Bayesian Inference for Switching Linear Dynamical Systems

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1 Model description

A switching linear dynamical system – also known as *switching state space model* – is defined, borrowing the notation from Linderman et al. [3], by the set of discrete-time stochastic equations

$$x_t = A_{z_t} x_{t-1} + b_{z_t} + v_t, (1$$

$$y_t = C_{z_t} x_t + d_{z_t} + w_t, (2)$$

where $v_t \in \mathbb{R}^M$ and $w_t \in \mathbb{R}^N$ are Gaussian-distributed random vectors with mean zero and variance Q_{z_t} , S_{z_t} respectively. The vectors $y_t \in \mathbb{R}^N$, $t = 1, \ldots, T$ may represent a time series of observations, while the $x_t \in \mathbb{R}^M$ are a set of continuous latent states linked together by linear dynamics defined by the matrices $A_k \in \mathbb{R}^{M \times M}$ and bias vectors $b_k \in \mathbb{R}^M$. The transformation between x and y is also linear, through the matrices $C_k \in \mathbb{R}^{N \times M}$ and bias vectors $d_k \in \mathbb{R}^N$.

The linear parameters A_k , b_k , C_k , d_k form a discrete set of K elements, and a discrete latent variable $z_t \in \{1, \ldots, K\}$ sets the specific instances in use at time step t. The variable z evolves over time as a Markov process, meaning that z_t is conditionally independent of all previous states except for its immediate predecessor z_{t-1} :

$$p(z_t \mid z_{t-1}, z_{t-2}, \dots, z_1) = p(z_t \mid z_{t-1}).$$
 (3)

We will denote the probability to transition from $z_{t-1} = j$ to $z_t = k$ with π_{jk} . The transition $z_{t-1} \to z_t$ effectively modifies the linear dynamics from x_{t-1} to x_t and the linear transformation from x_t to y_t , switching from one regime to another

Given a set of data points y_t , the goal is then to infer the posterior distribution of the parameter set

$$\vartheta = \{\pi_k, A_k, b_k, Q_k, C_k, d_k, S_k\},\tag{4}$$

where $x_{1:T}$ denotes the whole sequence x_1, x_2, \ldots, x_T , and π_k the kth row of the transition matrix.

2 Implementation in Stan

To set up a Monte Carlo sampling scheme, we chose to work with the Stan [5] programming language and its implementation in R through the package rstan [4]. Sampling in Stan is done by default using a variant of the Hamiltonian Monte Carlo scheme called 'No-U-Turn sampler' or NUTS [1].

The model cannot be implemented directly as it is, in the sense of specifying a categorical likelihood for the transition $z_t \to z_{t+1}$ and Gaussian likelihoods for $x_t \mid x_{t-1}$ and $y_t \mid x_t$, because Stan does not allow the definition of integer parameters: so, one should marginalize over the hidden discrete states. Besides Stan's limitations in this regard, the resulting strategy – known in the literature as forward algorithm – is more efficient than the straightforward implementation in sampling low-probability states, and is commonly used in similar inference problems involving hidden Markov models or other state space models [2].

2.1 The forward algorithm

The basic idea behind the forward algorithm is to exploit a recursive relationship to build the full likelihood: indeed, consider the quantity

$$\gamma_t(k) := p(z_t = k, x_{1:t}, y_{1:t}).$$
 (5)

By summing over the z states at t-1 first and then using the chain rule repeatedly, we can write

$$\gamma_{t}(k) = \sum_{j=1}^{K} p(z_{t} = k, z_{t-1} = j, x_{1:t}, y_{1:t})$$

$$= p(y_{t} \mid z_{t} = k, x_{t}) p(x_{t} \mid z_{t} = k, x_{t-1})$$

$$\cdot \sum_{j=1}^{K} \pi_{jk} p(z_{t-1} = j, x_{1:t-1}, y_{1:t-1}),$$
(6)

where we have recognized the conditional probability $p(z_t = k \mid z_{t-1} = j, x_{1:t-1}, y_{1:t-1}) = p(z_t = k \mid z_{t-1} = j)$ as the element (j, k) of the transition matrix π . The first two terms outside the sum are the likelihoods of y_t and x_t , and because of the model definition they only depend on z_t , x_t and x_{t-1} . Also, they are simply Gaussian densities:

$$\mathcal{L}_k(y_t) := p(y_t \mid z_t = k, x_t) = \mathcal{N}(C_k x_t + d_k, S_k), \tag{7}$$

$$\mathcal{L}_k(x_t) := p(x_t \mid z_t = k, x_{t-1}) = \mathcal{N}(A_k x_{t-1} + b_k, Q_k).$$
 (8)

Then, the remaining terms in the sum in (6) are nothing else than $\gamma_{t-1}(j)$, giving us the recursive relation we needed:

$$\gamma_t(k) = \mathcal{L}_k(y_t) \,\mathcal{L}_k(x_t) \sum_{j=1}^K \pi_{jk} \gamma_{t-1}(j). \tag{9}$$

Indeed, to retrieve the full joint likelihood of the sequences $x_{1:T}$ and $y_{1:T}$ we only need to marginalize the γ at

$$p(x_{1:T}, y_{1:T}) = \sum_{k=1}^{K} p(z_T = k, x_{1:T}, y_{1:T}) = \sum_{k=1}^{K} \gamma_T(k).$$
 (10)

To recursively build γ_t up to time T we need $O(TK^2)$ operations, because of the double marginalization over z_t and z_{t-1} . To initialize the recursion,

$$\gamma_1(k) = p(z_1 = k, x_1, y_1)
= \mathcal{L}_1(y_1) p(x_1 | z_1 = k) p(z_1 = k).$$
(11)

The last two terms are the prior distributions on x_1 and z_1 . We chose a multivariate Gaussian for the first and a uniform distribution over the K states for the second.

At this point, we also need the prior distributions for the dynamical parameters. Following the suggestion from Linderman et al. [3], we chose matrix-normal-inverse-Wishart priors:

$$(A_k, b_k), Q_k \sim \text{MNIW}(M_x, \Omega_x, \Psi_x, \nu_x)$$
 (12)

$$(C_k, d_k), S_k \sim \text{MNIW}(M_y, \Omega_y, \Psi_y, \nu_y).$$
 (13)

Here $M_x \in \mathbb{R}^{M \times (M+1)}$ and $M_y \in \mathbb{R}^{N \times (M+1)}$ are the mean matrices of the matrix normals, $\Omega_x, \Omega_y \in \mathbb{R}^{(M+1) \times (M+1)}$ their between-column covariance matrices, while $\Psi_x \in \mathbb{R}^{M \times M}$ and $\Psi_y \in \mathbb{R}^{N \times N}$ are the scale matrices of the inverse Wisharts and ν_x, ν_y their degrees of freedom. The returned random matrices with M+1 columns are then split between the matrices A_k and C_k and their corresponding bias vectors b_k and d_k .

2.2 Reconstructing the hidden states

Since we marginalize out the z sequence during the sampling procedure, we need a way to recover them probabilistically. One way to do it is to search a posteriori for the most likely hidden sequence of z states conditioned to the observed $y_{1:T}$ and inferred $x_{1:T}$. This is done through the so-called 'Viterbi algorithm' [5], which is based on a recursive relation much like the forward algorithm.

Indeed, consider the quantity

$$\eta_t(k) \coloneqq \arg\max_{z_{1:t-1}} p(z_{1:t-1}, z_t = k, x_{1:t}, y_{1:t}).$$
(14)

If we proceed similarly to (6), and using also the fact that

$$\max_{a,b} [f(a)g(a,b)] = \max_{a} [f(a)\max_{b} g(a,b)],$$
 (15)

we can expand the definition as

$$\eta_t(k) = \underset{j \in \{1, \dots, K\}}{\operatorname{arg max}} [p(y_t \mid z_t = k, x_t) p(x_t \mid z_t = k, x_{t-1})]$$

$$\pi_{jk} \arg \max_{z_{1:t-2}} p(z_{1:t-2}, z_{t-1} = j, x_{1:t-1}, y_{1:t-1})]. \quad (16)$$

the last time step T over the discrete states $k = 1, 2, \dots, K$: We then recognize the last factor as η at the previous time step t-1, giving us the recursive relation

$$\eta_t(k) = \mathcal{L}_k(y_t) \mathcal{L}_k(x_t) \underset{j \in \{1, \dots, K\}}{\arg \max} [\pi_{jk} \, \eta_{t-1}(j)].$$
 (17)

Regarding the initial value, there is no z_{t-1} to maximize over, so

$$\eta_1(k) = p(z_1 = k, x_1, y_1),$$
(18)

which is the same initialization of $\gamma_1(k)$ as shown in the previous section, Eq. (11). Once we have the value of η at time T we can maximize over z_T and recover the maximumprobability sequence $\hat{z}_{1:T}$:

$$\hat{z}_{1:T} = \arg\max_{z_{1:T}} p(z_{1:T}, x_{1:T}, y_{1:T}) = \arg\max_{k \in \{1, \dots, K\}} \eta_T(k).$$
 (19)

The procedure is thus substantially the same as the forward algorithm, but with maximization replacing summation.

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