Efficient smoothness selection for nonparametric Markov-switching models via quasi restricted maximum likelihood

Jan-Ole Koslik

St Andrews, February 5, 2025



- PhD student in the Statistics and Data Analysis Group at Bielefeld University
- mostly working on hidden Markov models and their relatives





minutes

HMM — model formulation



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- hidden state process selects which distribution is active at time t
- state process is a Markov chain



minutes

Reminder: Markov chains

Markovian state process is fully characterised by the initial distribution

$$\boldsymbol{\delta}^{(1)} = ig(\mathsf{Pr}(\mathcal{S}_1 = 1), \dots, \mathsf{Pr}(\mathcal{S}_1 = N)ig)$$

and the transition probabilities

$$\gamma_{ij}^{(t)} = \Pr(S_{t+1} = j \mid S_t = i),$$

which we summarise in the transition probability matrix (t.p.m.)

$$\boldsymbol{\Gamma}^{(t)} = (\gamma_{ij}^{(t)})_{i,j=1,\dots,N}.$$

Estimating HMMs

We can efficiently calculate the HMM likelihood using the forward algorithm

$$\mathcal{L}(\boldsymbol{ heta}) = \boldsymbol{\delta}^{(1)} \boldsymbol{P}(x_1) \boldsymbol{\Gamma}^{(1)} \boldsymbol{P}(x_2) \boldsymbol{\Gamma}^{(2)} \cdot \ldots \cdot \boldsymbol{\Gamma}^{(\mathcal{T}-1)} \boldsymbol{P}(x_{\mathcal{T}}) \mathbf{1},$$

where $\boldsymbol{P}(x_t) = \operatorname{diag}(f_1(x_t), \ldots, f_N(x_t)).$

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These approximate the gradient via finite differencing.



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 → difficult as we can't do state-specific EDA
- potential covariate effects typically modelled using **linear** predictors \rightarrow may result in us missing interesting relationships



= complicated

 \rightarrow often substantial lack of fit

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Selecting the Number of States in Hidden Markov Models: Pragmatic Solutions Illustrated Using Animal Movement

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Obvious alternative: nonparametric approach using penalised splines

An informal introduction to penalised splines

Idea: Perform **basis expansion** in x and represent smooth function s(x) as a linear combination of fixed basis functions $B_k(x)$

$$s(x) = b_1B_1(x) + b_2B_2(x) + \ldots + b_kB_k(x) = \boldsymbol{b}^{\mathsf{T}}\boldsymbol{B}(x)$$



For example, when using **B-Splines** $B_k(x)$ is a **piecewise** polynomial



and zero outside the outer knots.

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$$\lambda \int s''(x)^2 \ dx = \lambda b^{\mathsf{T}} S b,$$

where **S** is fixed penalty matrix with entries $S_{ij} = \int B_i''(x)B_j''(x) dx$.

Nonparametric emission distributions



Langrock, Kneib, Sohn, and DeRuiter, 2015

Markov-switching GAM



Langrock, Kneib, Glennie, and Michelot, 2017

Smooth covariate effects on the state process



Feldmann et al., 2023

So this is where my spline adventure begins...

- fairly applied project: $\gamma_{ii}^{(t)} \sim$ s(time of day, day)
- implementation with **fixed** penalisation straightforward



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But how to find a good **penalty strength**??

JOURNAL ARTICLE

Nonparametric Inference in Hidden Markov Models Using P-Splines @

Roland Langrock 🖾, <u>Thomas Kneib</u>, Alexander Sohn, Stacy L. DeRuiter

Biometrics, Volume 71, Issue 2, June 2015, Pages 520–528, https://doi.org/10.1111/biom.12282 Published: 13 January 2015 Article history ▼

Markov-switching generalized additive models

Roland Langrock¹ · Thomas Kneib² · Richard Glennie³ · Théo Michelot⁴



SPECIAL ISSUE ARTICLE 🔂 Full Access

Spline-based nonparametric inference in general stateswitching models

Roland Langrock 🔀, Timo Adam, Vianey Leos-Barajas, Sina Mews, David L. Miller, Yannis P. Papastamatiou

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Ideally: define the penalised log-likelihood \rightarrow automatic smoothness-selection

Comput Stat (2012) 27:757–777 DOI 10.1007/s00180-011-0289-6

ORIGINAL PAPER

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- splines as random effects?
- Laplace approximation?



Why are splines random effects?

Simple setting involving one penalised spline and no unpenalised parameters:

$$\ell_{\rho}(\boldsymbol{b};\lambda) = \ell(\boldsymbol{b}) - \frac{1}{2}\lambda \boldsymbol{b}^{\mathsf{T}} \boldsymbol{S} \boldsymbol{b},$$

• coefficients
$$\boldsymbol{b} = (b_1, \dots, b_k)$$

- fixed penalty matrix \boldsymbol{S}
- penalty strength λ

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Penalised likelihood function

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We might as well assume $m{b}\sim\mathcal{N}(m{0},m{S}^{-1}/\lambda)$ and add the missing normalisation constant^1

$$\mathcal{L}_{j}(\boldsymbol{b};\lambda) = \mathcal{L}(\boldsymbol{b}) \cdot (2\pi)^{-k/2} \det(\lambda \boldsymbol{S})^{1/2} \exp\left(-\frac{1}{2}\lambda \boldsymbol{b}^{\mathsf{T}} \boldsymbol{S} \boldsymbol{b}\right)$$

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giving the joint log-likelihood

$$\ell_j(\boldsymbol{b}; \lambda) = \ell(\boldsymbol{b}) - \frac{k}{2}\log(2\pi) + \frac{1}{2}\log\det(\lambda \boldsymbol{S}) - \frac{1}{2}\lambda \boldsymbol{b}^{\mathsf{T}} \boldsymbol{S} \boldsymbol{b}$$

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How to estimate models with random effects?

Joint likelihood of the data and the random effect as a function of λ

 $f_{\lambda}(\mathbf{x}, \mathbf{b}) = f(\mathbf{x} \mid \mathbf{b}) \cdot f_{\lambda}(\mathbf{b}),$

with $f(\boldsymbol{x} \mid \boldsymbol{b}) = \mathcal{L}(\boldsymbol{b})$ and $f_{\lambda}(\boldsymbol{b}) \sim \mathcal{N}(\boldsymbol{0}, \boldsymbol{S}^{-1}/\lambda)$.

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which we would like to maximise to find the **MLE** $\hat{\lambda}$.

Marginal ML



Marginal ML



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In reality, this integral is intractable!

What can we do?

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The Laplace approximation

• find the mode (in **b**) of the joint log-likelihood

$$\ell_j(\boldsymbol{b},\lambda) = -\frac{k}{2}\log(2\pi) + \frac{1}{2}\log\det(\lambda S) + \ell(\boldsymbol{b}) - \frac{1}{2}\lambda \boldsymbol{b}^{\mathsf{T}}\boldsymbol{S}\boldsymbol{b}$$

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• second-order Taylor approximation around the mode:

$$\ell_{approx}(\boldsymbol{b},\lambda) = \ell_j(\hat{\boldsymbol{b}},\lambda) - \frac{1}{2}(\boldsymbol{b}-\hat{\boldsymbol{b}})^{\mathsf{T}}\boldsymbol{J}(\lambda)(\boldsymbol{b}-\hat{\boldsymbol{b}}),$$

where $\boldsymbol{J}(\lambda) = -\nabla^2 \ell_j(\hat{\boldsymbol{b}}, \lambda)$ is the (negative) Hessian of $\ell_j(\boldsymbol{b}, \lambda)$ w.r.t. \boldsymbol{b} at $\hat{\boldsymbol{b}}(\lambda)$.



• now exponentiate to obtain likelihood and integrate out $m{b}$

$$\exp(\ell_j(\hat{\boldsymbol{b}},\lambda)) \cdot \int \exp(-\frac{1}{2}(\boldsymbol{b}-\hat{\boldsymbol{b}})^{\mathsf{T}}\boldsymbol{J}(\lambda)(\boldsymbol{b}-\hat{\boldsymbol{b}})) d\boldsymbol{b}$$

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• hence, approximate marginal log-likelihood becomes

$$\ell(\lambda) pprox \ell_j(\hat{m{b}}, \lambda) + rac{k}{2}\log(2\pi) - rac{1}{2}\log\det(m{J}(\lambda))$$



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- but each evaluation of marginal log-likelihood requires inner optimisation w.r.t. $b \rightarrow$ leads to nested optimisation in general

Schellhase and Kauermann (2012) start with the (approximate) marginal log-likelihood

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Writing out our joint log-likelihood, we have as our marginal log-likelihood

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which we can now (partially) differentiate w.r.t. λ

$$\frac{\partial}{\partial \lambda} \Big(\frac{1}{2} \log \det(\lambda S) \Big) = \frac{\partial}{\partial \lambda} \Big(\frac{1}{2} \log \big(\lambda^k \det(S) \big) \Big) = \frac{\partial}{\partial \lambda} \Big(\frac{k}{2} \log(\lambda) + \frac{1}{2} \log \det(S) \Big) = \frac{k}{2\lambda}$$

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$$\frac{\partial}{\partial \lambda} \Big(-\frac{1}{2} \log \det \big(\boldsymbol{J}(\lambda) \big) \Big) = -\frac{1}{2} \mathrm{tr} \big(\boldsymbol{J}(\lambda)^{-1} \boldsymbol{S} \big)$$

Hence in total

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from which we can construct the estimating equation (omitting the details)

$$\lambda = \frac{\operatorname{tr}(\boldsymbol{J}(\lambda)^{-1}\boldsymbol{J}(\lambda=0))}{\hat{\boldsymbol{b}}^{\mathsf{T}}\boldsymbol{S}\hat{\boldsymbol{b}}} = \frac{\operatorname{dof}(\lambda)}{\hat{\boldsymbol{b}}^{\mathsf{T}}\boldsymbol{S}\hat{\boldsymbol{b}}}.$$

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which yields the iterative procedure:

- 1. fit model via penalised ML
- 2. calculate Hessian at optimum
- 3. update penalty strength
- 4. repeat 1.-3. until convergence



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- So why did it work??

The full setting

$$\ell_{\mathcal{P}}(\boldsymbol{a}, \boldsymbol{b}; \boldsymbol{\lambda}) = \ell(\boldsymbol{a}, \boldsymbol{b}) - rac{1}{2} \sum_{i} \lambda_{i} \boldsymbol{b}_{i}^{\mathsf{T}} \boldsymbol{S} \boldsymbol{b}_{i}$$

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Problem: If we integrate out **b**, marginal likelihood is more complicated beause of **a**.

Solution: Assume $a \sim \mathcal{N}(\mathbf{0}, \infty)$ and integrate out both a and $b \rightarrow$ restricted maximum likelihood (REML), Laird and Ware, 1982

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 \rightarrow smoothness selection procedure that makes nonparametric HMMs feasible!



Here comes RTMB (Kristensen, 2024) with **automatic differentiation**, natively supporting the full Laplace approximation for models written in plain R code



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- user only needs to specify penalised negative log-likelihood
- **qREML** + **AD** \rightarrow efficiency skyrocketed!

Practical usage

library(LaMa)

pnll <- function(par){ # penalised negative log-likelihood
 ... # computing the negative log-likelihood
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mod <- qreml(pnll, par, dat, random = "splinePars")</pre>

Real-data example

- Western Australia, extreme seasonal changes
- seven bull sharks tagged (14K observations)
- temperature, depth, and acceleration data
- response: overall dynamic body acceleration
- 2-state HMM: low and high activity



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- $\gamma_{ii}^{(t)} \sim s(tod_t) + AvgTemp_t * s(tod_t)$ (parametric in original paper)
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- model fit takes \sim 5 minutes (32 penalised fits until convergence)



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time of day

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- models with i.i.d. random effects can be fitted using the same approach
- RTMB will become an extremely valuable tool for fitting complex models

