and different kinds of deterministic approximation methods that are analytically inexact but less computationally demanding. We will only focus on the latter category.

Mention EKF

3.2.1 Gaussian filtering and smoothing

One approach to forming Gaussian approximations is to assume a Gaussian probability density function with mean and variance that match the actual ones (Ito 2000; Särkkä 2006). Let

$$\mathbf{a} \sim N(\mathbf{m}, \boldsymbol{\Sigma}_a) \quad (26) \quad \{\text{eq:pa}\}$$

$$\mathbf{b} | \mathbf{a} \sim N(\mathbf{f}(\mathbf{a}), \boldsymbol{\Sigma}_{b|a}) \quad (27) \quad \{\text{eq:pb_given_a}\}$$

then

$$p(\mathbf{a}, \mathbf{b}) = N(\mathbf{m}, \boldsymbol{\Sigma}_a) N(\mathbf{f}(\mathbf{a}), \boldsymbol{\Sigma}_{b|a}) \quad (28) \quad \{\text{eq:joint_non}\}$$

is only Gaussian if $\mathbf{f}(\mathbf{a})$ is linear. Let the Gaussian approximation to (28) be

$$p\left(\begin{bmatrix} \mathbf{a} \\ \mathbf{b} \end{bmatrix}\right) \approx \mathcal{N}\left(\begin{bmatrix} \boldsymbol{\mu}_a \\ \boldsymbol{\mu}_b \end{bmatrix}, \begin{bmatrix} \boldsymbol{\Sigma}_{aa} & \boldsymbol{\Sigma}_{ab} \\ \boldsymbol{\Sigma}_{ba} & \boldsymbol{\Sigma}_{bb} \end{bmatrix}\right) \quad (29) \quad \text{\texttt{\{eq:joint_gau\}}}$$

Then since the marginal distributions of a Gaussian distribution are also Gaussian, we have to have

$$\boldsymbol{\mu}_a = \mathbf{m} \quad (30)$$

$$\boldsymbol{\Sigma}_{aa} = \boldsymbol{\Sigma}_a \quad (31)$$

$$\boldsymbol{\mu}_b = \int \mathbf{b} p(\mathbf{b}) \, d\mathbf{b} \quad (32) \quad \text{\texttt{\{eq:mub\}}}$$

$$\boldsymbol{\Sigma}_{bb} = \int (\mathbf{b} - \boldsymbol{\mu}_b)(\mathbf{b} - \boldsymbol{\mu}_b)^T p(\mathbf{b}) \, d\mathbf{b} \quad (33) \quad \text{\texttt{\{eq:varb\}}}$$

Both (32) and (33) can be written in terms of (26) and (27). To see this, let us rewrite (32) as

$$\begin{aligned} \boldsymbol{\mu}_b &= \int \mathbf{b} p(\mathbf{b}) \, d\mathbf{b} \\ &= \int \mathbf{b} \int p(\mathbf{b}|\mathbf{a}) p(\mathbf{a}) \, d\mathbf{a} \, d\mathbf{b} \\ &= \int \int \mathbf{b} p(\mathbf{b}|\mathbf{a}) \, d\mathbf{b} p(\mathbf{a}) \, d\mathbf{a} \\ &= \int \mathbf{f}(\mathbf{a}) \mathcal{N}(\mathbf{m}, \boldsymbol{\Sigma}_a) \, d\mathbf{a} \end{aligned} \quad (34) \quad \text{\texttt{\{eq:mean_int\}}}$$

and (33) as

$$\begin{aligned} \boldsymbol{\Sigma}_{bb} &= \int \mathbf{b} \mathbf{b}^T p(\mathbf{b}) \, d\mathbf{b} - \boldsymbol{\mu}_b \boldsymbol{\mu}_b^T \\ &= \int \mathbf{f}(\mathbf{a}) \mathbf{f}(\mathbf{a})^T p(\mathbf{a}) \, d\mathbf{a} - \boldsymbol{\mu}_b \boldsymbol{\mu}_b^T \\ &\quad + \int \int [(\mathbf{b} - \mathbf{f}(\mathbf{a}))(\mathbf{b} - \mathbf{f}(\mathbf{a}))^T] p(\mathbf{b}|\mathbf{a}) \, d\mathbf{b} p(\mathbf{a}) \, d\mathbf{a} \\ &= \int (\mathbf{f}(\mathbf{a}) - \boldsymbol{\mu}_b)(\mathbf{f}(\mathbf{a}) - \boldsymbol{\mu}_b)^T \mathcal{N}(\mathbf{m}, \boldsymbol{\Sigma}_a) \, d\mathbf{a} + \boldsymbol{\Sigma}_{b|a}. \end{aligned} \quad (35) \quad \text{\texttt{\{eq:var_int\}}}$$

Finally, the cross-covariance $\Sigma_{ab} = \Sigma_{ba}^T$ similarly reads

$$\begin{aligned}
\Sigma_{ab} &= \int \int (\mathbf{a} - \boldsymbol{\mu}_a)(\mathbf{b} - \boldsymbol{\mu}_b)^T p(\mathbf{a}, \mathbf{b}) \, d\mathbf{a} \, d\mathbf{b} \\
&= \int \int (\mathbf{a} - \boldsymbol{\mu}_a)(\mathbf{b} - \boldsymbol{\mu}_b)^T p(\mathbf{a}) p(\mathbf{b}|\mathbf{a}) \, d\mathbf{a} \, d\mathbf{b} \\
&= \int (\mathbf{a} - \boldsymbol{\mu}_a) \left(\int \mathbf{b} p(\mathbf{b}|\mathbf{a}) \, d\mathbf{b} - \boldsymbol{\mu}_b \right)^T p(\mathbf{a}) \, d\mathbf{a} \\
&= \int (\mathbf{a} - \mathbf{m})(\mathbf{f}(\mathbf{a}) - \boldsymbol{\mu}_b)^T N(\mathbf{m}, \Sigma_a) \, d\mathbf{a}
\end{aligned} \tag{36} \quad \text{\texttt{\{eq:cross_cov\}}}$$

To see how this idea can be used to form a Gaussian approximation to (??), let us rewrite (??) as

$$\begin{aligned}
p(\mathbf{x}_{k-1}, \mathbf{x}_k | \mathbf{Y}) &= p(\mathbf{x}_{k-1} | \mathbf{x}_k, \mathbf{Y}_{1:k-1}) p(\mathbf{x}_k | \mathbf{Y}) \\
&= \frac{p(\mathbf{x}_{k-1}, \mathbf{x}_k | \mathbf{Y}_{1:k-1}) p(\mathbf{x}_k | \mathbf{Y})}{p(\mathbf{x}_k | \mathbf{Y}_{1:k-1})},
\end{aligned} \tag{37} \quad \text{\texttt{\{eq:joint_smooth\}}}$$

where the dependance on the current estimate of the parameter $\hat{\boldsymbol{\theta}}_j$ is suppressed for clarity. Since the Gaussian approximation to (??) will be calculated by forward (filtering) and backward (smoothing) recursions, let us assume that we already have available the Gaussian approximation

$$p(\mathbf{x}_{k-1} | \mathbf{Y}_{1:k-1}, \hat{\boldsymbol{\theta}}_j) \approx N(\mathbf{m}_{k-1|k-1}, \mathbf{P}_{k-1|k-1}). \tag{38}$$

The Gaussian approximation to

$$p(\mathbf{x}_{k-1}, \mathbf{x}_k | \mathbf{Y}_{1:k-1}) = N(\mathbf{x}_k | \mathbf{f}(\mathbf{x}_{k-1}), \mathbf{Q}) p(\mathbf{x}_{k-1} | \mathbf{Y}_{1:k-1}) \tag{39}$$

is then given by application of equations (34), (35) and (36)

$$\mathbf{m}_{k|k-1} = \int \mathbf{f}(\mathbf{x}_{k-1}) N(\mathbf{x}_{k-1} | \mathbf{m}_{k-1|k-1}, \mathbf{P}_{k-1|k-1}) \, d\mathbf{x}_{k-1} \tag{40} \quad \text{\texttt{\{eq:prediction_m\}}}$$

$$\begin{aligned}
\mathbf{P}_{k|k-1} &= \int (\mathbf{f}(\mathbf{x}_{k-1}) - \mathbf{m}_{k|k-1})(\mathbf{f}(\mathbf{x}_{k-1}) - \mathbf{m}_{k|k-1})^T \\
&\quad N(\mathbf{x}_{k-1} | \mathbf{m}_{k-1|k-1}, \mathbf{P}_{k-1|k-1}) \, d\mathbf{x}_{k-1} + \mathbf{Q}
\end{aligned} \tag{41} \quad \text{\texttt{\{eq:prediction_P\}}}$$

$$\begin{aligned}
\mathbf{C}_{k-} &= \int (\mathbf{x}_{k-1} - \mathbf{m}_{k-1|k-1})(\mathbf{f}(\mathbf{x}_{k-1}) - \mathbf{m}_{k|k-1})^T \\
&\quad N(\mathbf{x}_{k-1} | \mathbf{m}_{k-1|T}, \mathbf{P}_{k-1|T}) \, d\mathbf{x}_{k-1}
\end{aligned} \tag{42} \quad \text{\texttt{\{eq:prediction_C\}}}$$

so that the approximation is

$$p(\mathbf{x}_{k-1}, \mathbf{x}_k | \mathbf{Y}_{1:k-1}, \hat{\boldsymbol{\theta}}_j) \approx N\left(\begin{bmatrix} \mathbf{m}_{k-1|k-1} \\ \mathbf{m}_{k|k-1} \end{bmatrix}, \begin{bmatrix} \mathbf{P}_{k-1|k-1} & \mathbf{C}_{k-} \\ \mathbf{C}_{k-}^T & \mathbf{P}_{k|k-1} \end{bmatrix}\right) \tag{43} \quad \text{\texttt{\{eq:joint_pred\}}}$$

In order to calculate (40) and (41) we also need a Gaussian approximation for the joint distribution of the current state and measurement given the previous measurements

$$\begin{aligned} p(\mathbf{x}_k, \mathbf{y}_k | \mathbf{Y}_{1:k-1}) &= N(\mathbf{y}_k | \mathbf{h}(\mathbf{x}_k), \mathbf{R}) p(\mathbf{x}_k | \mathbf{Y}_{1:k-1}) \\ &\approx N\left(\begin{bmatrix} \mathbf{m}_{k|k-1} \\ \boldsymbol{\mu}_k \end{bmatrix}, \begin{bmatrix} \mathbf{P}_{k|k-1} & \mathbf{C}_k \\ \mathbf{C}_k^T & \mathbf{S}_k \end{bmatrix}\right). \end{aligned} \quad (44) \quad \text{\texttt{\{eq:joint_upd\}}}$$

Applying equations (34), (35) and (36) again, we get

$$\boldsymbol{\mu}_k = \int \mathbf{h}(\mathbf{x}_k) N(\mathbf{x}_k | \mathbf{m}_{k|k-1}, \mathbf{P}_{k|k-1}) d\mathbf{x}_k \quad (45) \quad \text{\texttt{\{eq:update_me\}}}$$

$$\mathbf{S}_k = \int (\mathbf{h}(\mathbf{x}_k) - \boldsymbol{\mu}_k)(\mathbf{h}(\mathbf{x}_k) - \boldsymbol{\mu}_k)^T N(\mathbf{x}_k | \mathbf{m}_{k|k-1}, \mathbf{P}_{k|k-1}) d\mathbf{x}_k + \mathbf{R} \quad (46) \quad \text{\texttt{\{eq:update_va\}}}$$

$$\mathbf{C}_k = \int (\mathbf{x}_k - \mathbf{m}_{k|k-1})(\mathbf{h}(\mathbf{x}_k) - \boldsymbol{\mu}_k)^T N(\mathbf{x}_k | \mathbf{m}_{k|k-1}, \mathbf{P}_{k|k-1}) d\mathbf{x}_k \quad (47) \quad \text{\texttt{\{eq:update_co\}}}$$

and by using the well known formula for calculating the conditional distribution of jointly Gaussian variables we have

$$\mathbf{m}_{k|k} = \mathbf{m}_{k|k-1} + \mathbf{C}_k \mathbf{S}_k^{-1} (\mathbf{y}_k - \boldsymbol{\mu}_k) \quad (48) \quad \text{\texttt{\{eq:update_me\}}}$$

$$\mathbf{P}_{k|k} = \mathbf{P}_{k|k-1} - \mathbf{C}_k \mathbf{S}_k^{-1} \mathbf{C}_k^T. \quad (49) \quad \text{\texttt{\{eq:update_va\}}}$$

Again using the formula for the conditional of jointly Gaussian variables we get from (43)

$$p(\mathbf{x}_{k-1} | \mathbf{x}_k, \mathbf{Y}_{1:k-1}) \approx N(\mathbf{m}_2, \mathbf{P}_2) \quad (50)$$

$$\mathbf{G}_{k-1} = \mathbf{C}_{k-1} \mathbf{P}_{k|k-1}^{-1} \quad (51)$$

$$\mathbf{m}_2 = \mathbf{m}_{k-1|k-1} + \mathbf{G}_{k-1} (\mathbf{x}_k - \mathbf{m}_{k|k-1}) \quad (52)$$

$$\mathbf{P}_2 = \mathbf{P}_{k-1|k-1} - \mathbf{G}_{k-1} \mathbf{P}_{k|k-1} \mathbf{G}_{k-1}^T \quad (53)$$

and then finally we can write the Gaussian approximation to the joint distribution of consecutive states given all the measurements as

$$\begin{aligned} p(\mathbf{x}_{k-1}, \mathbf{x}_k | \mathbf{Y}) &= p(\mathbf{x}_{k-1} | \mathbf{x}_k, \mathbf{Y}_{1:k-1}) p(\mathbf{x}_k | \mathbf{Y}) \\ &\approx N\left(\begin{bmatrix} \mathbf{m}_{k-1|T} \\ \mathbf{m}_{k|T} \end{bmatrix}, \begin{bmatrix} \mathbf{P}_{k-1|T} & \mathbf{D}_k \\ \mathbf{D}_k^T & \mathbf{P}_{k|T} \end{bmatrix}\right) \end{aligned} \quad (54)$$

where

$$\mathbf{D}_k = \mathbf{G}_{k-1} \mathbf{P}_{k|T} \quad (55)$$

$$\mathbf{m}_{k-1|T} = \mathbf{m}_{k-1|k-1} + \mathbf{G}_{k-1} (\mathbf{m}_{k|T} - \mathbf{m}_{k|k-1}) \quad (56)$$

$$\mathbf{P}_{k-1|T} = \mathbf{P}_{k-1|k-1} + \mathbf{G}_{k-1} (\mathbf{P}_{k|T} - \mathbf{P}_{k|k-1}) \mathbf{G}_{k-1}^T \quad (57)$$

3.2.2 Quadrature and cubature

(Arasaratnam et al. 2009)

3.2.3 Gauss-Hermite Kalman Filter and Smoother

(Ito 2000)

3.2.4 Unscented Kalman Filter and Smoother

(Julier et al. 1997; Merwe 2004)

3.2.5 Cubature Kalman Filter and Smoother

(Arasaratnam et al. 2009; Arasaratnam et al. 2011; Jia et al. 2012)

4 Parameter estimation

4.1 Maximum likelihood and maximum a posteriori estimation

In the Bayesian sense the complete answer to the parameter estimation problem is the marginal posterior probability of the parameters given the measurements, which is given by Bayes' rule as

$$p(\boldsymbol{\theta}|\mathbf{Y}) = \frac{p(\mathbf{Y}|\boldsymbol{\theta})p(\boldsymbol{\theta})}{p(\mathbf{Y})}. \quad (58) \quad \{\text{eq:param_pos}\}$$

Computing the posterior distribution of the parameters is usually intractable. A much easier problem is finding a suitable *point estimate* $\hat{\boldsymbol{\theta}}$. This effectively means that we don't need to worry about the normalizing term $p(\mathbf{Y})$, since it's constant with respect to the parameters. A point estimate that maximizes the posterior distribution is called a *maximum a posteriori* (MAP) estimate. Since the logarithm is a strictly monotonic function, maximizing a function is the same as maximizing its logarithm. Thus the MAP estimate $\boldsymbol{\theta}^*$ is given by

why? examples?

$$\boldsymbol{\theta}_{\text{MAP}} = \operatorname{argmax}_{\boldsymbol{\theta}} \left[\underbrace{\log p(\mathbf{Y}|\boldsymbol{\theta})}_{\ell(\boldsymbol{\theta})} + \log p(\boldsymbol{\theta}) \right] \quad (59) \quad \{\text{eq:MAP}\}$$

In the case of a flat (constant and thus improper) prior distribution, $p(\boldsymbol{\theta})$, the MAP estimate converges to the *maximum likelihood* (ML) estimate

$$\boldsymbol{\theta}_{\text{ML}} = \operatorname{argmax}_{\boldsymbol{\theta}} [\ell(\boldsymbol{\theta})] \quad (60) \quad \{\text{eq:ML}\}$$

Going further we will only be concerned with finding the ML estimate, but it should be remembered that both of the methods we consider can be extended to the estimation of the MAP estimate in a straightforward fashion.

Explain MAP in both cases

Among different point estimates, the maximum likelihood estimator has good statistical properties. Let us denote the true parameter value, the value that the data was generated with, with $\boldsymbol{\theta}_\star$ and let T denote the amount of observations. Then provided that some conditions of not very restricting nature hold, we can state the following asymptotic properties for the ML estimate $\boldsymbol{\theta}_{\text{ML}}$:

Modify to reflect p.465 in Cappé

Strong consistency

An important property for an estimator, which says that the estimator tends to the true value as the amount of data tends to infinity:

$$\ell_T(\boldsymbol{\theta}_{\text{ML}}) \xrightarrow{\text{a.s.}} \ell(\boldsymbol{\theta}_\star), \quad \text{when } T \rightarrow \infty, \quad (61)$$

where ℓ_T is the likelihood function after T measurements and ℓ is a continuous deterministic function with a unique global maximum at $\boldsymbol{\theta}_\star$.

Asymptotic normality

This property gives us the means to compute asymptotic error bounds for the estimate:

$$\sqrt{T}(\boldsymbol{\theta}_{\text{ML}} - \boldsymbol{\theta}_\star) \xrightarrow{D} \mathcal{N}(\mathbf{0}, \mathcal{I}^{-1}(\boldsymbol{\theta}_\star)), \quad \text{when } T \rightarrow \infty, \quad (62) \quad \{\text{tablelabel}\}$$

where $\mathcal{I}(\boldsymbol{\theta}_\star)$ is the *Fischer information matrix* evaluated at $\boldsymbol{\theta}_\star$

Efficiency

When the amount of information tends to infinity, the ML-estimate achieves the Cramér-Rao lower bound, i.e. no other consistent estimator has lower asymptotic mean-squared-error.

4.1.1 Identifiability

Intuitively, any parameters $\boldsymbol{\theta}, \boldsymbol{\theta}' \in \Theta$ cannot be distinguished from each other with maximum likelihood estimation if

$$p(\mathbf{Y}|\boldsymbol{\theta}) = p(\mathbf{Y}|\boldsymbol{\theta}'), \quad (63)$$

i.e., if the same data can arise with two (or more) separate parameter values. (Haykin 2001; Cappé et al. 2005)

Elaborate on identifiability

4.2 Gradient based nonlinear optimization

{sec:grad}

This is the classical way of solving the parameter estimation problem. It consists of computing the gradient of the log-likelihood function $\ell(\boldsymbol{\theta})$ and then using some non-linear optimization method to find a *local* maximum to it (Mbalawata et al. 2011; Cappé et al. 2005). An efficient non-linear optimization algorithm is the scaled conjugate gradient method (Mbalawata et al. 2011).

By marginalizing the joint distribution of equation (23) we get

$$p(\mathbf{y}_k | \mathbf{y}_{1:k-1}, \boldsymbol{\theta}) = \mathcal{N}(\mathbf{y}_k; \mathbf{H}\mathbf{m}_{k|k-1}, \mathbf{S}_k). \quad (64)$$

Applying equation (19) and taking the logarithm then gives

$$\ell(\boldsymbol{\theta}) = -\frac{1}{2} \sum_{k=1}^T \log |\mathbf{S}_k| - \frac{1}{2} \sum_{k=1}^T (\mathbf{y}_k - \mathbf{H}\mathbf{m}_{k|k-1})^T \mathbf{S}_k^{-1} (\mathbf{y}_k - \mathbf{H}\mathbf{m}_{k|k-1}) + C, \quad (65) \quad \{\text{eq:logLH}\}$$

where C is a constant that doesn't depend on $\boldsymbol{\theta}$ and thus can be ignored in the maximization. Employing an efficient numerical optimization method generally requires that the gradient of the objective function is available. There are at least two seemingly quite different methods for computing the gradient of $\ell(\boldsymbol{\theta})$. The first one proceeds straightforwardly by taking the partial derivatives of $\ell(\boldsymbol{\theta})$. As will soon be demonstrated, this leads to some additional recursive formulas which allow computing the gradient in parallel with the Kalman filter. The second method needs the smoothing distributions with the cross-timestep covariances of equation (??) and it can be easily computed with the expectation maximization machinery that will be introduced later. These two methods can be proved to compute the exact same quantity. At this point we will focus on the first one.

In order to calculate the gradient of $\ell(\boldsymbol{\theta})$, we can take the partial derivatives of it w.r.t every parameter θ_i in $\boldsymbol{\theta}$:

$$\begin{aligned} \frac{\partial \ell(\boldsymbol{\theta})}{\partial \theta_i} = & -\frac{1}{2} \sum_{k=1}^T \text{Tr} \left(\mathbf{S}_k^{-1} \frac{\partial \mathbf{S}_k}{\partial \theta_i} \right) \\ & + \sum_{k=1}^T \left(\mathbf{H}_k \frac{\partial \mathbf{m}_{k|k-1}}{\partial \theta_i} \right)^T \mathbf{S}_k^{-1} (\mathbf{y}_k - \mathbf{H}\mathbf{m}_{k|k-1}) \\ & + \frac{1}{2} \sum_{k=1}^T (\mathbf{y}_k - \mathbf{H}\mathbf{m}_{k|k-1})^T \mathbf{S}_k^{-1} \left(\frac{\partial \mathbf{S}_k}{\partial \theta_i} \right) \mathbf{S}_k^{-1} (\mathbf{y}_k - \mathbf{H}\mathbf{m}_{k|k-1}) \end{aligned} \quad (66) \quad \{\text{eq:dlogLH}\}$$

From the Kalman filter recursions (22) we find out that

$$\frac{\partial \mathbf{S}_k}{\partial \theta_i} = \mathbf{H} \frac{\partial \mathbf{P}_{k|k-1}}{\partial \theta_i} \mathbf{H}^T + \frac{\partial \mathbf{R}}{\partial \theta_i} \quad (67)$$

so that we're left with the task of determining the partial derivatives for $\mathbf{m}_{k|k-1}$ and $\mathbf{P}_{k|k-1}$:

$$\frac{\partial \mathbf{m}_{k|k-1}}{\partial \theta_i} = \frac{\partial \mathbf{A}}{\partial \theta_i} \mathbf{m}_{k-1|k-1} + \mathbf{A} \frac{\partial \mathbf{m}_{k-1|k-1}}{\partial \theta_i} \quad (68) \quad \{\text{eq:m_pred_pd}\}$$

$$\begin{aligned} \frac{\partial \mathbf{P}_{k|k-1}}{\partial \theta_i} = & \frac{\partial \mathbf{A}}{\partial \theta_i} \mathbf{P}_{k-1|k-1} \mathbf{A}^T + \mathbf{A} \frac{\partial \mathbf{P}_{k-1|k-1}}{\partial \theta_i} \mathbf{A}^T \\ & + \mathbf{A} \mathbf{P}_{k-1|k-1} \left(\frac{\partial \mathbf{A}}{\partial \theta_i} \right)^T + \frac{\partial \mathbf{Q}}{\partial \theta_i} \end{aligned} \quad (69) \quad \{\text{eq:P_pred_pd}\}$$

as well as for $\mathbf{m}_{k|k}$ and $\mathbf{P}_{k|k}$:

$$\frac{\partial \mathbf{K}_k}{\partial \theta_i} = \frac{\partial \mathbf{P}_{k|k-1}}{\partial \theta_i} \mathbf{H}^T \mathbf{S}_k^{-1} - \mathbf{P}_{k|k-1} \mathbf{H}^T \mathbf{S}_k^{-1} \frac{\partial \mathbf{S}_k}{\partial \theta_i} \mathbf{S}_k^{-1} \quad (70) \quad \{\text{eq:K_pd}\}$$

$$\frac{\partial \mathbf{m}_{k|k}}{\partial \theta_i} = \frac{\partial \mathbf{m}_{k|k-1}}{\partial \theta_i} + \frac{\partial \mathbf{K}_k}{\partial \theta_i} (\mathbf{y}_k - \mathbf{H} \mathbf{m}_{k|k-1}) - \mathbf{K}_k \mathbf{H} \frac{\partial \mathbf{m}_{k|k-1}}{\partial \theta_i} \quad (71) \quad \{\text{eq:m_pd}\}$$

$$\frac{\partial \mathbf{P}_{k|k}}{\partial \theta_i} = \frac{\partial \mathbf{P}_{k|k-1}}{\partial \theta_i} - \frac{\partial \mathbf{K}_k}{\partial \theta_i} \mathbf{S}_k \mathbf{K}_k^T - \mathbf{K}_k \frac{\partial \mathbf{S}_k}{\partial \theta_i} \mathbf{K}_k^T - \mathbf{K}_k^T \mathbf{S}_k \left(\frac{\partial \mathbf{K}_k}{\partial \theta_i} \right)^T \quad (72) \quad \{\text{eq:P_pd}\}$$

Equations (68), (69), (70), (71) and (72) together specify a recursive algorithm for computing (66) that can be run alongside the Kalman filter recursions. As noted in Cappé et al. (2005), these equations are sometimes known as the *sensitivity equations* and they are derived at least in Gupta et al. (1974) and Mbalawata et al. (2011).

Elaborate on nonlinear programming

4.3 Expectation maximization (EM)

The expectation maximization (EM) algorithm (Dempster et al. 1977) is a general method for finding ML and MAP estimates in probabilistic models with missing data or latent variables (Bishop 2006; Barber 2012). As will be seen, instead of maximizing (65) directly, EM alternates between forming a variational lower bound and maximizing it. We shall use $\langle \cdot \rangle_q \equiv \int \cdot q(z) dz$ to denote the expectation over some arbitrary distribution $q(z)$. Let us introduce a “variational” distribution $\tilde{p}(\mathbf{X}|\boldsymbol{\theta}')$ over the states, parameterized with $\boldsymbol{\theta}'$ (not necessarily related to $\boldsymbol{\theta}$). Toting now that $p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}) = p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta})/p(\mathbf{Y}|\boldsymbol{\theta})$ and that $\ell(\boldsymbol{\theta}) \equiv \log p(\mathbf{Y}|\boldsymbol{\theta})$ is independent of \mathbf{X} we can then perform the following decomposition on the log likelihood:

$$\begin{aligned} \ell(\boldsymbol{\theta}) &= \log p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}) - \log p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}) \\ &= \langle \log p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}) \rangle_{\tilde{p}} - \langle \log p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}) \rangle_{\tilde{p}} \\ &= \underbrace{\left\langle \log p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}) - \log \tilde{p}(\mathbf{X}|\boldsymbol{\theta}') \right\rangle_{\tilde{p}}}_{\mathcal{L}(\tilde{p}, \boldsymbol{\theta})} - \underbrace{\left\langle \log p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}) - \log \tilde{p}(\mathbf{X}|\boldsymbol{\theta}') \right\rangle_{\tilde{p}}}_{\text{KL}(\tilde{p}||p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}))} \quad (73) \quad \{\text{eq:1LH_decom}\} \end{aligned}$$

The important step here is taking the expectation over $\tilde{p}(\mathbf{X}|\boldsymbol{\theta}')$, since the *complete-data log-likelihood* $\log p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta})$ cannot be evaluated as \mathbf{X} is unobserved. Since $\text{KL}(\tilde{p}||p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}))$, the *Kullback-Leibler divergence* between $\tilde{p}(\mathbf{X}|\boldsymbol{\theta}')$ and $p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta})$, is always nonnegative, we see that

$$\ell(\boldsymbol{\theta}) \geq \mathcal{L}(\tilde{p}, \boldsymbol{\theta}) \quad (74) \quad \{\text{eq:em_lh_lb}\}$$

with equality when

$$\tilde{p}(\mathbf{X}|\boldsymbol{\theta}') = p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}), \quad (75) \quad \{\text{eq:em_maxq}\}$$

i.e. the posterior distribution of the states with equal parameter value $\boldsymbol{\theta}' = \boldsymbol{\theta}$. Considered as a functional of only \tilde{p} , clearly $\mathcal{L}(\tilde{p}, \boldsymbol{\theta})$ is maximized and $\text{KL}(\tilde{p}||p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}))$

vanishes by (75). The nonnegativeness of the Kullback-Leibler divergence can be proved by noting that $-\log$ is a convex function and so *Jensen's inequality* can be applied (Bishop 2006).

Let us take a closer at the the first term in (73) with $\tilde{p}(\mathbf{X}|\boldsymbol{\theta}') = p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}')$:

$$\mathcal{L}\left(p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}'), \boldsymbol{\theta}\right) = \underbrace{\left\langle \log p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}) \right\rangle_{p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}')}}_{\mathfrak{L}(\boldsymbol{\theta}', \boldsymbol{\theta})} - \left\langle \log p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}') \right\rangle_{p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}')} . \quad (76) \quad \{\text{eq:completed}\}$$

Clearly the latter term (the differential entropy of $p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}')$) is constant with respect to $\boldsymbol{\theta}$, so that maximizing \mathcal{L} with respect to $\boldsymbol{\theta}$ amounts to maximizing $\mathfrak{L}(\boldsymbol{\theta}', \boldsymbol{\theta})$, the *expected complete-data log-likelihood*, with respect to $\boldsymbol{\theta}$.

We are now ready to define the EM algorithm, which produces a series of estimates $\{\boldsymbol{\theta}_j\}$ to the parameter $\boldsymbol{\theta}$ starting from an initial guess $\boldsymbol{\theta}_0$. The two alternating steps of the algorithm are:

E-step

Given the current estimate $\boldsymbol{\theta}_j$ of the parameters, compute

$$\tilde{p}_{j+1} = \arg \max_{\tilde{p}} \mathcal{L}(\tilde{p}, \boldsymbol{\theta}_j) . \quad (77) \quad \{\text{eq:EM_E}\}$$

As stated, the maximum is obtained with $\tilde{p}_{j+1} = p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}_j)$, the posterior distribution of the states given the current parameter estimate. After the maximization we have

$$\begin{aligned} \ell(\boldsymbol{\theta}_j) &= \mathcal{L}\left(p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}_j), \boldsymbol{\theta}_j\right) \\ &= \mathfrak{L}(\boldsymbol{\theta}_j, \boldsymbol{\theta}_j) + \text{const} \end{aligned} \quad (78) \quad \{\text{eq:EM_E_lh}\}$$

M-step

Set

$$\begin{aligned} \boldsymbol{\theta}_{j+1} &= \arg \max_{\boldsymbol{\theta}} \mathcal{L}(\tilde{p}_{j+1}, \boldsymbol{\theta}) \\ &= \arg \max_{\boldsymbol{\theta}} \mathfrak{L}(\boldsymbol{\theta}_j, \boldsymbol{\theta}) . \end{aligned} \quad (79) \quad \{\text{eq:EM_M}\}$$

We are now in a position to formulate the so called *fundamental inequality of EM* (Cappé et al. 2005):

$$\ell(\boldsymbol{\theta}_{j+1}) - \ell(\boldsymbol{\theta}_j) \geq \mathcal{L}\left(p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}_j), \boldsymbol{\theta}_{j+1}\right) - \mathcal{L}\left(p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}_j), \boldsymbol{\theta}_j\right) \quad (80) \quad \{\text{eq:fundament}\}$$

which is just the combination of (74) and (78). But it highlights the fact that *the likelihood is increased or unchanged with every new estimate $\boldsymbol{\theta}_{j+1}$* . Also following from (80) is the fact that if the iterations stop at a certain point, i.e. $\boldsymbol{\theta}_{l+1} = \boldsymbol{\theta}_l$ at iteration l , then $\mathcal{L}(p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}_l), \boldsymbol{\theta})$ must be maximal at $\boldsymbol{\theta}_l$ and so the gradients of the lower bound and of the likelihood must be zero. Thus $\boldsymbol{\theta}_l$ is a *stationary point* of $\ell(\boldsymbol{\theta})$, i.e a local maximum or a saddle point.

4.3.1 EM as a special case of variational Bayes

(Barber 2012; Jordan 1998) Variational Bayes (VB) is a fully Bayesian methodology where one seeks for an approximation to the parameter posterior

Ensure that notation matches prev chapter

$$p(\boldsymbol{\theta}|\mathbf{Y}) \propto \int_{\mathcal{X}} p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}) d\mathbf{X} p(\boldsymbol{\theta}) \quad (81) \quad \text{\texttt{\{tablelabel\}}}$$

As mentioned earlier, finding this distribution is commonly intractable, so in VB we assume a factorized form for the joint posterior of states and parameters

$$p(\mathbf{X}, \boldsymbol{\theta}|\mathbf{Y}) \approx q(\mathbf{X}) q(\boldsymbol{\theta}) \quad (82) \quad \text{\texttt{\{eq:VB_factor\}}}$$

and the task is then to find the best approximation with respect to the KL divergence between the true posterior and the approximation

$$\text{KL}(q(\mathbf{X}) q(\boldsymbol{\theta}) \| p(\mathbf{X}, \boldsymbol{\theta}|\mathbf{Y})) = \langle q(\mathbf{X}) \rangle_{q(\mathbf{X})} + \langle q(\mathbf{X}) \rangle_{q(\mathbf{X})} - \langle p(\mathbf{X}, \boldsymbol{\theta}|\mathbf{Y}) \rangle_{q(\mathbf{X})q(\boldsymbol{\theta})}. \quad (83) \quad \text{\texttt{\{eq:KL_VB\}}}$$

Using $p(\mathbf{X}, \boldsymbol{\theta}|\mathbf{Y}) = p(\mathbf{X}, \boldsymbol{\theta}, \mathbf{Y}) / p(\mathbf{Y})$ equation (83) gives

$$\ell(\boldsymbol{\theta}) \geq \langle p(\mathbf{X}, \boldsymbol{\theta}, \mathbf{Y}) \rangle_{q(\mathbf{X})q(\boldsymbol{\theta})} - \langle q(\mathbf{X}) \rangle_{q(\mathbf{X})} - \langle q(\mathbf{X}) \rangle_{q(\mathbf{X})} \quad (84) \quad \text{\texttt{\{eq:VB_bound\}}}$$

and thus minimizing the KL divergence is equivalent to finding the tightest lower bound to the log likelihood. Analogously to EM, minimizing the KL divergence is done iteratively keeping $q(\boldsymbol{\theta})$ fixed and minimizing w.r.t $q(\mathbf{X})$ in the “E”-step and vice versa in the “M”-step:

E-step

$$q^{\text{new}}(\mathbf{X}) = \arg \max_{q(\mathbf{X})} \left(\text{KL} \left(q(\mathbf{X}) q^{\text{old}}(\boldsymbol{\theta}) \| p(\mathbf{X}, \boldsymbol{\theta}|\mathbf{Y}) \right) \right) \quad (85) \quad \text{\texttt{\{eq:VB_E\}}}$$

M-step

$$q^{\text{new}}(\boldsymbol{\theta}) = \arg \max_{q(\boldsymbol{\theta})} \left(\text{KL} \left(q^{\text{new}}(\mathbf{X}) q(\boldsymbol{\theta}) \| p(\mathbf{X}, \boldsymbol{\theta}|\mathbf{Y}) \right) \right) \quad (86) \quad \text{\texttt{\{eq:VB_M\}}}$$

Let us then suppose that we only wish to find the MAP point estimate $\boldsymbol{\theta}^*$. This can be accomplished by assuming a delta function form $q(\boldsymbol{\theta}) = \delta(\boldsymbol{\theta}, \boldsymbol{\theta}^*)$ for the parameter factor in the joint distribution of states and parameters (82). With this assumption equation (84) becomes

$$p(\mathbf{Y}|\boldsymbol{\theta}^*) \geq \langle p(\mathbf{X}, \boldsymbol{\theta}^*, \mathbf{Y}) \rangle_{q(\mathbf{X})q(\boldsymbol{\theta})} - \langle q(\mathbf{X}) \rangle_{q(\mathbf{X})} + \text{const} \quad (87) \quad \text{\texttt{\{eq:VB_MAP_bo\}}}$$

and the “M”-step (86) can then be written as

$$\boldsymbol{\theta}^* = \arg \max_{\boldsymbol{\theta}} \left(\langle \log p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}) \rangle_{q(\mathbf{X})} + \log p(\boldsymbol{\theta}) \right). \quad (88) \quad \text{\texttt{\{tablelabel\}}}$$

If the point estimate is plugged in the “E”-step equation (85) we have

$$q^{\text{new}}(\mathbf{X}) \propto p(\mathbf{X}, \mathbf{Y}|\boldsymbol{\theta}^*) \propto p(\mathbf{X}|\mathbf{Y}, \boldsymbol{\theta}^*) \quad (89) \quad \text{\texttt{\{tablelabel\}}}$$

4.3.2 Partial E and M steps

4.4 EM in SSMs

Let us then look at how to apply EM to a SSM of the form (5). First of all, from the factorization in (3), the complete-data log-likelihood becomes

{sec:EM_SSM}

$$\begin{aligned}
 p(\mathbf{X}, \mathbf{Y} | \boldsymbol{\theta}) = & -\frac{1}{2} (\mathbf{x}_0 - \boldsymbol{\mu}_0)^\top \boldsymbol{\Sigma}_0^{-1} (\mathbf{x}_0 - \boldsymbol{\mu}_0) - \frac{1}{2} \log |\boldsymbol{\Sigma}_0| \\
 & - \frac{1}{2} \sum_{k=1}^T (\mathbf{x}_k - \mathbf{f}(\mathbf{x}_{k-1}))^\top \mathbf{Q}^{-1} (\mathbf{x}_k - \mathbf{f}(\mathbf{x}_{k-1})) - \frac{T}{2} \log |\mathbf{Q}| \\
 & - \frac{1}{2} \sum_{k=1}^T (\mathbf{y}_k - \mathbf{h}(\mathbf{x}_k))^\top \mathbf{R}^{-1} (\mathbf{y}_k - \mathbf{h}(\mathbf{x}_k)) - \frac{T}{2} \log |\mathbf{R}| \\
 & + \text{const}
 \end{aligned}$$

Taking the expectation w.r.t. $p(\mathbf{X} | \mathbf{Y}, \boldsymbol{\theta}')$ (assumed implicitly in the notation), applying the identity $\mathbf{a}^\top \mathbf{C} \mathbf{b} = \text{Tr}[\mathbf{a}^\top \mathbf{C} \mathbf{b}] = \text{Tr}[\mathbf{C} \mathbf{b} \mathbf{a}^\top]$, denoting $\mathbf{f}_{k-1} \equiv \mathbf{f}(\mathbf{x}_{k-1})$ and $\mathbf{h}_k \equiv \mathbf{h}(\mathbf{x}_k)$ and dropping the constant terms we get

$$\begin{aligned}
 \mathcal{L}(\boldsymbol{\theta}', \boldsymbol{\theta}) = & -\frac{1}{2} \text{Tr} \left[\boldsymbol{\Sigma}_0^{-1} \left\langle (\mathbf{x}_0 - \boldsymbol{\mu}_0) (\mathbf{x}_0 - \boldsymbol{\mu}_0)^\top \right\rangle \right] - \frac{1}{2} \log |\boldsymbol{\Sigma}_0| \\
 & - \frac{1}{2} \text{Tr} \left[\mathbf{Q}^{-1} \sum_{k=1}^T \left\langle (\mathbf{x}_k - \mathbf{f}_{k-1}) (\mathbf{x}_k - \mathbf{f}_{k-1})^\top \right\rangle \right] - \frac{T}{2} \log |\mathbf{Q}| \\
 & - \frac{1}{2} \text{Tr} \left[\mathbf{R}^{-1} \sum_{k=1}^T \left\langle (\mathbf{y}_k - \mathbf{h}_k) (\mathbf{y}_k - \mathbf{h}_k)^\top \right\rangle \right] - \frac{T}{2} \log |\mathbf{R}|
 \end{aligned} \tag{90}$$

{eq:ec1LH}

Let us denote the three expectations in equation (90) with

$$\mathbf{I}_1 = \left\langle (\mathbf{x}_0 - \boldsymbol{\mu}_0) (\mathbf{x}_0 - \boldsymbol{\mu}_0)^\top \right\rangle \tag{91}$$

$$= \int_{\mathcal{X} \times T} (\mathbf{x}_0 - \boldsymbol{\mu}_0) (\mathbf{x}_0 - \boldsymbol{\mu}_0)^\top p(\mathbf{X} | \mathbf{Y}, \boldsymbol{\theta}') d\mathbf{X}$$

$$= \int_{\mathcal{X}} (\mathbf{x}_0 - \boldsymbol{\mu}_0) (\mathbf{x}_0 - \boldsymbol{\mu}_0)^\top p(\mathbf{x}_0 | \mathbf{Y}, \boldsymbol{\theta}') d\mathbf{x}_0 \tag{92}$$

{eq:I1_general}

$$\mathbf{I}_2 = \sum_{k=1}^T \int_{\mathcal{X} \times 2} (\mathbf{x}_k - \mathbf{f}_{k-1}) (\mathbf{x}_k - \mathbf{f}_{k-1})^\top p \left(\begin{bmatrix} \mathbf{x}_k^\top & \mathbf{x}_{k-1}^\top \end{bmatrix}^\top \middle| \mathbf{Y}, \boldsymbol{\theta}' \right) d \begin{bmatrix} \mathbf{x}_k & \mathbf{x}_{k-1} \end{bmatrix}^\top \tag{93}$$

{eq:I2_general}

$$\mathbf{I}_3 = \sum_{k=1}^T \int_{\mathcal{X}} (\mathbf{y}_k - \mathbf{h}_k) (\mathbf{y}_k - \mathbf{h}_k)^\top p(\mathbf{x}_k | \mathbf{Y}, \boldsymbol{\theta}') d\mathbf{x}_k \tag{94}$$

{eq:I3_general}

It is clear then that in the E-step one needs to compute the $T + 1$ smoothing distributions, including the T cross-timestep distributions, since these will be needed

in the expectations. By applying the identity

$$\text{var}[\mathbf{x}] = \langle \mathbf{x}\mathbf{x}^T \rangle - \langle \mathbf{x} \rangle \langle \mathbf{x} \rangle^T, \quad (95)$$

we can already write the first expectation as

$$\mathbf{I}_1 = \mathbf{P}_{0|T} + (\mathbf{m}_{0|T} - \boldsymbol{\mu}_0)(\mathbf{m}_{0|T} - \boldsymbol{\mu}_0)^T. \quad (96) \quad \{\text{eq:I1}\}$$

This was a result of assuming the Gaussian prior distribution of equation (5c) and it is common for the more specific models we're assessing in the next chapters.

Since in this thesis we're assuming Gaussian noise, the M-step maximization equations for the complete noise covariance matrices \mathbf{Q} and \mathbf{R} will also be common. To derive these equations, we will derivate (90) with respect to these matrices. As can be seen from (90), the terms involving \mathbf{Q} or \mathbf{R} are analogous and so we will only the derivation for \mathbf{Q} and only the result for \mathbf{R} . It's easier to take the derivative with respect to \mathbf{Q}^{-1} :

$$\begin{aligned} \frac{\partial \mathcal{L}(\boldsymbol{\theta}', \boldsymbol{\theta})}{\partial \mathbf{Q}^{-1}} &= -\frac{1}{2} \frac{\partial}{\partial \mathbf{Q}^{-1}} \text{Tr} \left[\mathbf{Q}^{-1} \sum_{k=1}^T \left\langle (\mathbf{x}_k - \mathbf{f}_{k-1}) (\mathbf{x}_k - \mathbf{f}_{k-1})^T \right\rangle \right] \\ &\quad - \frac{T}{2} \frac{\partial}{\partial \mathbf{Q}^{-1}} \log |\mathbf{Q}| \\ &= -\frac{1}{2} \sum_{k=1}^T \left\langle (\mathbf{x}_k - \mathbf{f}_{k-1}) (\mathbf{x}_k - \mathbf{f}_{k-1})^T \right\rangle + \frac{T}{2} \mathbf{Q}, \end{aligned} \quad (97) \quad \{\text{eq:c1LH_pdQ}\}$$

where we have used formula 92 in Petersen et al. 2008 for the first derivative and formula 51 for the second. Setting (97) to zero we get the update equation for the next estimate of \mathbf{Q}

Why is it same for the inverse?

$$\begin{aligned} \mathbf{Q}_{j+1} &= \frac{1}{T} \sum_{k=1}^T \left\langle (\mathbf{x}_k - \mathbf{f}_{k-1}) (\mathbf{x}_k - \mathbf{f}_{k-1})^T \right\rangle \\ &= \frac{1}{T} \mathbf{I}_2 \end{aligned} \quad (98) \quad \{\text{eq:EM_M_Q}\}$$

The derivation of the update equation for the next estimate of \mathbf{R} is exactly analogous, giving

$$\begin{aligned} \mathbf{R}_{j+1} &= \frac{1}{T} \sum_{k=1}^T \left\langle (\mathbf{y}_k - \mathbf{h}_k) (\mathbf{y}_k - \mathbf{h}_k)^T \right\rangle \\ &= \frac{1}{T} \mathbf{I}_3 \end{aligned} \quad (99) \quad \{\text{eq:EM_M_R}\}$$

In the following more specific cases, we will only consider the two remaining expectations and the rest of the M-step maximization equations.

Let us then discuss a certain important aspect of the M-step maximization equations, with a consequence that is useful also for gradient based methods for ML.

Often times one is not interested in estimating any *full* matrix, with which we mean that the model might have been parameterized so that there are only scalar parameters embedded inside matrices or as part of \mathbf{f} or \mathbf{h} . In that case it is quite likely that the M-step maximization equations will not admit themselves to closed form analytical expressions. Rather the use of nonlinear optimization methods is needed and for that we would like have the gradient of \mathfrak{L} . This is fortunately rather straightforward. Let θ_i be an element of $\boldsymbol{\theta}$. Then

$$\begin{aligned}
\frac{\partial \mathfrak{L}(\boldsymbol{\theta}, \boldsymbol{\theta}')}{\partial \theta_i} = & \frac{1}{2} \text{Tr} \left[\boldsymbol{\Sigma}^{-1} \left(\frac{\partial \boldsymbol{\Sigma}}{\partial \theta_i} \boldsymbol{\Sigma}^{-1} \sum_{k=1}^T \left\langle (\mathbf{x}_0 - \boldsymbol{\mu}_0) (\mathbf{x}_0 - \boldsymbol{\mu}_0)^T \right\rangle_{\hat{\boldsymbol{\theta}}_j} \right. \right. \\
& \left. \left. + 2 \sum_{k=1}^T \left\langle \frac{\partial \boldsymbol{\mu}_0}{\partial \theta_i} (\mathbf{x}_0 - \boldsymbol{\mu}_0)^T \right\rangle - \frac{\partial \boldsymbol{\Sigma}}{\partial \theta_i} \right) \right] \\
& + \frac{1}{2} \text{Tr} \left[\mathbf{Q}^{-1} \left(\frac{\partial \mathbf{Q}}{\partial \theta_i} \mathbf{Q}^{-1} \sum_{k=1}^T \left\langle (\mathbf{x}_k - \mathbf{f}_{k-1}) (\mathbf{x}_k - \mathbf{f}_{k-1})^T \right\rangle \right. \right. \\
& \left. \left. + 2 \sum_{k=1}^T \left\langle \frac{\partial \mathbf{f}_{k-1}}{\partial \theta_i} (\mathbf{x}_k - \mathbf{f}_{k-1})^T \right\rangle - T \frac{\partial \mathbf{Q}}{\partial \theta_i} \right) \right] \\
& + \frac{1}{2} \text{Tr} \left[\mathbf{R}^{-1} \left(\frac{\partial \mathbf{R}}{\partial \theta_i} \mathbf{R}^{-1} \sum_{k=1}^T \left\langle (\mathbf{y}_k - \mathbf{h}_k) (\mathbf{y}_k - \mathbf{h}_k)^T \right\rangle \right. \right. \\
& \left. \left. + 2 \sum_{k=1}^T \left\langle \frac{\partial \mathbf{h}_k}{\partial \theta_i} (\mathbf{y}_k - \mathbf{h}_k)^T \right\rangle - T \frac{\partial \mathbf{R}}{\partial \theta_i} \right) \right].
\end{aligned} \tag{100} \quad \text{\texttt{\{eq:dLB_nonli}}}$$

What is interesting about this quantity is that it can also be used to compute the gradient of the log-likelihood, i.e the score, itself. From equation (73) it can be seen rather easily, that the partial derivative of the log-likelihood evaluated at $\hat{\boldsymbol{\theta}}$ is given by

$$\left. \frac{\partial \ell(\boldsymbol{\theta})}{\partial \theta_i} \right|_{\boldsymbol{\theta}=\hat{\boldsymbol{\theta}}} = \left. \frac{\partial \mathcal{L} \left(p(\mathbf{X} | \mathbf{Y}, \hat{\boldsymbol{\theta}}), \boldsymbol{\theta} \right)}{\partial \theta_i} \right|_{\boldsymbol{\theta}=\hat{\boldsymbol{\theta}}} = \left. \frac{\partial \mathfrak{L}(\hat{\boldsymbol{\theta}}, \boldsymbol{\theta})}{\partial \theta_i} \right|_{\boldsymbol{\theta}=\hat{\boldsymbol{\theta}}} \tag{101} \quad \text{\texttt{\{eq:EM_gradie}}}$$

Equation (101) is known as *Fisher's identity* (Cappé et al. 2005). It gives an alternative route for the score function computation.

elaborate on
score computa-
tion

4.4.1 EM in linear-Gaussian SSMs

(Shumway et al. 1982; Ghahramani 1996) Let us substitute $\mathbf{A}\mathbf{x}_{k-1}$ for \mathbf{f}_{k-1} and $\mathbf{H}\mathbf{x}_k$ for \mathbf{h}_k . Let us also denote by

$$p(\mathbf{x}_k, \mathbf{x}_{k-1} | \mathbf{Y}, \boldsymbol{\theta}) = \mathcal{N} \left(\begin{bmatrix} \mathbf{x}_k \\ \mathbf{x}_{k-1} \end{bmatrix} ; \mathbf{m}_{k,k-1|T}, \mathbf{P}_{k,k-1|T} \right), \quad (102) \quad \{\text{eq:pdf_smoot}\}$$

the joint smoothing distribution of \mathbf{x}_k and \mathbf{x}_{k-1} . Then by applying the manipulation

$$\begin{aligned} & \left\langle (\mathbf{x}_k - \mathbf{A}\mathbf{x}_{k-1}) (\mathbf{x}_k - \mathbf{A}\mathbf{x}_{k-1})^T \right\rangle \\ &= \begin{bmatrix} \mathbf{I} & -\mathbf{A} \end{bmatrix} \left\langle \begin{bmatrix} \mathbf{x}_k \\ \mathbf{x}_{k-1} \end{bmatrix} \begin{bmatrix} \mathbf{x}_k^T & \mathbf{x}_{k-1}^T \end{bmatrix} \right\rangle \begin{bmatrix} \mathbf{I} \\ -\mathbf{A}^T \end{bmatrix} \end{aligned} \quad (103)$$

we get

$$\begin{aligned} \mathbf{I}_2 &= \begin{bmatrix} \mathbf{I} & -\mathbf{A} \end{bmatrix} \sum_{k=1}^T \left(\mathbf{P}_{k,k-1|T} + \mathbf{m}_{k,k-1|T} \mathbf{m}_{k,k-1|T}^T \right) \begin{bmatrix} \mathbf{I} \\ -\mathbf{A}^T \end{bmatrix} \\ &= \begin{bmatrix} \mathbf{I} & -\mathbf{A} \end{bmatrix} \begin{bmatrix} \sum_{k=1}^T \langle \mathbf{x}_k \mathbf{x}_k^T \rangle & \sum_{k=1}^T \langle \mathbf{x}_k \mathbf{x}_{k-1}^T \rangle \\ \sum_{k=1}^T \langle \mathbf{x}_{k-1} \mathbf{x}_k^T \rangle & \sum_{k=1}^T \langle \mathbf{x}_{k-1} \mathbf{x}_{k-1}^T \rangle \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{A}^T \end{bmatrix} \\ &= \begin{bmatrix} \mathbf{I} & -\mathbf{A} \end{bmatrix} \begin{bmatrix} \bar{\mathbf{X}}_{00} & \bar{\mathbf{X}}_{01} \\ \bar{\mathbf{X}}_{01}^T & \bar{\mathbf{X}}_{11} \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{A}^T \end{bmatrix} \\ &= \bar{\mathbf{X}}_{00} - \mathbf{A} \bar{\mathbf{X}}_{01}^T - \bar{\mathbf{X}}_{01} \mathbf{A}^T + \mathbf{A} \bar{\mathbf{X}}_{11} \mathbf{A}^T \\ &= (\mathbf{A} - \bar{\mathbf{X}}_{01} \bar{\mathbf{X}}_{11}^{-1}) \bar{\mathbf{X}}_{11} (\mathbf{A} - \bar{\mathbf{X}}_{01} \bar{\mathbf{X}}_{11}^{-1})^T + \bar{\mathbf{X}}_{00} + \bar{\mathbf{X}}_{01} \bar{\mathbf{X}}_{11}^{-1} \bar{\mathbf{X}}_{01}^T. \end{aligned} \quad (104) \quad \{\text{eq:LGSSM_I2}\}$$

It's easy to see that the extremum value of the last line with respect to \mathbf{A} is obtained by setting

$$\mathbf{A}_{j+1} = \bar{\mathbf{X}}_{01} \bar{\mathbf{X}}_{11}^{-1}. \quad (105) \quad \{\text{eq:Amax}\}$$

For \mathbf{I}_3 we get

$$\mathbf{I}_3 = \begin{bmatrix} \mathbf{I} & -\mathbf{H} \end{bmatrix} \sum_{k=1}^T \begin{bmatrix} \mathbf{y}_k \mathbf{y}_k^T & \mathbf{y}_k \langle \mathbf{x}_k \rangle^T \\ \langle \mathbf{x}_k \rangle \mathbf{y}_k^T & \langle \mathbf{x}_k \mathbf{x}_k^T \rangle \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{H}^T \end{bmatrix} \quad (106) \quad \{\text{eq:LGSSM_I3}\}$$

$$= \begin{bmatrix} \mathbf{I} & -\mathbf{H} \end{bmatrix} \sum_{k=1}^T \begin{bmatrix} \bar{\mathbf{Y}}_{00} & \bar{\mathbf{C}}_{00} \\ \bar{\mathbf{C}}_{00}^T & \bar{\mathbf{X}}_{00} \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{H}^T \end{bmatrix}, \quad (107)$$

so with analogous manipulation as in (105) we get

$$\mathbf{H}_{j+1} = \bar{\mathbf{C}}_{00} \bar{\mathbf{X}}_{00}^{-1}. \quad (108) \quad \{\text{eq:EM_M_H}\}$$

All in all, the E-step of EM algorithm in linear-Gaussian SSM:s consists of computing the T joint distributions in equation (102) with the RTS smoother. Actually

this is not the only option, as for example in Elliott et al. (1999) a new kind of filter is presented that can compute the expectations with only forward recursions. After this, the M-step estimates are computed for \mathbf{Q} from equation (98), for \mathbf{R} from equation (99), for \mathbf{A} from equation (106) and for \mathbf{H} from equation (109). Also for evaluating the convergence, one needs to compare the current value of \mathcal{L} with the previous one. For this we can compute \mathbf{I}_1 from equation (96), \mathbf{I}_2 from equation (104) and \mathbf{I}_3 from equation (115).

index I:s

For working with structured matrices let us rewrite the gradient equation (100) in the linear-Gaussian case:

$$\begin{aligned}
\frac{\partial \mathcal{L}(\boldsymbol{\theta}, \boldsymbol{\theta}')}{\partial \theta_i} = & \frac{1}{2} \text{Tr} \left[\boldsymbol{\Sigma}^{-1} \left(\frac{\partial \boldsymbol{\Sigma}}{\partial \theta_i} \boldsymbol{\Sigma}^{-1} \sum_{k=1}^T \left\langle (\mathbf{x}_0 - \boldsymbol{\mu}_0) (\mathbf{x}_0 - \boldsymbol{\mu}_0)^T \right\rangle_{\hat{\boldsymbol{\theta}}_j} \right. \right. \\
& \left. \left. + 2 \sum_{k=1}^T \left\langle \frac{\partial \boldsymbol{\mu}_0}{\partial \theta_i} (\mathbf{x}_0 - \boldsymbol{\mu}_0)^T \right\rangle - \frac{\partial \boldsymbol{\Sigma}}{\partial \theta_i} \right) \right] \\
& + \frac{1}{2} \text{Tr} \left[\mathbf{Q}^{-1} \left(\frac{\partial \mathbf{Q}}{\partial \theta_i} \mathbf{Q}^{-1} \begin{bmatrix} \mathbf{I} & -\mathbf{A} \end{bmatrix} \begin{bmatrix} \bar{\mathbf{X}}_{00} & \bar{\mathbf{X}}_{01} \\ \bar{\mathbf{X}}_{01}^T & \bar{\mathbf{X}}_{11} \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{A}^T \end{bmatrix} \right. \right. \\
& \left. \left. + 2 \begin{bmatrix} \mathbf{0} & -\frac{\partial \mathbf{A}}{\partial \theta_i} \end{bmatrix} \begin{bmatrix} \bar{\mathbf{X}}_{00} & \bar{\mathbf{X}}_{01} \\ \bar{\mathbf{X}}_{01}^T & \bar{\mathbf{X}}_{11} \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{A}^T \end{bmatrix} - T \frac{\partial \mathbf{Q}}{\partial \theta_i} \right) \right] \\
& + \frac{1}{2} \text{Tr} \left[\mathbf{R}^{-1} \left(\frac{\partial \mathbf{R}}{\partial \theta_i} \mathbf{R}^{-1} \begin{bmatrix} \mathbf{I} & -\mathbf{H} \end{bmatrix} \begin{bmatrix} \bar{\mathbf{Y}}_{00} & \bar{\mathbf{C}}_{00} \\ \bar{\mathbf{C}}_{00}^T & \bar{\mathbf{X}}_{00} \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{H}^T \end{bmatrix} \right. \right. \\
& \left. \left. + 2 \begin{bmatrix} \mathbf{0} & -\frac{\partial \mathbf{H}}{\partial \theta_i} \end{bmatrix} \begin{bmatrix} \bar{\mathbf{Y}}_{00} & \bar{\mathbf{C}}_{00} \\ \bar{\mathbf{C}}_{00}^T & \bar{\mathbf{X}}_{00} \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{H}^T \end{bmatrix} - T \frac{\partial \mathbf{R}}{\partial \theta_i} \right) \right].
\end{aligned} \tag{110}$$

{eq:dLB_linear}

4.4.2 EM in nonlinear-Gaussian SSMs

As explained in section 3.2, in the nonlinear case the filtering and smoothing distributions cannot be computed exactly. Thus the E-step is also only approximate and the convergence guarantees of EM won't apply anymore. The proposed methods can nevertheless provide good results most of the time. In the fortunate case that the model is linear-in-the-parameters the M-step can be solved in closed form. This situation will be covered later in section 4.4.3. Currently we will assume however that the model is nonlinear in the parameters as well as in the states so that the simplest form to write the model is exactly (5).

The E-step in the EM-algorithm consists now of computing approximations to \mathbf{I}_1 , \mathbf{I}_2 and \mathbf{I}_3 from equations (96), (93) and (94). In the M-step one should a gradient

based nonlinear optimization algorithm to compute the next parameter estimate (79). The gradient can be computed with the equation (100).

4.4.3 EM in linear-in-the-parameters SSM:s

Suppose now that $\mathbf{f}(\mathbf{x}) : \mathcal{X} \rightarrow \mathcal{X}$ is a linear combination of vector valued functions $\boldsymbol{\rho}_k : \mathcal{X} \rightarrow \mathbb{R}^{d_{\Phi,k}}$, so that the parameters of \mathbf{f} , $\boldsymbol{\Phi}_k$, are matrices of size $d_x \times d_{\Phi,k}$. Then \mathbf{f} can be written as

$$\begin{aligned} \mathbf{f}(\mathbf{x}) &= \boldsymbol{\Phi}_1 \boldsymbol{\rho}_1(\mathbf{x}) + \cdots + \boldsymbol{\Phi}_m \boldsymbol{\rho}_m(\mathbf{x}) \\ &= \begin{bmatrix} \boldsymbol{\Phi}_1 & \cdots & \boldsymbol{\Phi}_m \end{bmatrix} \begin{bmatrix} \boldsymbol{\rho}_1(\mathbf{x}) \\ \vdots \\ \boldsymbol{\rho}_m(\mathbf{x}) \end{bmatrix} \\ &= \mathbf{A} \mathbf{g}(\mathbf{x}), \end{aligned} \tag{111}$$

so that \mathbf{A} is now a matrix of size $d_x \times \sum_{k=1}^m d_{\Phi,k}$ and $\mathbf{g} : \mathcal{X} \rightarrow \mathbb{R}^{\sum_{k=1}^m d_{\Phi,k}}$. Denoting $\mathbf{g}(\mathbf{x}_{k-1}) \equiv \mathbf{g}_{k-1}$ and $\mathbf{l}(\mathbf{x}_k) \equiv \mathbf{l}_k$ and following the derivation in equation (104) we have

$$\begin{aligned} \mathbf{I}_2 &= \begin{bmatrix} \mathbf{I} & -\mathbf{A} \end{bmatrix} \begin{bmatrix} \sum_{k=1}^T \langle \mathbf{x}_k \mathbf{x}_k^T \rangle & \sum_{k=1}^T \langle \mathbf{x}_k \mathbf{g}_{k-1}^T \rangle \\ \sum_{k=1}^T \langle \mathbf{g}_{k-1} \mathbf{x}_k^T \rangle & \sum_{k=1}^T \langle \mathbf{g}_{k-1} \mathbf{g}_{k-1}^T \rangle \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{A}^T \end{bmatrix} \\ &= \begin{bmatrix} \mathbf{I} & -\mathbf{A} \end{bmatrix} \begin{bmatrix} \bar{\mathbf{X}}_{00} & \bar{\mathbf{G}}_{01} \\ \bar{\mathbf{G}}_{01}^T & \bar{\mathbf{G}}_{11} \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{A}^T \end{bmatrix} \end{aligned} \tag{112}$$

Then similarly to (106)

$$\mathbf{A}_{j+1} = \bar{\mathbf{G}}_{01} \bar{\mathbf{G}}_{11}^{-1}. \tag{113}$$

Applying similar assumptions as for \mathbf{f} to \mathbf{h} , we can write

$$\begin{aligned} \mathbf{h}(\mathbf{x}) &= \boldsymbol{\Upsilon}_1 \boldsymbol{\pi}_1(\mathbf{x}) + \cdots + \boldsymbol{\Upsilon}_m \boldsymbol{\pi}_m(\mathbf{x}) \\ &= \begin{bmatrix} \boldsymbol{\Upsilon}_1 & \cdots & \boldsymbol{\Upsilon}_m \end{bmatrix} \begin{bmatrix} \boldsymbol{\pi}_1(\mathbf{x}) \\ \vdots \\ \boldsymbol{\pi}_m(\mathbf{x}) \end{bmatrix} \\ &= \mathbf{H} \mathbf{l}(\mathbf{x}), \end{aligned} \tag{114}$$

where the size of \mathbf{H} is now $d_x \times \sum_{k=1}^m d_{\Upsilon,k}$ and $\mathbf{l} : \mathcal{X} \rightarrow \mathbb{R}^{\sum_{k=1}^m d_{\Upsilon,k}}$. For \mathbf{I}_3 we get

$$\mathbf{I}_3 = \begin{bmatrix} \mathbf{I} & -\mathbf{H} \end{bmatrix} \sum_{k=1}^T \begin{bmatrix} \mathbf{y}_k \mathbf{y}_k^T & \mathbf{y}_k \langle \mathbf{l}_k \rangle^T \\ \langle \mathbf{l}_k \rangle \mathbf{y}_k^T & \langle \mathbf{l}_k \mathbf{l}_k^T \rangle \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{H}^T \end{bmatrix} \tag{115}$$

$$= \begin{bmatrix} \mathbf{I} & -\mathbf{H} \end{bmatrix} \sum_{k=1}^T \begin{bmatrix} \bar{\mathbf{Y}}_{00} & \bar{\mathbf{D}}_{00} \\ \bar{\mathbf{D}}_{00}^T & \bar{\mathbf{L}}_{00} \end{bmatrix} \begin{bmatrix} \mathbf{I} \\ -\mathbf{H}^T \end{bmatrix}, \tag{116}$$

and

$$\mathbf{H}_{j+1} = \bar{\mathbf{D}}_{00} \bar{\mathbf{L}}_{00}^{-1}. \tag{117}$$

4.5 Properties of the estimation algorithms

Both the direct method of section 4.2 and the EM algorithm of section 4.4 have their strengths and weaknesses. Neither of them can be said to eclipse the other in an absolute sense. In this section we will go through some features of the said algorithms found in the literature. Then a more detailed analysis will be performed on two essential aspects of any estimation algorithm: the convergence properties, i.e. when should one expect the algorithm to find a local maximum and the computational requirements of the algorithms.

Cappé et al. (2005) contains a list of arguments in favor of either of the methods. The following list includes those and additional points with comments:

Direct

- *No smoother needed* The log-likelihood, or an approximation to it, can be evaluated with forward-filtering
- *No M-step*. There is no need to figure out model-dependent maximization formulas even in nonlinear models
- *Faster convergence*. Advanced gradient-based optimization methods can reach convergence speeds that are close to quadratic

EM

- *Simple to implement*. This argument is often put forward in favor of the EM algorithm. However in practice when using the gradient-based method one would use any one of the off-the-shelf nonlinear optimizers and not re-implement one. Thus which one is easier to implement boils down to gradient computation. If the model is linear or linear-in-the-parameters the EM algorithm doesn't need any gradient information.
- *Parameter constraints*. In Cappé et al. (2005) it is argued that since the M-step maximization equations are so simple, including parameter constraints are easier in the EM algorithm. This again depends on if the model is linear or linear-in-the-parameters.
- *Parameterization independent*. This again depends on if one has to use gradient-based optimization in the M-step. If not, then the EM algorithm is parameterization independent. In the gradient-based method the gradient and the Hessian, and so the convergence, are affected by the parameterization.

4.5.1 Limit points

4.5.2 Convergence rates

Let us denote the mapping implicitly defined by the EM algorithm by $\mathbf{M} : \Theta \rightarrow \Theta$. Then if the iterates $\boldsymbol{\theta}_j$ converge to a point $\boldsymbol{\theta}_*$, we must have $\boldsymbol{\theta}_* = \mathbf{M}(\boldsymbol{\theta}_*)$.

4.5.3 Computational load

(Harvey 1990; Watson 1983; Cappé et al. 2005; Saatci 2011; Olsson et al. 2007; Salakhutdinov et al. 2003)

5 Results

5.1 Tracking a ballistic object on reentry

(Ristic et al. 2004)

5.2 fMRI signal component analysis

(Särkkä et al. 2012)

6 Discussion

6.1 Stability

(Haykin 2001)

6.2 Dual and joint filtering

(Haykin 2001)

6.3 Particle filtering approaches

(Kantas et al. 2009; Doucet et al. 2001; Lindsten 2010)

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