The Whittle-Levinson-Durbin Recursion

R. J. Kinnear

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Abstract

A brief overview of the Levinson-Durbin recursion for estimating autoregressive time series models is given. Whittle's generalized (multivariate) version is also expounded upon. We are essentially summarizing content from [1], [2], [3].

It will be seen that the algorithm itself provides deep insights into the structure of autoregressive time series models, and provides an indispensable algorithm in practice.

Contents

Todo list

1 introduction

Suppose we have a process $x(t) \in \mathbb{R}^n$ generated by the all-pole model

$$\sum_{\tau=0}^{p} A(\tau)x(t-\tau) = v_t, \tag{1}$$

where $B(0) = I_n$ and v_t is a temporally uncorrelated driving sequence with $\mathbb{E}[v_t] = 0$. This is closely connected to fitting $VAR(p)$ time series models

$$\widehat{x}(t) = \sum_{\tau=1}^{p} B(\tau)x(t-\tau), \tag{2}$$

where one can estimate A and then simply drop $A(0) = I$ and take $B = -A$.

If we observe only $x(t)$, how do we determine $A(\tau)$? An answer is provided by the Levinson-Durbin recursion. This algorithm is extremely efficient as it allows one to fit a sequence of $VAR(p)$ models for every $p = 1, \ldots, p_{\max}$ all for the cost of inverting a single toeplitz (or block-toeplitz) matrix. This implies that there is effectively no additional cost for performing model selection (i.e. choosing p) over and above what it costs to fit a single $VAR(p_{\max})$ model.

1.1 Yule-Walker Equations

The Levinson-Durbin recursion is essentially an efficient procedure for solving the Yule-Walker equations in one dimension, and Whittle's generalization extends to the multivariate case. The Yule-Walker equations are simply

$$\sum_{\tau=0}^{p} A(\tau)R(s-\tau) = \delta_s \Sigma_v; s = 0, 1, \ldots, p, \tag{3}$$

which describes the relationship between the coefficients A of Equation (2) and the covariance sequence $R(\tau) = \mathbb{E}[x(t)x(t-\tau)^{\mathsf{T}}]$. They can be derived easily by taking the model (2) and multiplying it on the right by $x(t-\tau)^{\mathsf{T}}$ and computing the expectation.

There is a close relationship between the Yule-Walker equations and Toeplitz matrices: If one is to write out equation 3 in a large matrix format, the resulting system is a toeplitz (or block-toeplitz in the multivariate case) system[1]

$$\mathbf{RA} = e_1 \otimes \Sigma_v, \tag{4}$$

where $\mathbf{A} = [I \ A(1)^{\mathsf{T}} \ \cdots \ A(p)^{\mathsf{T}}]^{\mathsf{T}}$ and $\mathbf{R}$ is a (symmetric) block-toeplitz matrix consisting of blocks[2] $[\mathbf{R}]_{s\tau} = R(s-\tau)^{\mathsf{T}}$, and $e_1 \otimes \Sigma_v = [\Sigma_v \ 0 \ \cdots \ 0]$. It is critical to notice that the variables in this equation are $A(1), \ldots, A(p)$ *and* Σ_v, so it is not written in the usual "$Ax = b$" format, but is still a linear equation. In the unidimensional case we can write

$$Ra = \sigma_v e_1, \tag{5}$$

where R is a bona-fide toeplitz matrix.

[1] The symbol $\otimes$ indicates the Kronecker product

[2] One must keep careful track of transposes.

1.2 Estimating Covariances

Given a finite sample of data $\{x(t)\}_{t=1}^{T}$, we can treat this as an infinitely extended sequence $\widetilde{x}(t)$ where $\widetilde{x}(t) = 0$ for $t \leq 0$ or $t > T$ (i.e. a rectangularly windowed sequence) and then estimate the covariance via

$$\widehat{R}(\tau) = \frac{1}{T} \sum_{t=\tau+1}^{p} x(t)x(t-\tau)^{\mathsf{T}}. \tag{6}$$

It is critical to use this particular covariance estimator in order to ensure that $R(\tau)$ is a positive (semi-)definite sequence, that is, the Toeplitz matrix $\mathbf{R}$ satisfies $\mathbf{R} \succeq 0$.

2 The Recursions

2.1 The Levinson-Durbin Recursion

We will first write down the Levinson-Durbin recursion, which is the unidimensional method for solving equation 3. We will consider the "API" for this Algorithm as taking as input a sequence of $p+1$ covariance estimates $\big(r(0), r(1), \ldots, r(p)\big)$ for a unidimensional ($n = 1$) time series $x(t)$, and returning $p+1$ variance estimates $\sigma_0^2, \sigma_1^2, \ldots, \sigma_p^2$, as well as a sequence $\mathbf{b}_0, \mathbf{b}_1, \ldots, \mathbf{b}_p$ where $\mathbf{b}_k \in \mathbb{R}^k$ provides coefficients for an $AR(k)$ model of order k, where the estimated mean squared error of this model is given by σ_k^2

Double check results for σ_k^2

$$\begin{aligned} \widehat{x}_k(t) &= \sum_{\tau=1}^{k} b_k(\tau)x(t-\tau), \\ \sigma_k^2 &= \frac{1}{T}\sum_{t=1}^{T}(x(t) - \widehat{x}_k(t))^2 \\ &= \mathbb{E}(x(t) - \widehat{x}_k(t))^2 + O\big(\frac{1}{\sqrt{T}}\big). \end{aligned} \tag{7}$$

The algorithm is also applicable when $x(t) \in \mathbb{C}^n$, therefore in Algorithm 1 $*$ indicates complex conjugate. It is important to keep in mind that $r(-\tau) = r(\tau)^*$

2.1.1 Properties

The algorithm runs in $O(p^2)$ time (whereas standard matrix inversion requires $O(p^3)$ time). As well there are a number of remarkable properties associated to Algorithm 1:

1. $|b_k(k)| \leq 1$ if and only if $r(0), \ldots, r(k)$ is positive semi-definite for $k = 1, \ldots, p$
2. $\sigma_k^2 \geq 0$ if and only if $r(0), \ldots, r(k)$ is positive semi-definite for $k = 1, \ldots, p$
3. The $AR(k)$ model obtained from $\mathbf{b}_k$ is stable

Algorithm 1: Levinson-Durbin Recursion

input : Covariance Sequence $r(0), \ldots, r(p)$
output : AR coefficients $\mathbf{b}_1, \ldots, \mathbf{b}_p$ and error estimates $\sigma_0^2, \ldots, \sigma_p^2$
initialize: $a_0(0) = 1$
$\sigma_0^2 = r(0)$

for $k = 0, \ldots, p-1$ **do**
  $\gamma = \sum_{\tau=0}^{k} a_k(\tau) r(k - \tau + 1)$
  $a_{k+1}(k+1) = -\gamma / \sigma_k^2$ `# Reflection Coefficient`
  $a_{k+1}(0) = 1$
  for $\tau = 1, \ldots, k$ **do**
    $a_{k+1}(\tau) = a_k(\tau) + a_{k+1}(k+1) a_k^*(k - \tau + 1)$ `# Copy to next array`
  $\sigma_{k+1}^2 = \sigma_k^2 (1 - |a_{k+1}(k+1)|^2)$
  $\mathbf{b}_{k+1} = \big(- a_{k+1}(1), \ldots, -a_{k+1}(k+1) \big)$ `# Convert to VAR Coefficients`
  assert $\sum_{\tau=0}^{k+1} a_{k+1}(\tau) r(s - \tau) = 0$; for $s = 1, \ldots, k+1$ `# Verify`
return $\big(\mathbf{b}_1, \ldots, \mathbf{b}_p\big)$, $(\sigma_0^2, \ldots, \sigma_p^2)$

2.2 Whittle's Generalization

Whittle [3] generalized Algorithm 1 to the multivariate case. This generalization is non-trivial and requires both a *forwards* set of coefficients $A(\tau)$, but also a *backwards* set of coefficients $\bar{A}(\tau)$ corresponding to the anti-causal system

$$\sum_{\tau=0}^{p} \bar{A}(\tau) x(t+\tau) = \bar{v}_t. \tag{8}$$

The most direct reason that the Levinson-Durbin recursion does not immediately generalize is simply because matrix multiplication is not commutative.

The algorithm will consume a sequence of covariance matrices $R(0), \ldots, R(p)$, and return a sequence $\mathbf{B}_1, \ldots, \mathbf{B}_p$ of $VAR(k)$ model coefficients, where $\mathbf{B}_k = \big(B_k(1), \ldots, B_k(k)\big)$ as well as a sequence $\Sigma_0, \ldots, \Sigma_p$ of error variance matrices where

Double check results for Σ_k

$$\begin{aligned} \widehat{x}_k(t) &= \sum_{\tau=1}^{k} B_k(\tau) x(t-\tau), \\ \Sigma_k &= \frac{1}{T} \sum_{t=1}^{T} (x(t) - \widehat{x}_k(t))(x(t) - \widehat{x}_k(t))^{\mathsf{T}} \\ &= \mathbb{E}[(x(t) - \widehat{x}_k(t))(x(t) - \widehat{x}_k(t))^{\mathsf{T}}] + O\big(\frac{1}{\sqrt{T}}\big). \end{aligned} \tag{9}$$

Keeping in mind that $R(-\tau) = R(\tau)^{\mathsf{H}}$, we have

Algorithm 2: Whittle-Levinson-Durbin Recursion

input : Covariance Sequence $R(0), \ldots, R(p)$
output : AR coefficients $\mathbf{B}_1, \ldots, \mathbf{B}_p$ and error estimates $\Sigma_0, \ldots, \Sigma_p$
initialize: $A_0(0) = I, \bar{A}_0(0) = I$
$\Sigma_0 = R(0), \bar{\Sigma}_0 = R(0)$

for $k = 0, \ldots, p-1$ **do**
  $\Gamma = \sum_{\tau=0}^{k} A_k(\tau) R(k - \tau + 1)$
  $\bar{\Gamma} = \sum_{\tau=0}^{k} \bar{A}_k(\tau) R(\tau - k - 1)$
  $A_{k+1}(k+1) = -\Gamma \bar{\Sigma}_k^{-1}$ `# Use 'cholesky' and 'cho_solve' to invert`
  $\bar{A}_{k+1}(k+1) = -\bar{\Gamma} \Sigma_k^{-1}$
  $A_{k+1}(0) = I, \bar{A}_{k+1}(0) = I$
  for $\tau = 1, \ldots, k$ **do**
    $A_{k+1}(\tau) = A_k(\tau) + A_{k+1}(k+1) \bar{A}_k(k - \tau + 1)$ `# Copy to next array`
    $\bar{A}_{k+1}(\tau) = \bar{A}_k(\tau) + \bar{A}_{k+1}(k+1) A_k(k - \tau + 1)$
  $\Sigma_{k+1} = \Sigma_k + A_{k+1}(k+1) \bar{\Gamma}$ `# Update the error variance`
  $\bar{\Sigma}_{k+1} = \bar{\Sigma}_k + \bar{A}_{k+1}(k+1) \Gamma$
  $\mathbf{B}_{k+1} = \big(-A_{k+1}(1), \ldots, -A_{k+1}(k+1) \big)$ `# Convert to VAR coefficients`
  assert $\sum_{\tau=0}^{k+1} A_{k+1}(\tau) R(s - \tau) = 0;$ for $s = 1, \ldots, k+1$ `# Verify`
  assert $\sum_{\tau=0}^{k+1} \bar{A}_{k+1}(\tau) R(\tau - s) = 0;$ for $s = 1, \ldots, k+1$
return $\big(\mathbf{B}_1, \ldots, \mathbf{B}_p\big)$, $(\Sigma_0, \ldots, \Sigma_p)$

3 Efficient Model Order Selection

Given a collection of data, there are a variety of methods to choose the filter order p. For example, the Bayesian Information Criteria (BIC) provides a "meta-optimization" objective which trades off between error reduction and model complexity. Choosing p by the BIC criterion dictates that we minimize over p the following:

$$BIC(p) = T \log\det \Sigma_p + n^2 p \log T. \tag{10}$$

This can be carried out by a simple direct search on each model order between 1 and some prescribed $p_{\max}$. In practice it is sufficient to pick $p_{\max}$ ad-hoc or via some simple heuristic (e.g. plotting the sequence $BIC(p)$ over p). Conveniently, the Levinson-Durbin algorithm provides to us the sequence of error matrices Σ_p all in one go, so choosing p by this criteria takes time equivalent to simply fitting a single $VAR(p_{\max})$ model.

It must finally be pointed out that this is not a perfect strategy for model order selection, and is likely only useful when n is quite small; it is merely one that is efficiently facilitated by the LD algorithm. One can in principle make use of the same BIC criterion to further search for sparse matrices that provide a good fit for the data, but this becomes a much different problem.