

# Overview over Dynamical Statistical Models – with a view towards application in biological sciences

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## 1 Introduction

This paper is intended to give an overview over some aspects of dynamical statistical models (hereafter abbreviated DSMs) with a view towards applications in biological sciences. The

specific view we have is to applications where data are measured online and where interest is utilizing data as they arrive (as opposed to waiting until a whole batch of data is available for an analysis).

To set the scene, we let  $y_t$  denote a measurement at time  $t$ . The observations available up to including time  $t$  is denoted either as  $D_t$  or as  $y_{1:t}$ . Observations available up to but not including time  $t$  are denoted  $D_{<t}$  or  $y_{<t}$ . In the models we consider there are unknown parameters, generally denoted by  $\theta$ . Following this, we are interested in models in which a new observation  $y_t$  measured at time  $t$  can readily be incorporated together with previous observations  $y_{<t}$  to provide an updated estimate of a quantity of interest to be used in e.g. a prediction problems

The approaches to DSMs in the engineering, statistical and biological communities differ in many respects, but there is a gain to be made by combining the views.

## 2 Example: New Hampshire temperatures

The data are recordings of the mean annual temperature in degrees Fahrenheit in New Haven, Connecticut, from 1912 to 1971. Data are available in R. A plot of data is shown in Figure 1.

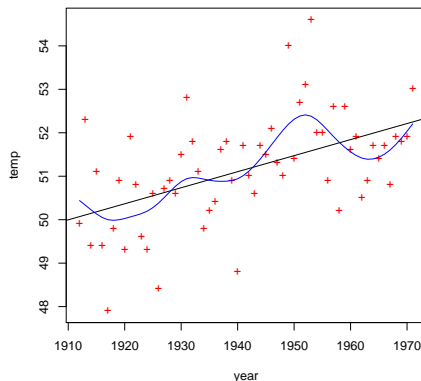


Figure 1: New Hampshire temperatures data with linear trend and smoothed spline added.

For simplicity we recode the year as  $t = 1, 2, \dots, 60$ .

## 3 Regression and recursive least squares

Consider a regression setting model,

$$y_t \approx \alpha + \beta t \tag{1}$$

Suppose the task is to make predictions, and we focus on making 1-step forecasts. That is given information  $D_t$  we wish to predict the temperature in year  $t + 1$ .

### 3.1 Recursive least squares

A naive approach is to refit the model to all available observations  $D_t$  every time a new measurement  $y_t$  is made. Let  $\theta = (\alpha, \beta)$  and let  $\hat{\theta}_t$  denote the estimate of  $\theta$  based on data  $D_t$ . Thus  $\hat{\theta}$  is obtained by minimizing the sum-of-squares,

$$SS = \sum_{t'=1}^t (y_{t'} - (\alpha + \beta t'))^2. \quad (2)$$

In a naive implementation of this approach we will have to store all data, which might be a problem in some applications. However it is possible (and in linear models it is quite simple) to avoid this: Instead we can find  $\hat{\theta}_t$  from  $\hat{\theta}_{t-1}$  and  $(y_t, t)$ . This is called *recursive least squares* method.

### 3.2 Recursive least squares with forgetting factor

Note that with this method is that observations are “weighted equally” in (2); that is observations made in the far past are given the same weight as recent observations. This may not be appropriate. A very practical alternative is instead to minimize a weighted sum-of-squares,

$$SS = \sum_{t'=1}^t \kappa^{t-t'} (y_{t'} - (\alpha + \beta t'))^2. \quad (3)$$

for some constant  $\kappa$  where  $0 < \kappa \leq 1$ . This constant sometimes called a *forgetting factor* as it describes “how fast the past is forgotten” (down weighted) in the regression. Small values of  $\kappa$  means that the past is forgotten quickly. It is worth noticing that  $\kappa$  can not be estimated directly as part of a least squares procedure (letting  $\kappa$  go to zero means that  $SS$  goes to zero no matter the choice of  $\theta$ ). However, one can estimate  $\kappa$  along these lines: Suppose a batch of data is available. Let  $\hat{y}_{t+1} = \hat{\alpha}_t + \hat{\beta}_t(t+1)$  be the 1-step prediction of  $y_{t+1}$ . Note that  $\hat{y}_{t+1}$  depends on  $\theta, \kappa$  so we might write  $\hat{y}_{t+1} = \hat{y}_{t+1}(\theta, \kappa)$ . Then the squared prediction error  $(y_{t+1} - \hat{y}_{t+1}(\theta, \kappa))^2$  is a function of  $(\theta, \kappa)$ . We can then estimate  $(\theta, \kappa)$  by minimizing (the non-linear) function:

$$SS = \sum_{t'=2}^t (y_{t+1} - \hat{y}_{t+1}(\theta, \kappa))^2. \quad (4)$$

Based on this we can obtain an estimate  $\hat{\kappa}$  which can be used in the dynamic update in (3). To some people this approach may appear too much *ad hoc* while other find it perfectly acceptable.

See [\[Wikipedia\]](#) for a details and references.

## 4 Exponential smoothing and the Holt–Winters filter

In this section we discuss exponential smoothing and the Holt–Winters filter.

### 4.1 Exponential smoothing

Suppose there is no obvious trend in data (which does not apply to the New Hampshire temperature data), such that data can be described as “level + noise”. A *one sided exponential smoothing* is obtained as

$$\alpha_t = \kappa y_t + (1 - \kappa)\alpha_{t-1} \text{ with } \alpha_1 = y_1 \quad (5)$$

for some constant  $\kappa$  where  $0 < \kappa \leq 1$ . The 1-step ahead prediction is then simply  $\hat{\alpha}_t + 1 = \alpha_t$ . It is illustrative to write (5) as

$$\alpha_t = \alpha_{t-1} + \kappa(y_t - \hat{\alpha}_t) \text{ with } \alpha_1 = y_1 \quad (6)$$

where  $(y_t - \hat{\alpha}_t)$  can be regarded as the 1-step prediction error. Note that (5) has the property that data are incorporated as they arrive. That is, once  $\alpha_{t-1}$  is calculated there is no need to store  $y_t$  (or any of the preceeding values for that matter). (The term exponential comes from the fact that observations from the past are downweight at an exponential rate.) Clearly,  $\kappa$  can not be estimated by minimizing the sum-of-squares,  $\sum_t (y_t - \alpha_t)$  because that sum is zero if  $\kappa$  is zero. The parameter  $\kappa$  can instead be estimated by minimizing the 1-step prediction error, i.e. by minimizing  $\sum_t (y_t - \hat{\alpha}_t)^2$  where  $\hat{\alpha}_t$  is the 1-step prediction based on data up to time  $t - 1$ .

See [Brockwell and Davis \(2002\)](#) and [\[Wikipedia\]](#) for a details and references.

### 4.2 Holt–Winters filter

If instead there is a (linear) trend in data (as seems to be the case for the New Hampshire data), an alternative is the Holt–Winters filter. (This filter applies to other situations as well, but we shall here focus on the linear case).

Consider a situation where we have data  $D_t$  and wish to make prediction at time  $t + h$ . The Holt–Winters filter does so as:

$$\hat{y}_{t+h} = \alpha_t + \beta_t h \quad (7)$$

The parameters  $\alpha$  and  $\beta$  are updated as

$$\alpha_t = \kappa y_t + (1 - \kappa)(\alpha_{t-1} + \beta_{t-1}) \quad (8)$$

$$\beta_t = \delta(\alpha_t - \alpha_{t-1}) + (1 - \delta)(\beta_{t-1}) \quad (9)$$

Hence there are now two smoothing parameters/forgetting factors, namely  $\kappa$  and  $\delta$ . Note that setting  $\delta = 0$  implies that (8) reduces to (5).

The smoothing parameters are usually estimated from a batch of data based on minimizing the 1-step prediction error.

See [Winters \(1960\)](#) and [Brockwell and Davis \(2002\)](#) for details and references.

## 5 State space models

Linear state space models (SSMs) are traditionally specified as consisting of

$$y_t = F_t^\top \theta_t + v_t, \quad v_t \sim N(0, V_t) \quad (10)$$

$$\theta_t = G_t \theta_{t-1} + w_t, \quad w_t \sim N(0, W_t) \quad (11)$$

$$\theta_0 \sim N(m_0, C_0) \quad (12)$$

known as respectively *observation equation*, the *system equation* and the *initial distribution* (of  $\theta_0$ ). The term  $\lambda_t = F_t^\top \theta_t = \mathbb{E}(y_t | \theta_t)$  is the *signal*. Thus, technically the 6-tuple  $(F_t, V_t, G_t, W_t, m_0, C_0)$  defines a linear state space model.

The state space model is a *probabilistic model* for all  $\theta_t$ s and all  $y_t$ s. An alternative (but equivalent) formulation is

$$p(y, \theta) = p(\theta_0) \prod_t p(\theta_t | \theta_{t-1}) p(y_t | \theta_t)$$

where

$$p(\theta_t | \theta_{t-1}) : N(G_t \theta_{t-1}, W_t), \quad (13)$$

$$p(y_t | \theta_t) : N(F_t^\top \theta_t, V_t), \quad (14)$$

### 5.1 Examples of SSMs

Many classical statistical models can be formulated as an SSM. Consider an autoregression of order 1, i.e.

$$y_t = \kappa y_{t-1} + v_t \quad (15)$$

This model can be put in state space form as

$$y_t = \alpha_t \quad (16)$$

$$\alpha_t = \kappa \alpha_{t-1} + v_t \quad (17)$$

## 6 The Kalman filter

Based on observed data we can then calculate (the distribution of) the parameters  $\theta_t$ . The Kalman filter (named after its inventor, Rudolf Kalman) is, roughly speaking, an efficient algorithm for recursively estimating  $\theta_t$  when new observations become available. The filter was developed in papers by [Kalman \(1960\)](#) and [Kalman and S. \(1961\)](#).

A wide variety of Kalman filters have now been developed, from Kalman's original formulation, now called the simple Kalman filter, to the extended Kalman filter, the information filter, a variety of square-root filters, unscented filter, particle filters.

See [\[Wikipedia\]](#) for an overview and references.

The Kalman filter works by “updating  $\theta_t$  as new observations arrive”, i.e. for  $t = 1, \dots, T$ : Given is that  $\theta_{t-1}|D_{t-1} \sim N(m_{t-1}, C_{t-1})$ . Then

$$\theta_t|D_{t-1} \sim N(\overbrace{G_t, m_{t-p}}^{a_t}, \overbrace{G_t C_{t-1} G_t^\top + W_t}^{R_t}) \quad (18)$$

$$y_t|D_{t-1} \sim N(\overbrace{F_t^\top a_t}^{f_t}, \overbrace{F_t^\top R_t F_t + V_t}^{Q_t}) \quad (19)$$

$$\theta_t|D_t \sim N(\underbrace{a_t + R_t F_t Q_t^{-1} (y_t - f_t)}_{A_t}, \underbrace{R_t - A_t Q_t A_t^\top}_{C_t}) \quad (20)$$

Hence, the only thing needed to be stored from time  $t-1$  to time  $t$  is  $(m_{t-1}, C_{t-1})$ . Here  $f_t$  is the 1-step forecast,  $e_t$  is the *1-step forecast error* while  $A_t$  is the *Kalman gain*.

For the normal distribution an alternative formulation is as follows: Since  $y_t = F_t^\top \theta_t + v_t$  we have  $\text{Cov}(y_t, \theta_t) = F_t \text{Var}(\theta_t)$  and so  $\text{Cov}(y_t, \theta_t|D_{t-1}) = F_t^\top R_t$ . Hence

$$\begin{pmatrix} y_t \\ \theta_t \end{pmatrix} | D_{t-1} \sim N \left( \begin{bmatrix} f_t \\ a_t \end{bmatrix} \begin{bmatrix} Q_t & F_t^\top R_t \\ R_t F_t & R_t \end{bmatrix} \right) \quad (21)$$

### 6.1 The likelihood

The state space model depends on unknown parameters through  $V_t$  and  $W_t$ . There are several ways of estimating these; a brute force approach is the following: We have

$$p(y) = \prod_t p(y_t | y_{t-1}, \dots, y_1) = \prod_t p(y_t | D_{t-1})$$

We have  $y_t|D_{t-1} \sim N(f_t, Q_t)$  so the log likelihood  $\log L_t$  for a single observation is

$$\log L_t = -\frac{d}{2} \log(2\pi) - \frac{1}{2} (\det(Q_t) + e_t^\top Q_t^{-1} e_t)$$

because  $L_t = (2\pi)^{-d/2} \det(Q_t)^{-1/2} \exp(-\frac{1}{2} e_t^\top Q_t^{-1} e_t)$ . Hence the entire log likelihood is

$$\log L = -n \frac{d}{2} \log(2\pi) - \frac{1}{2} \sum_t (\det(Q_t) + e_t^\top Q_t^{-1} e_t)$$

This likelihood can be maximized numerically, but it is often not entirely trivial to get this to work in specific applications.

## 6.2 Online outlier detection

Suppose  $y_t$  is  $p$  dimensional. Based on  $D_{t-1}$  the distribution of  $y_t$  is  $y_t|D_{t-1} \sim N(f_t, Q_t)$  so the prediction error is

$$e_t = y_t - f_t \sim N(0, Q_t)$$

That means that when  $y_t$  is observed but before we update the equations, we can see how “likely”  $y_t$  is by noticing

$$ssd_t = e_t'(Q_t)^{-1} e_t \sim \chi_p^2$$

This provides a tool for on-line detection of outliers: A large value of  $ssd_t$  suggests that  $y_t$  is an “unlikely observation”, that is, it might be erroneous.

## 7 Connecting exponential smoothing and state space models

There is a close connection between exponential smoothing and SSMs. Consider exponential smoothing as set out in Section 4 with

$$\alpha_t = \alpha_{t-1} + \kappa(y_t - \hat{\alpha}_t) \text{ with } \alpha_1 = y_1 \quad (22)$$

We can think of  $e_t = (y_t - \hat{\alpha}_t)$  as a prediction error (which initially is zero). It is then trivially true that  $y_t = \alpha_{t-1} + (y_t - \alpha_{t-1}) = \alpha_{t-1} + e_t$ . Combining this with (22) gives

$$y_t = \alpha_{t-1} + e_t \quad (23)$$

$$\alpha_t = \alpha_{t-1} + \kappa e_t \quad (24)$$

which has a resemblance with the general form of a linear state space model in (12). There are however two important differences: The error term  $e_t$  appears both in the observation and system equation and there has been made no assumption about the distribution of the error term. If we assume the error terms to be normal and independent, then we are

almost at (12). The fact that the error term appears in both the observation and system equation only changes the filter equations slightly; the Kalman filter can still be used.

This can be put in a more general setting: Let

$$y_t = F_t^\top \theta_{t-1} + e_t, \quad e_t \sim N(0, V_t) \quad (25)$$

$$\theta_t = G_t \theta_{t-1} + \kappa e_t \quad (26)$$

where  $\kappa$  now is a vector in which each element is in  $[0; 1]$ . From this we find that

$$y_t = F_t^\top \theta_{t-1} + e_t, \quad e_t \sim N(0, V_t) \quad (27)$$

$$\theta_t = G_t \theta_{t-1} + \kappa(y_t - F_t^\top \theta_{t-1}) = D \theta_{t-1} + \kappa y_t, \quad (28)$$

say. So  $\theta_t$  is hence a weighted sum of  $\theta_{t-1}$  and  $t_t$ .

## 8 Discussion

Purists would often feel uncomfortable with the methods set out in Sections 3 and 4. They are not per se based on an assumption of an underlying statistical model. We can obtain parameter estimates, but the way they are obtained appears quite ad hoc and hence we do not know “how good they are”. We can make predictions, but we do not know how good they are either. On the other hand, the methods are quite easy to work with in practice and are also easily implemented.

The state space models on the other hand appear attractive to some people because they are based on a proper statistical model, the parameter estimates have well understood interpretations (for example, (20) gives the conditional distribution of  $\theta_t$  given  $D_t$ ), the distribution of the prediction errors is known etc. In practice however, it is usually tricky to estimate the unknown parameters and therefore one often ends up by stipulating reasonable values.

With a view towards biology there are some additional reservations to be made. In physical/engineering applications there are often (but certainly not always) differential equations etc. which describe a natural form of the system equation. For example that  $d\theta/dt = h(\theta, t)$  which can be translated into the approximation that  $\theta_t \approx \theta_{t-1} + h(\theta_{t-1}, t-1)$  which yields the system equation. In biological sciences such quantitative descriptions are more rare. Therefore, it is often necessary to stipulate a system equation which essentially provides a smoothing of the state vector. It is in some cases therefore not clear whether it is worthwhile to invoke the entire “Kalman machinery” for that purpose (other simpler methods exist for that).



## 9 Acknowledgements

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