

Dynamic models

- 6 Dynamic models
 - Dependent data
 - The $AR(p)$ model
 - The $MA(q)$ model
 - Hidden Markov models

Dependent data

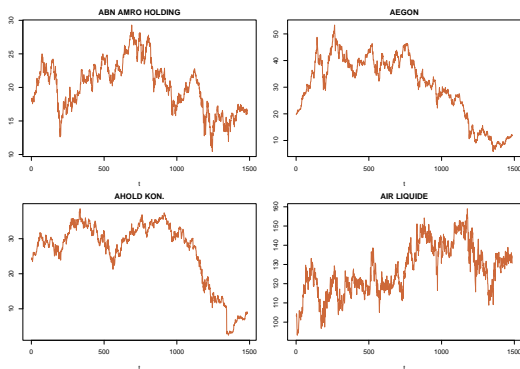
Huge portion of real-life data involving dependent datapoints

Example (Capture-recapture)

- capture histories
- capture sizes

Eurostoxx 50

First four stock indices of the financial index Eurostoxx 50



Markov chain

Stochastic process $(x_t)_{t \in \mathcal{T}}$ where distribution of x_t given the past values $\mathbf{x}_{0:(t-1)}$ only depends on x_{t-1} .

Homogeneity: distribution of x_t given the past constant in $t \in \mathcal{T}$.

Corresponding likelihood

$$\ell(\theta | \mathbf{x}_{0:T}) = f_0(x_0 | \theta) \prod_{t=1}^T f(x_t | x_{t-1}, \theta)$$

[Homogeneity means f independent of t]

Stationarity constraints

Difference with the independent case: *stationarity* and *causality* constraints put restrictions on the parameter space

Stationarity processes

Definition (Stationary stochastic process)

$(x_t)_{t \in \mathcal{T}}$ is stationary if the joint distributions of (x_1, \dots, x_k) and $(x_{1+h}, \dots, x_{k+h})$ are the same for all h, k 's.

It is *second-order stationary* if, given the autocovariance function

$$\gamma_x(r, s) = \mathbb{E}[\{x_r - \mathbb{E}(x_r)\}\{x_s - \mathbb{E}(x_s)\}], \quad r, s \in \mathcal{T},$$

then

$$\mathbb{E}(x_t) = \mu \quad \text{and} \quad \gamma_x(r, s) = \gamma_x(r + t, s + t) \equiv \gamma_x(r - s)$$

for all $r, s, t \in \mathcal{T}$.

Imposing or not imposing stationarity

Bayesian inference on a non-stationary process can be [formally] conducted

Debate

From a Bayesian point of view, to impose the *stationarity* condition is objectionable: stationarity requirement on finite datasets artificial *and/or* datasets themselves should indicate whether the model is stationary

Reasons for imposing stationarity: asymptotics (Bayes estimators are not necessarily convergent in non-stationary settings) causality, identifiability and ... common practice.

Unknown stationarity constraints

Practical difficulty: for complex models, stationarity constraints get quite involved to the point of being unknown in some cases

The AR(1) model

Case of linear Markovian dependence on the last value

$$x_t = \mu + \rho(x_{t-1} - \mu) + \epsilon_t, \epsilon_t \stackrel{\text{i.i.d.}}{\sim} \mathcal{N}(0, \sigma^2)$$

If $|\rho| < 1$, $(x_t)_{t \in \mathbb{Z}}$ can be written as

$$x_t = \mu + \sum_{j=0}^{\infty} \rho^j \epsilon_{t-j}$$

and this is a stationary representation.

Stationary but...

If $|\rho| > 1$, alternative stationary representation

$$x_t = \mu - \sum_{j=1}^{\infty} \rho^{-j} \epsilon_{t+j}.$$

This stationary solution is criticized as artificial because x_t is correlated with *future* white noises $(\epsilon_t)_{s>t}$, unlike the case when $|\rho| < 1$.

Non-causal representation...

Standard constraint

© Customary to restrict AR(1) processes to the case $|\varrho| < 1$

Thus use of a uniform prior on $[-1, 1]$ for ϱ

Exclusion of the case $|\varrho| = 1$ that leads to a random walk because the process is then a random walk [*no stationary solution*]

The AR(p) model

Conditional model

$$x_t | x_{t-1}, \dots \sim \mathcal{N} \left(\mu + \sum_{i=1}^p \varrho_i (x_{t-i} - \mu), \sigma^2 \right)$$

- Generalisation of AR(1)
- Among the most commonly used models in dynamic settings
- More challenging than the static models (stationarity constraints)
- Different models depending on the processing of the starting value x_0

Stationarity + causality

Stationarity constraints in the prior as a restriction on the values of θ .

Theorem

AR(p) model second-order stationary and causal iff the roots of the polynomial

$$\mathcal{P}(x) = 1 - \sum_{i=1}^p \rho_i x^i$$

are all outside the unit circle

Initial conditions

Unobserved initial values can be processed in various ways

- ① All \mathbf{x}_{-i} 's ($i > 0$) set equal to μ , for computational convenience
- ② Under stationarity and causality constraints, $(x_t)_{t \in \mathbb{Z}}$ has a stationary distribution: Assume $\mathbf{x}_{-p:-1}$ distributed from stationary $\mathcal{N}_p(\mu \mathbf{1}_p, \mathbf{A})$ distribution
Corresponding marginal likelihood

$$\int \sigma^{-T} \prod_{t=0}^T \exp \left\{ \frac{-1}{2\sigma^2} \left(x_t - \mu - \sum_{i=1}^p \varrho_i (x_{t-i} - \mu) \right)^2 \right\} f(\mathbf{x}_{-p:-1} | \mu, \mathbf{A}) d\mathbf{x}_{-p:-1},$$

Initial conditions (cont'd)

- ③ Condition instead on the initial *observed* values $\mathbf{x}_{0:(p-1)}$

$$\ell^c(\mu, \varrho_1, \dots, \varrho_p, \sigma | \mathbf{x}_{p:T}, \mathbf{x}_{0:(p-1)}) \propto \sigma^{-T} \prod_{t=p}^T \exp \left\{ - \left(x_t - \mu - \sum_{i=1}^p \varrho_i (x_{t-i} - \mu) \right)^2 / 2\sigma^2 \right\} .$$

Prior selection

For AR(1) model, Jeffreys' prior associated with the stationary representation is

$$\pi_1^J(\mu, \sigma^2, \rho) \propto \frac{1}{\sigma^2} \frac{1}{\sqrt{1 - \rho^2}}.$$

Extension to higher orders quite complicated (ρ part)!

Natural conjugate prior for $\theta = (\mu, \rho_1, \dots, \rho_p, \sigma^2)$:

normal distribution on $(\mu, \rho_1, \dots, \rho_p)$ and inverse gamma distribution on σ^2

... and for constrained ρ 's?

Stationarity constraints

Under stationarity constraints, complex parameter space: each value of ϱ needs to be checked for roots of corresponding polynomial with modulus less than 1

E.g., for an AR(2) process with autoregressive polynomial $\mathcal{P}(u) = 1 - \varrho_1 u - \varrho_2 u^2$, constraint is

$$\varrho_1 + \varrho_2 < 1, \quad \varrho_1 - \varrho_2 < 1 \quad \text{and} \quad |\varrho_2| < 1.$$

A first useful reparameterisation

Durbin–Levinson recursion proposes a *reparameterisation* from the parameters ϱ_i to the *partial autocorrelations*

$$\psi_i \in [-1, 1]$$

which allow for a uniform prior on the hypercube.

Partial autocorrelation defined as

$$\psi_i = \text{corr} \left(x_t - \mathbb{E}[x_t | x_{t+1}, \dots, x_{t+i-1}], \right. \\ \left. x_{t+i} - \mathbb{E}[x_{t+i} | x_{t+1}, \dots, x_{t+i-1}] \right)$$

[see also Yule-Walker equations]

Durbin–Levinson recursion

Transform

- 1 Define $\varphi^{ii} = \psi_i$ and $\varphi^{ij} = \varphi^{(i-1)j} - \psi_i \varphi^{(i-1)(i-j)}$, for $i > 1$ and $j = 1, \dots, i-1$.
- 2 Take $\varrho_i = \varphi^{pi}$ for $i = 1, \dots, p$.

Stationarity & priors

For AR(1) model, Jeffreys' prior associated with the stationary representation is

$$\pi_1^J(\mu, \sigma^2, \rho) \propto \frac{1}{\sigma^2} \frac{1}{\sqrt{1 - \rho^2}}.$$

Within the non-stationary region $|\rho| > 1$, Jeffreys' prior is

$$\pi_2^J(\mu, \sigma^2, \rho) \propto \frac{1}{\sigma^2} \frac{1}{\sqrt{|1 - \rho^2|}} \sqrt{\left| 1 - \frac{1 - \rho^{2T}}{T(1 - \rho^2)} \right|}.$$

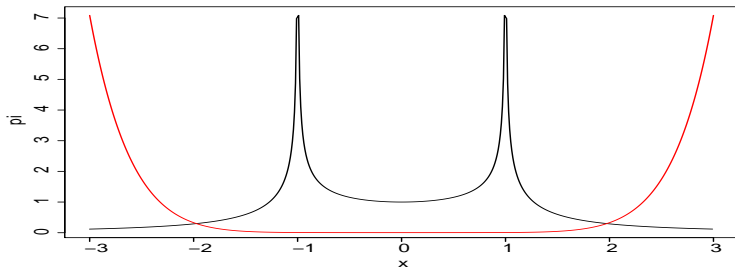
The dominant part of the prior is the non-stationary region!

Alternative prior

The reference prior π_1^J is only defined when the stationary constraint holds.

Idea Symmetrise to the region $|\varrho| > 1$

$$\pi^B(\mu, \sigma^2, \varrho) \propto \frac{1}{\sigma^2} \begin{cases} 1/\sqrt{1-\varrho^2} & \text{if } |\varrho| < 1, \\ 1/|\varrho|\sqrt{\varrho^2-1} & \text{if } |\varrho| > 1, \end{cases}$$



MCMC consequences

When devising an MCMC algorithm, use the Durbin-Levinson recursion to end up with single normal simulations of the ψ_i 's since the ϱ_j 's are linear functions of the ψ_i 's

Root parameterisation

◀ Skip Durbin back Lag polynomial representation

$$\left(\text{Id} - \sum_{i=1}^p \varrho_i B^i \right) x_t = \epsilon_t$$

with (inverse) roots

$$\prod_{i=1}^p (\text{Id} - \lambda_i B) x_t = \epsilon_t$$

Closed form expression of the likelihood as a function of the (inverse) roots

Uniform prior under stationarity

Stationarity The λ_i 's are within the unit circle if in \mathbb{C} [complex numbers] and within $[-1, 1]$ if in \mathbb{R} [real numbers]

Naturally associated with a flat prior on either the unit circle or $[-1, 1]$

$$\frac{1}{\lfloor k/2 \rfloor + 1} \prod_{\lambda_i \in \mathbb{R}} \frac{1}{2} \mathbb{I}_{|\lambda_i| < 1} \prod_{\lambda_i \notin \mathbb{R}} \frac{1}{\pi} \mathbb{I}_{|\lambda_i| < 1}$$

where $\lfloor k/2 \rfloor + 1$ number of possible cases

⚡ Term $\lfloor k/2 \rfloor + 1$ is important for reversible jump applications

MCMC consequences

In a Gibbs sampler, each λ_{i^*} can be simulated conditionally on the others since

$$\prod_{i=1}^p (\text{Id} - \lambda_i B) x_t = y_t - \lambda_{i^*} y_{t-1} = \epsilon_t$$

where

$$Y_t = \prod_{i \neq i^*} (\text{Id} - \lambda_i B) x_t$$

Metropolis-Hastings implementation

- ① use the prior π itself as a proposal on the (inverse) roots of \mathcal{P} , selecting one or several roots of \mathcal{P} to be simulated from π ;
- ② acceptance ratio is likelihood ratio
- ③ need to watch out for real/complex dichotomy

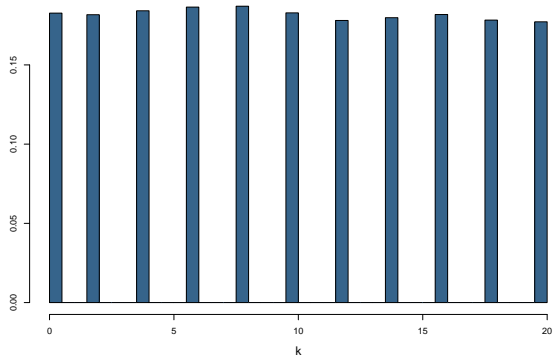
A [paradoxical] reversible jump implementation

- Define “model” \mathfrak{M}_{2k} ($0 \leq k \leq \lfloor p/2 \rfloor$) as corresponding to a number $2k$ of complex roots $0 \leq k \leq \lfloor p/2 \rfloor$)
- Moving from model \mathfrak{M}_{2k} to model \mathfrak{M}_{2k+2} means that two real roots have been replaced by two conjugate complex roots.
- Propose jump from \mathfrak{M}_{2k} to \mathfrak{M}_{2k+2} with probability $1/2$ and from \mathfrak{M}_{2k} to \mathfrak{M}_{2k-2} with probability $1/2$ [*boundary exceptions*]
- accept move from \mathfrak{M}_{2k} to \mathfrak{M}_{2k+} or -2 with probability

$$\frac{\ell^c(\mu, \varrho_1^*, \dots, \varrho_p^*, \sigma | \mathbf{x}_{p:T}, \mathbf{x}_{0:(p-1)})}{\ell^c(\mu, \varrho_1, \dots, \varrho_p, \sigma | \mathbf{x}_{p:T}, \mathbf{x}_{0:(p-1)})} \wedge 1,$$

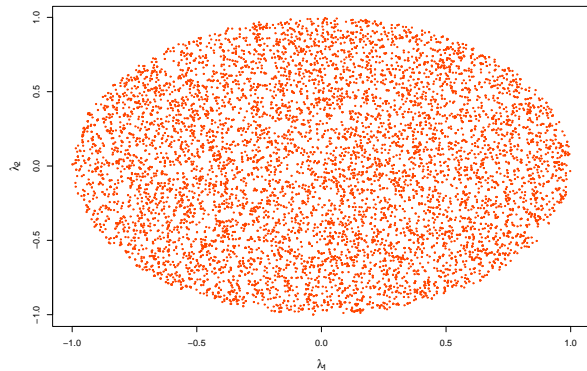
Checking your code

Try with no data and recover the prior



Checking your code

Try with no data and recover the prior



Order estimation

Typical setting for model choice: determine order p of $AR(p)$ model

Roots [may] change drastically from one p to the other.

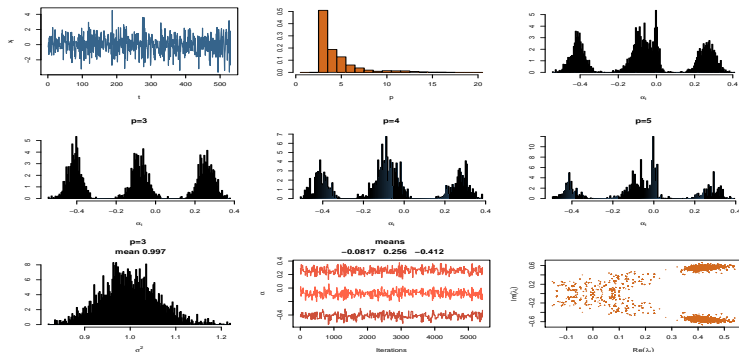
No difficulty from the previous perspective: recycle above reversible jump algorithm

AR(?) reversible jump algorithm

Use (purely birth-and-death) proposals based on the uniform prior

- $k \rightarrow k+1$ [Creation of real root]
- $k \rightarrow k+2$ [Creation of complex root]
- $k \rightarrow k-1$ [Deletion of real root]
- $k \rightarrow k-2$ [Deletion of complex root]

Reversible jump output



$AR(3)$ simulated dataset of 530 points (*upper left*) with true parameters α_i $(-0.1, 0.3, -0.4)$ and $\sigma = 1$. *First histogram* associated with p , following histograms with the α_i 's, for different values of p , and of σ^2 . *Final graph*: scatterplot of the complex roots. *One before last*: evolution of $\alpha_1, \alpha_2, \alpha_3$.

The MA(q) model

Alternative type of time series

$$x_t = \mu + \epsilon_t - \sum_{j=1}^q \vartheta_j \epsilon_{t-j}, \quad \epsilon_t \sim \mathcal{N}(0, \sigma^2)$$

Stationary but, for identifiability considerations, the polynomial

$$Q(x) = 1 - \sum_{j=1}^q \vartheta_j x^j$$

must have all its roots outside the unit circle.

Identifiability

Example

For the MA(1) model, $x_t = \mu + \epsilon_t - \vartheta_1 \epsilon_{t-1}$,

$$\text{var}(x_t) = (1 + \vartheta_1^2)\sigma^2$$

can also be written

$$x_t = \mu + \tilde{\epsilon}_{t-1} - \frac{1}{\vartheta_1} \tilde{\epsilon}_t, \quad \tilde{\epsilon} \sim \mathcal{N}(0, \vartheta_1^2 \sigma^2),$$

Both pairs (ϑ_1, σ) & $(1/\vartheta_1, \vartheta_1 \sigma)$ lead to alternative representations of the *same* model.

Properties of MA models

- Non-Markovian model (but special case of hidden Markov)
- Autocovariance $\gamma_x(s)$ is null for $|s| > q$

Representations

$\mathbf{x}_{1:T}$ is a normal random variable with constant mean μ and covariance matrix

$$\Sigma = \begin{pmatrix} \sigma^2 & \gamma_1 & \gamma_2 & \dots & \gamma_q & 0 & \dots & 0 & 0 \\ \gamma_1 & \sigma^2 & \gamma_1 & \dots & \gamma_{q-1} & \gamma_q & \dots & 0 & 0 \\ & & & \ddots & & & & & \\ 0 & 0 & 0 & \dots & 0 & 0 & \dots & \gamma_1 & \sigma^2 \end{pmatrix},$$

with ($|s| \leq q$)

$$\gamma_s = \sigma^2 \sum_{i=0}^{q-|s|} \vartheta_i \vartheta_{i+|s|}$$

Not manageable in practice [*large T's*]

Representations (contd.)

Conditional on past $(\epsilon_0, \dots, \epsilon_{-q+1})$,

$$L(\mu, \vartheta_1, \dots, \vartheta_q, \sigma | x_{1:T}, \epsilon_0, \dots, \epsilon_{-q+1}) \propto \sigma^{-T} \prod_{t=1}^T \exp \left\{ - \left(x_t - \mu + \sum_{j=1}^q \vartheta_j \hat{\epsilon}_{t-j} \right)^2 / 2\sigma^2 \right\},$$

where $(t > 0)$

$$\hat{\epsilon}_t = x_t - \mu + \sum_{j=1}^q \vartheta_j \hat{\epsilon}_{t-j}, \quad \hat{\epsilon}_0 = \epsilon_0, \quad \dots, \quad \hat{\epsilon}_{1-q} = \epsilon_{1-q}$$

Recursive definition of the likelihood, still costly $O(T \times q)$

Recycling the AR algorithm

Same algorithm as in the AR(p) case when modifying the likelihood

Simulation of the past noises ϵ_{-i} ($i = 1, \dots, q$) done via a Metropolis-Hastings step with target

$$f(\epsilon_0, \dots, \epsilon_{-q+1} | \mathbf{x}_{1:T}, \mu, \sigma, \boldsymbol{\vartheta}) \propto \prod_{i=-q+1}^0 e^{-\epsilon_i^2/2\sigma^2} \prod_{t=1}^T e^{-\tilde{\epsilon}_t^2/2\sigma^2},$$

Representations (contd.)

Encompassing approach for general time series models

State-space representation

$$\mathbf{x}_t = G\mathbf{y}_t + \boldsymbol{\varepsilon}_t, \quad (1)$$

$$\mathbf{y}_{t+1} = F\mathbf{y}_t + \boldsymbol{\xi}_t, \quad (2)$$

(1) is the *observation equation* and (2) is the *state equation*

Note

As seen below, this is a special case of hidden Markov model

MA(q) state-space representation

For the MA(q) model, take

$$\mathbf{y}_t = (\epsilon_{t-q}, \dots, \epsilon_{t-1}, \epsilon_t)'$$

and then

$$\mathbf{y}_{t+1} = \begin{pmatrix} 0 & 1 & 0 & \dots & 0 \\ 0 & 0 & 1 & \dots & 0 \\ & & & \dots & \\ 0 & 0 & 0 & \dots & 1 \\ 0 & 0 & 0 & \dots & 0 \end{pmatrix} \mathbf{y}_t + \epsilon_{t+1} \begin{pmatrix} 0 \\ 0 \\ \vdots \\ 0 \\ 1 \end{pmatrix}$$

$$x_t = \mu - (\vartheta_q \quad \vartheta_{q-1} \quad \dots \quad \vartheta_1 \quad -1) \mathbf{y}_t.$$

MA(q) state-space representation (cont'd)

Example

For the MA(1) model, observation equation

$$x_t = (1 \quad 0)\mathbf{y}_t$$

with

$$\mathbf{y}_t = (y_{1t} \quad y_{2t})'$$

directed by the state equation

$$\mathbf{y}_{t+1} = \begin{pmatrix} 0 & 1 \\ 0 & 0 \end{pmatrix} \mathbf{y}_t + \epsilon_{t+1} \begin{pmatrix} 1 \\ \vartheta_1 \end{pmatrix} .$$

ARMA extension

ARMA(p, q) model

$$x_t - \sum_{i=1}^p \varrho_i x_{t-1} = \mu + \epsilon_t - \sum_{j=1}^q \vartheta_j \epsilon_{t-j}, \quad \epsilon_t \sim \mathcal{N}(0, \sigma^2)$$

Identical stationarity and identifiability conditions for both groups
($\varrho_1, \dots, \varrho_p$) and ($\vartheta_1, \dots, \vartheta_q$)

Reparameterisation

Identical root representations

$$\prod_{i=1}^p (\text{Id} - \lambda_i B) x_t = \prod_{i=1}^q (\text{Id} - \eta_i B) \epsilon_t$$

State-space representation

$$\mathbf{x}_t = x_t = \mu - \begin{pmatrix} \vartheta_{r-1} & \vartheta_{r-2} & \dots & \vartheta_1 & -1 \end{pmatrix} \mathbf{y}_t$$

and

$$\mathbf{y}_{t+1} = \begin{pmatrix} 0 & 1 & 0 & \dots & 0 \\ 0 & 0 & 1 & \dots & 0 \\ & & & \dots & \\ 0 & 0 & 0 & \dots & 1 \\ \varrho_r & \varrho_{r-1} & \varrho_{r-2} & \dots & \varrho_1 \end{pmatrix} \mathbf{y}_t + \epsilon_{t+1} \begin{pmatrix} 0 \\ 0 \\ \vdots \\ 0 \\ 1 \end{pmatrix},$$

under the convention that $\varrho_m = 0$ if $m > p$ and $\vartheta_m = 0$ if $m > q$.

Bayesian approximation

Quasi-identical MCMC implementation:

- 1 Simulate $(\varrho_1, \dots, \varrho_p)$ conditional on $(\vartheta_1, \dots, \vartheta_q)$ and μ
- 2 Simulate $(\vartheta_1, \dots, \vartheta_q)$ conditional on $(\varrho_1, \dots, \varrho_p)$ and μ
- 3 Simulate (μ, σ) conditional on $(\varrho_1, \dots, \varrho_p)$ and $(\vartheta_1, \dots, \vartheta_q)$

© Code can be recycled almost as is!

Hidden Markov models

Generalisation both of a mixture and of a state space model.

Example

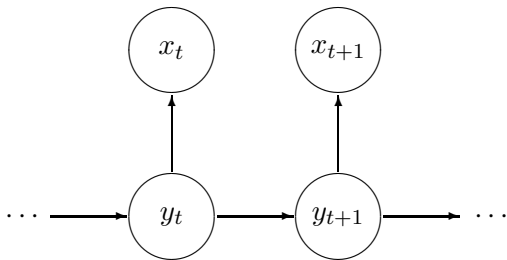
Extension of a *mixture* model with Markov dependence

$$x_t | z, x_j \text{ } j \neq t \sim \mathcal{N}(\mu_{z_t}, \sigma_{z_t}^2), \quad P(z_t = u | z_j, j < t) = p_{z_{t-1}u},$$

$$(u = 1, \dots, k)$$

⚡ Label switching also strikes in this model!

Generic dependence graph



$$(x_t, y_t) | \mathbf{x}_{0:(t-1)}, \mathbf{y}_{0:(t-1)} \sim f(y_t | y_{t-1}) f(x_t | y_t)$$

Definition

Observable series $\{\mathbf{x}_t\}_{t \geq 1}$ associated with a second process $\{y_t\}_{t \geq 1}$, with a finite set of N possible values such that

1. indicators Y_t have an homogeneous **Markov dynamic**

$$p(y_t | \mathbf{y}_{1:t-1}) = p(y_t | y_{t-1}) = \mathbb{P}_{y_{t-1} y_t}$$

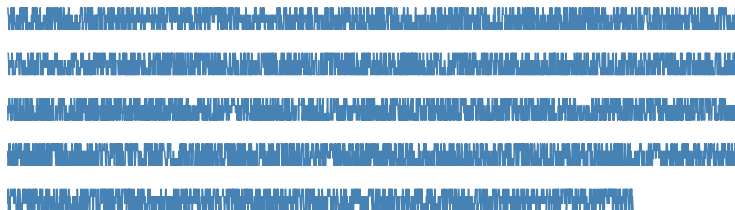
where $\mathbf{y}_{1:t-1}$ denotes the sequence $\{y_1, y_2, \dots, y_{t-1}\}$.

2. Observables x_t are independent conditionally on the indicators y_t

$$p(\mathbf{x}_{1:T} | \mathbf{y}_{1:T}) = \prod_{t=1}^T p(x_t | y_t)$$

Dnadataset

DNA sequence [made of A, C, G, and T's] corresponding to a complete HIV genome where A, C, G, and T have been recoded as 1, ..., 4.



Possible modeling by a two-state hidden Markov model with

$$\mathcal{Y} = \{1, 2\} \quad \text{and} \quad \mathcal{X} = \{1, 2, 3, 4\}$$

Parameterization

- For the Markov bit, *transition matrix*

$$\mathbb{P} = [p_{ij}] \quad \text{where} \quad \sum_{j=1}^N p_{ij} = 1$$

and *initial distribution*

$$\varrho = \varrho \mathbb{P}$$

- for the observables,

$$f_i(x_t) = p(x_t | y_t = i) = f(x_t | \theta_i)$$

usually within the same parametrized class of distributions.

Finite case

When both hidden and observed chains are finite, with

$\mathcal{Y} = \{1, \dots, \kappa\}$ and $\mathcal{X} = \{1, \dots, k\}$, parameter θ made up of p probability vectors $\mathbf{q}^1 = (q_1^1, \dots, q_k^1), \dots, \mathbf{q}^\kappa = (q_1^\kappa, \dots, q_k^\kappa)$

Joint distribution of $(x_t, y_t)_{0 \leq t \leq T}$

$$p_{y_0} q_{x_0}^{y_0} \prod_{t=1}^T p_{y_{t-1}y_t} q_{x_t}^{y_t},$$

Bayesian inference in the finite case

Posterior of (θ, \mathbb{P}) given $(x_t, y_t)_t$ factorizes as

$$\pi(\theta, \mathbb{P}) \propto_{y_0} \prod_{i=1}^{\kappa} \prod_{j=1}^k (q_j^i)^{n_{ij}} \times \prod_{i=1}^{\kappa} \prod_{j=1}^p p_{ij}^{m_{ij}},$$

where n_{ij} # of visits to state j by the x_t 's when the corresponding y_t 's are equal to i and m_{ij} # of transitions from state i to state j on the hidden chain $(y_t)_{t \in \mathbb{N}}$

Under a flat prior on p_{ij} 's and q_j^i 's, posterior distributions are [almost] Dirichlet [*initial distribution side effect*]

MCMC implementation

Finite State HMM Gibbs Sampler

Initialization:

- ① Generate random values of the p_{ij} 's and of the q_j^i 's
- ② Generate the hidden Markov chain $(y_t)_{0 \leq t \leq T}$ by $(i = 1, 2)$

$$\mathbb{P}(y_t = i) \propto \begin{cases} p_{ii} q_{x_0}^i & \text{if } t = 0, \\ p_{y_{t-1}i} q_{x_t}^i & \text{if } t > 0, \end{cases}$$

and compute the corresponding sufficient statistics

MCMC implementation (cont'd)

Finite State HMM Gibbs Sampler

Iteration m ($m \geq 1$):

- 1 Generate

$$(p_{i1}, \dots, p_{i\kappa}) \sim \mathcal{D}(1 + n_{i1}, \dots, 1 + n_{i\kappa})$$

$$(q_1^i, \dots, q_k^i) \sim \mathcal{D}(1 + m_{i1}, \dots, 1 + m_{ik})$$

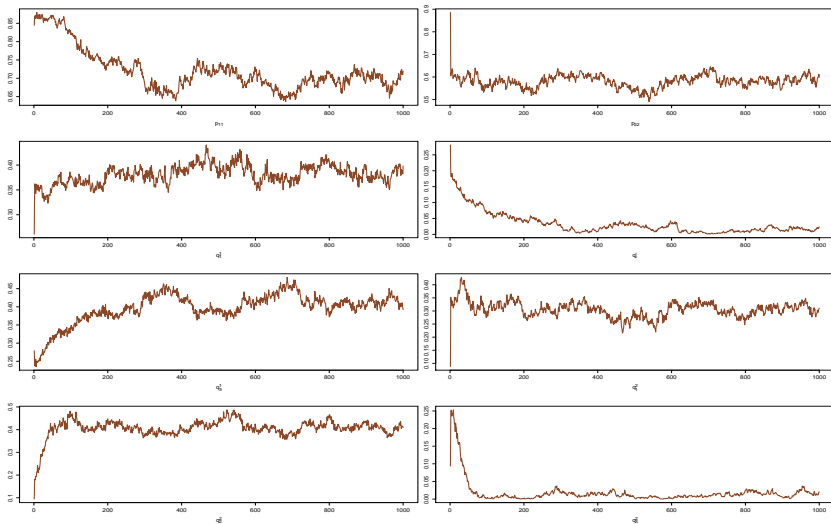
and correct for missing initial probability by a MH step with acceptance probability q'_{y_0}/q_{y_0}

- 2 Generate successively each y_t ($0 \leq t \leq T$) by

$$\mathbb{P}(y_t = i | x_t, y_{t-1}, y_{t+1}) \propto \begin{cases} p_{ii} q_{x_1}^i p_{iy_1} & \text{if } t = 0, \\ p_{y_{t-1}i} q_{x_t}^i p_{iy_{t+1}} & \text{if } t > 0, \end{cases}$$

and compute corresponding sufficient statistics

Dnadatast



Forward-Backward formulae

Existence of a (magical) recurrence relation that provides the observed likelihood function in manageable computing time
Called *forward-backward* or *Baum–Welch* formulas

Observed likelihood computation

Likelihood of the *complete model* simple:

$$\ell^c(\boldsymbol{\theta}|\mathbf{x}, \mathbf{y}) = \prod_{t=2}^T p_{y_{t-1}y_t} f(x_t|\theta_{y_t})$$

but likelihood of the *observed model* is not:

$$\ell(\boldsymbol{\theta}|\mathbf{x}) = \sum_{\mathbf{y} \in \{1, \dots, \kappa\}^T} \ell^c(\boldsymbol{\theta}|\mathbf{x}, \mathbf{y})$$

© $O(\kappa^T)$ complexity

Forward-Backward paradox

It is possible to express the (observed) likelihood $L^O(\boldsymbol{\theta}|\mathbf{x})$ in

$$O(T^2 \times \kappa)$$

computations, based on the Markov property of the pair (x_t, y_t) .

► [Direct to backward smoothing](#)

Conditional distributions

We have

$$p(\mathbf{y}_{1:t} | \mathbf{x}_{1:t}) = \frac{f(x_t | y_t) p(\mathbf{y}_{1:t} | \mathbf{x}_{1:(t-1)})}{p(x_t | \mathbf{x}_{1:(t-1)})}$$

[Smoothing/Bayes]

and

$$p(\mathbf{y}_{1:t} | \mathbf{x}_{1:(t-1)}) = k(y_t | \mathbf{y}_{t-1}) p(\mathbf{y}_{1:(t-1)} | \mathbf{x}_{1:(t-1)})$$

[Prediction]

where $k(y_t | \mathbf{y}_{t-1}) = p_{y_{t-1} y_t}$ associated with the matrix \mathbb{P} and

$$f(x_t | y_t) = f(x_t | \theta_{y_t})$$

Update of predictive

Therefore

$$\begin{aligned} p(\mathbf{y}_{1:t} | \mathbf{x}_{1:t}) &= \frac{p(y_t | \mathbf{x}_{1:(t-1)}) f(x_t | y_t)}{p(x_t | \mathbf{x}_{1:(t-1)})} \\ &= \frac{f(x_t | y_t) k(y_t | \mathbf{y}_{t-1})}{p(x_t | \mathbf{x}_{1:(t-1)})} p(\mathbf{y}_{1:(t-1)} | \mathbf{x}_{1:(t-1)}) \end{aligned}$$

with the same order of complexity for $p(\mathbf{y}_{1:t} | \mathbf{x}_{1:t})$ as for $p(x_t | \mathbf{x}_{1:(t-1)})$

Propagation and actualization equations

$$p(y_t | \mathbf{x}_{1:(t-1)}) = \sum_{\mathbf{y}_{1:(t-1)}} p(\mathbf{y}_{1:(t-1)} | \mathbf{x}_{1:(t-1)}) k(y_t | y_{t-1})$$

[Propagation]

and

$$p(y_t | x_{1:t}) = \frac{p(y_t | \mathbf{x}_{1:(t-1)}) f(x_t | y_t)}{p(x_t | \mathbf{x}_{1:(t-1)})}.$$

[Actualization]

Forward-backward equations (1)

Evaluation of

$$p(y_t | \mathbf{x}_{1:T}) \quad t \leq T$$

by *forward-backward algorithm*

Denote $t \leq T$

$$\gamma_t(i) = P(y_t = i | x_{1:T})$$

$$\alpha_t(i) = p(\mathbf{x}_{1:t}, y_t = i)$$

$$\beta_t(i) = p(\mathbf{x}_{t+1:T} | y_t = i)$$

Recurrence relations

Then

$$\begin{cases} \alpha_1(i) &= f(x_1|y_1 = i) \varrho_i \\ \alpha_{t+1}(j) &= f(x_{t+1}|y_{t+1} = j) \sum_{i=1}^{\kappa} \alpha_t(i) p_{ij} \end{cases}$$

[Forward]

$$\begin{cases} \beta_T(i) &= 1 \\ \beta_t(i) &= \sum_{j=1}^{\kappa} p_{ij} f(x_{t+1}|y_{t+1} = j) \beta_{t+1}(j) \end{cases}$$

[Backward]

and

$$\gamma_t(i) = \frac{\alpha_t(i) \beta_t(i)}{\sum_{j=1}^{\kappa} \alpha_t(j) \beta_t(j)}$$

Extension of the recurrence relations

For

$$\xi_t(i, j) = P(y_t = i, y_{t+1} = j | \mathbf{x}_{1:T}) \quad i, j = 1, \dots, \kappa,$$

we also have

$$\xi_t(i, j) = \frac{\alpha_t(i) \mathbb{P}_{ij} f(x_{t+1} | y_t = j) \beta_{t+1}(j)}{\sum_{i=1}^{\kappa} \sum_{j=1}^{\kappa} \alpha_t(i) \mathbb{P}_{ij} f(x_{t+1} | y_{t+1} = j) \beta_{t+1}(j)}$$

Overflows and underflows

⚡ On-line scalings of the $\alpha_t(i)$'s and $\beta_T(i)$'s for each t by

$$c_t = 1 / \sum_{i=1}^{\kappa} \alpha_t(i) \quad \text{and} \quad d_t = 1 / \sum_{i=1}^{\kappa} \beta_t(i)$$

avoid overflows or/and underflows for large datasets

Backward smoothing

Recursive derivation of conditionals

We have

$$p(y_s | y_{s-1}, \mathbf{x}_{1:t}) = p(y_s | y_{s-1}, \mathbf{x}_{s:t})$$

[Markov property!]

Therefore ($s = T, T - 1, \dots, 1$)

$$p(y_s | y_{s-1}, \mathbf{x}_{1:T}) \propto k(y_s | y_{s-1}) f(x_s | y_s) \sum_{y_{s+1}} p(y_{s+1} | y_s, \mathbf{x}_{1:T})$$

[Backward equation]

with

$$p(y_T | y_{T-1}, \mathbf{x}_{1:T}) \propto k(y_T | y_{T-1}) f(x_T | y_T).$$

End of the backward smoothing

The first term is

$$p(y_1 | \mathbf{x}_{1:t}) \propto \pi(y_1) f(x_1 | y_1) \sum_{y_2} p(y_2 | y_1, \mathbf{x}_{1:t}),$$

with π stationary distribution of \mathbb{P}

The conditional for y_s needs to be defined for each of the κ values of y_{s-1}

© $O(t \times \kappa^2)$ operations

Details

Need to introduce unnormalized version of the conditionals $p(y_t|y_{t-1}, \mathbf{x}_{0:T})$ such that

$$p_T^*(y_T|y_{T-1}, \mathbf{x}_{0:T}) = p_{y_{T-1}y_T} f(x_T|y_T)$$

$$p_t^*(y_t|y_{t-1}, \mathbf{x}_{1:T}) = p_{y_{t-1}y_t} f(x_t|y_t) \sum_{i=1}^{\kappa} p_{t+1}^*(i|y_t, \mathbf{x}_{1:T})$$

$$p_0^*(y_0|\mathbf{x}_{0:T}) = \varrho_{y_0} f(x_0|y_0) \sum_{i=1}^{\kappa} p_1^*(i|y_0, \mathbf{x}_{0:t})$$

Likelihood computation

Bayes formula

$$p(\mathbf{x}_{1:T}) = \frac{p(\mathbf{x}_{1:T}|\mathbf{y}_{1:T})p(\mathbf{y}_{1:T})}{p(\mathbf{y}_{1:T}|\mathbf{x}_{1:T})}$$



gives a representation of the likelihood based on the forward–backward formulae and an arbitrary sequence $\mathbf{x}_{1:T}^o$ (since the l.h.s. does *not* depend on $\mathbf{x}_{1:T}$).

Obtained through the p_t^* 's as

$$p(\mathbf{x}_{0:T}) = \sum_{i=1}^{\kappa} p_1^*(i|\mathbf{x}_{0:T})$$

Prediction filter

If

$$\varphi_t(i) = p(y_t = i | \mathbf{x}_{1:t-1})$$

Forward equations

$$\begin{aligned}\varphi_1(j) &= p(y_1 = j) \\ \varphi_{t+1}(j) &= \frac{1}{c_t} \sum_{i=1}^{\kappa} f(x_t | y_t = i) \varphi_t(i) p_{ij} \quad (t \geq 1)\end{aligned}$$

where

$$c_t = \sum_{k=1}^{\kappa} f(x_t | y_t = k) \varphi_t(k),$$

Likelihood computation (2)

Follows the same principle as the backward equations

The (log-)likelihood is thus

$$\begin{aligned}\log p(\mathbf{x}_{1:t}) &= \sum_{r=1}^t \log \left[\sum_{i=1}^{\kappa} p(x_t, y_t = i | \mathbf{x}_{1:(r-1)}) \right] \\ &= \sum_{r=1}^t \log \left[\sum_{i=1}^{\kappa} f(x_t | y_t = i) \varphi_t(i) \right]\end{aligned}$$