

Lecture 10


ARMA Models: Estimation, Diagnostics, and Model Selection

Reading: Bowerman, O'Connell, and Koehler (2005): Chapter 10.1-10.2; Cryer and Chen (2008): Chapter 7.3-7.5; Chapter 8.1

MATH 4070: Regression and Time-Series Analysis

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ARMA Models: Estimation, Diagnostics, and Model Selection



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Parameter Estimation
Model Diagnostics and Selection


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Notes

Agenda

- 1 Parameter Estimation
- 2 Model Diagnostics and Selection

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Notes

Estimation of the ARMA Process Parameters

Suppose we choose an ARMA(p, q) model for a zero-mean $\{\eta_t\}$

- Need to estimate the $p + q + 1$ parameters:
 - AR component $\{\phi_1, \dots, \phi_p\}$
 - MA component $\{\theta_1, \dots, \theta_q\}$
 - $\text{Var}(Z_t) = \sigma^2$
- One strategy:
 - Do some preliminary estimation of the model parameters (e.g., via [Yule-Walker](#) estimates)
 - Follow-up with [maximum likelihood estimation](#) with Gaussian assumption

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Notes

The Yule-Walker Method

Suppose η_t is a **causal** AR(p) process

$$\eta_t - \phi_1 \eta_{t-1} - \dots - \phi_p \eta_{t-p} = Z_t$$

To estimate the parameters $\{\phi_1, \dots, \phi_p\}$, we use a **method of moments** estimation scheme:

- Let $h = 0, 1, \dots, p$. We multiply η_{t-h} to both sides

$$\eta_t \eta_{t-h} - \phi_1 \eta_{t-1} \eta_{t-h} - \dots - \phi_p \eta_{t-p} \eta_{t-h} = Z_t \eta_{t-h}$$

- Taking expectations:

$$\mathbb{E}(\eta_t \eta_{t-h}) - \phi_1 \mathbb{E}(\eta_{t-1} \eta_{t-h}) - \dots - \phi_p \mathbb{E}(\eta_{t-p} \eta_{t-h}) = \mathbb{E}(Z_t \eta_{t-h}),$$

we get

$$\gamma(h) - \phi_1 \gamma(h-1) - \dots - \phi_p \gamma(h-p) = \mathbb{E}(Z_t \eta_{t-h})$$

Notes

The Yule-Walker Equations

- When $h = 0$, $\mathbb{E}(Z_t \eta_{t-h}) = \text{Cov}(Z_t, \eta_t) = \sigma^2$ (Why?)

Therefore, we have

$$\gamma(0) - \sum_{j=1}^p \phi_j \gamma(j) = \sigma^2$$

- When $h > 0$, Z_t is uncorrelated with η_{t-h} (because the **assumption of causality**), thus $\mathbb{E}(Z_t \eta_{t-h}) = 0$ and we have

$$\gamma(h) - \sum_{j=1}^p \phi_j \gamma(h-j) = 0, \quad h = 1, 2, \dots, p$$

- The **Yule-Walker estimates** are the solution of these equations when we replace $\gamma(h)$ by $\hat{\gamma}(h)$

Notes

The Yule-Walker Equations in Matrix Form

Let $\hat{\phi} = (\hat{\phi}_1, \dots, \hat{\phi}_p)^T$ be an estimate for $\phi = (\phi_1, \dots, \phi_p)^T$ and let

$$\hat{\Gamma} = \begin{bmatrix} \hat{\gamma}(0) & \hat{\gamma}(1) & \dots & \hat{\gamma}(p-1) \\ \hat{\gamma}(1) & \hat{\gamma}(0) & \dots & \hat{\gamma}(p-2) \\ \vdots & \vdots & \ddots & \vdots \\ \hat{\gamma}(p-1) & \hat{\gamma}(p-2) & \dots & \hat{\gamma}(0) \end{bmatrix}$$

Then the **Yule-Walker estimates** of ϕ and σ^2 are

$$\hat{\phi} = \hat{\Gamma}^{-1} \hat{\gamma},$$

and

$$\hat{\sigma}^2 = \hat{\gamma}(0) - \hat{\phi}^T \hat{\gamma},$$

where $\hat{\gamma} = (\hat{\gamma}(1), \dots, \hat{\gamma}(p))^T$

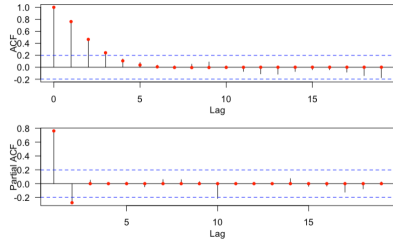
Notes

Lake Huron Example in R

```

#### R
YW_est <- ar(lm$residuals, aic = F, order.max = 2, method = "yw")
# plot sample and estimated acf/pacf
par(las = 1, mgp = c(2.2, 1, 0), mar = c(3.6, 3.6, 0.6, 0.6), mfrow = c(2, 1))
acf(lm$residuals)
acf_YWest <- ARMAacf(ar = YW_est$ar, lag.max = 23)
points(0:23, acf_YWest, col = "red", pch = 16, cex = 0.8)
pacf(lm$residuals)
pacf_YWest <- ARMAacf(ar = YW_est$ar, lag.max = 23, pacf = T)
points(1:23, pacf_YWest, col = "red", pch = 16, cex = 0.8)

```



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Notes

Remarks on the Yule-Walker Method

- For large sample size, Yule-Walker estimator have (approximately) the same sampling distribution as maximum likelihood estimator (MLE), but with small sample size Yule-Walker estimator can be far less efficient than the MLE
- The Yule-Walker method is a poor procedure for MA(q) and ARMA(p, q) processes with $q > 0$ (see Cryer Chan 2008, p. 150-151)
- We move on the more versatile and popular method for estimating ARMA(p, q) parameters—maximum likelihood estimation¹

¹See Least Squares Estimation in Chapter 7.2 of Cryer and Chan (2008).

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Maximum Likelihood Estimation

- The setup:
 - Model: $X = (X_1, X_2, \dots, X_n)$ has joint probability density function $f(x; \omega)$ where $\omega = (\omega_1, \omega_2, \dots, \omega_p)$ is a vector of p parameters
 - Data: $x = (x_1, x_2, \dots, x_n)$

- The likelihood function is defined as the the “likelihood” of the data, x , given the parameters, ω

$$L_n(\omega) = f(x; \omega)$$

- The maximum likelihood estimate (MLE) is the value of ω which maximizes the likelihood, $L_n(\omega)$, of the data x :

$$\hat{\omega} = \underset{\omega}{\operatorname{argmax}} L_n(\omega).$$

It is equivalent (and often easier) to maximize the log likelihood,

$$\ell_n(\omega) = \log L_n(\omega)$$

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Notes

The MLE for an i.i.d. Gaussian Process

Suppose $\{X_t\}$ be a Gaussian i.i.d. process with mean μ and variance σ^2 . We observe a time series

$$\mathbf{x} = (x_1, \dots, x_n)^T.$$

- The likelihood function is

$$\begin{aligned} L_n(\mu, \sigma^2) &= f(\mathbf{x}|\mu, \sigma^2) \\ &= \prod_{t=1}^n f(x_t|\mu, \sigma) \\ &= \prod_{t=1}^n \left\{ \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left[-\frac{(x_t - \mu)^2}{2\sigma^2}\right] \right\} \\ &= (2\pi)^{-n/2} (\sigma^2)^{-n/2} \exp\left[-\frac{\sum_{t=1}^n (x_t - \mu)^2}{2\sigma^2}\right] \end{aligned}$$

- The log-likelihood function is

$$\begin{aligned} \ell_n(\mu, \sigma^2) &= \log L_n(\mu, \sigma^2) \\ &= -\frac{n}{2} \log(2\pi) - \frac{n}{2} \log(\sigma^2) - \frac{\sum_{t=1}^n (x_t - \mu)^2}{2\sigma^2} \end{aligned}$$

$$\Rightarrow \hat{\mu}_{MLE} = \frac{\sum_{t=1}^n X_t}{n} = \bar{X}, \quad \hat{\sigma}_{MLE}^2 = \frac{\sum_{t=1}^n (X_t - \bar{X})^2}{n}$$

Notes

Likelihood for Stationary Gaussian Time Series Models

Suppose $\{X_t\}$ be a mean zero stationary Gaussian time series with ACVF $\gamma(h)$. If $\gamma(h)$ depends on p parameters,

$$\boldsymbol{\omega} = (\omega_1, \dots, \omega_p)$$

- The likelihood of the data $\mathbf{x} = (x_1, \dots, x_n)$ given the parameters $\boldsymbol{\omega}$ is

$$L_n(\boldsymbol{\omega}) = (2\pi)^{-n/2} |\boldsymbol{\Gamma}|^{-1/2} \exp\left(-\frac{1}{2} \mathbf{x}^T \boldsymbol{\Gamma}^{-1} \mathbf{x}\right),$$

where $\boldsymbol{\Gamma}$ is the covariance matrix of $\mathbf{X} = (X_1, \dots, X_n)^T$, $|\boldsymbol{\Gamma}|$ is the determinant of the matrix $\boldsymbol{\Gamma}$, and $\boldsymbol{\Gamma}^{-1}$ is the inverse of the matrix $\boldsymbol{\Gamma}$

- The log-likelihood is

$$\ell_n(\boldsymbol{\theta}) = -\frac{n}{2} \log(2\pi) - \frac{1}{2} \log |\boldsymbol{\Gamma}| - \frac{1}{2} \mathbf{x}^T \boldsymbol{\Gamma}^{-1} \mathbf{x}$$

Typically need to solve it numerically

Notes

Decomposing Joint Density into Conditional Densities

A joint distribution can be represented as the product of conditionals and a marginal distribution

- The simple version for $n = 2$ is:

$$f(x_1, x_2) = f(x_2|x_1)f(x_1)$$

- Extending for general n we get the following expression for the likelihood:

$$L_n(\boldsymbol{\theta}) = f(\mathbf{x}; \boldsymbol{\theta}) = f(x_1) \prod_{t=2}^n f(x_t|x_{t-1}, \dots, x_1; \boldsymbol{\theta}),$$

and the log-likelihood is

$$\ell_n(\boldsymbol{\theta}) = \log f(\mathbf{x}; \boldsymbol{\theta}) = \log(f(x_1)) + \sum_{t=2}^n \log f(x_t|x_{t-1}, \dots, x_1; \boldsymbol{\theta}).$$

Notes

AR(1) Log-likelihood

Let $\{\eta_1, \eta_2, \dots, \eta_n\}$ be a realization of a zero-mean stationary AR(1) Gaussian time series. Let $\theta = (\phi, \sigma^2)$

$$\ell_n(\theta) = \underbrace{\log(f(\eta_1))}_{\ell_{n,1}} + \underbrace{\sum_{t=2}^n \log f(\eta_t | \eta_{t-1}, \dots, \eta_1; \theta)}_{\ell_{n,2}}$$

Note that for $t \geq 2$, $f(\eta_t | \eta_{t-1}, \dots, \eta_1) = f(\eta_t | \eta_{t-1})$, where $[\eta_t | \eta_{t-1}] \sim N(\phi\eta_{t-1}, \sigma^2) \Rightarrow \ell_{n,2} =$

$$-\frac{(n-1)}{2} \log 2\pi - \frac{(n-1)}{2} \log \sigma^2 - \frac{\sum_{t=2}^n (\eta_t - \phi\eta_{t-1})^2}{2\sigma^2}$$

Also, we know $[\eta_1] \sim N(0, \frac{\sigma^2}{1-\phi^2}) \Rightarrow \ell_{1,n} =$

$$-\frac{\log 2\pi}{2} - \frac{\log \sigma^2}{2} + \frac{\log(1-\phi^2)}{2} - \frac{(1-\phi^2)\eta_1^2}{2\sigma^2}$$

$$\Rightarrow \ell_n(\theta) = -\frac{n}{2} \log 2\pi - \frac{n}{2} \log \sigma^2 - \frac{\sum_{t=2}^n (\eta_t - \phi\eta_{t-1})^2}{2\sigma^2} + \frac{\log(1-\phi^2)}{2} - \frac{(1-\phi^2)\eta_1^2}{2\sigma^2}$$

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AR(1) Log-likelihood Cont'd

$$\ell_n(\theta) = -\frac{n}{2} \log 2\pi - \frac{n}{2} \log \sigma^2 + \frac{\log(1-\phi^2)}{2} - \frac{S(\phi)}{2\sigma^2},$$

where $S(\phi) = \sum_{t=2}^n (\eta_t - \phi\eta_{t-1})^2 + (1-\phi^2)\eta_1^2$

- For given value of ϕ , $\ell_n(\phi, \sigma^2)$ can be maximized analytically with respect to σ^2

$$\hat{\sigma}^2 = \frac{S(\hat{\phi})}{n}$$

- Estimation of ϕ can be simplified by maximizing the **conditional sum-of-squares** ($\sum_{t=2}^n (\eta_t - \phi\eta_{t-1})^2$)
- Standard errors** can be obtained by computing the inverse of the **Hessian matrix**: $\text{Var}(\hat{\theta}) = H(\hat{\theta})^{-1}$, where $H(\theta) = \frac{\partial^2 \ell_n(\theta)}{\partial \theta \partial \theta^T}$

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Notes

arima in R with the Lake Huron Example

arima: ARIMA Modelling of Time Series

Description

Fits an ARIMA model to a univariate time series.

Usage

```
arima(x, order = c(p, d, q), m0,
      residuals = list(order = c(p, d, q), m0, period = 1),
      fit.method = "ML",
      fit.control = list(),
      fixed = NULL, full = FALSE,
      method = c("ML", "REML", "GLS", "REML", "GLS", "REML"),
      model = c("ARMA", "ARIMA", "ARMA", "ARIMA"),
      plot.method = "none",
      plot.control = list(), main = NULL)
```

```
## {r}
(ML_est1 <- arima(Lm$residuals, order = c(2, 0, 0), method = "ML"))
##
```

Call:

```
arima(x = Lm$residuals, order = c(2, 0, 0), method = "ML")
```

Coefficients:

```
ar1 ar2 intercept
1.0047 -0.2919 0.0197
s.e. 0.0977 0.1004 0.2350
```

sigma^2 estimated as 0.4571: log likelihood = -101.25, aic = 210.5

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Notes

Inference for the ARMA Parameters

Motivating example: What is an approximate 95% CI for ϕ_1 in an AR(1) model?

- Let $\phi = (\phi_1, \dots, \phi_p)$ and $\theta = (\theta_1, \dots, \theta_q)$ denote the ARMA parameters (excluding σ^2), and let $\hat{\phi}$ and $\hat{\theta}$ be the ML estimates of ϕ and θ . Then for "large" n , $(\hat{\phi}, \hat{\theta})$ have approximately a **joint normal** distribution:

$$\begin{bmatrix} \hat{\phi} \\ \hat{\theta} \end{bmatrix} \sim N\left(\begin{bmatrix} \phi \\ \theta \end{bmatrix}, \frac{V(\phi, \theta)}{n}\right)$$

- $V(\phi, \theta)$ is a known $(p+q) \times (p+q)$ matrix depending on the ARMA parameters

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Notes

$V(\phi, \theta)$ for AR Processes

- For an AR(p) process

$$V(\phi) = \sigma^2 \Gamma^{-1},$$

where Γ is the $p \times p$ covariance matrix of the series (η_1, \dots, η_p)

- AR(1) process:

$$V(\phi_1) = 1 - \phi_1^2$$

- AR(2) process:

$$V(\phi_1, \phi_2) = \begin{bmatrix} 1 - \phi_2^2 & -\phi_1(1 + \phi_2) \\ -\phi_1(1 + \phi_2) & 1 - \phi_2^2 \end{bmatrix}$$

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Other Examples of $V(\phi, \theta)$

- MA(1) process:

$$V(\theta_1) = 1 - \theta_1^2$$

- MA(2) process:

$$V(\theta_1, \theta_2) = \begin{bmatrix} 1 - \theta_2^2 & \theta_1(1 - \theta_2) \\ \theta_1(1 - \theta_2) & 1 - \theta_2^2 \end{bmatrix}$$

- Casual and invertible ARMA(1,1) process

$$V(\phi, \theta) = \frac{1 + \phi\theta}{(\phi + \theta)^2} \begin{bmatrix} (1 - \phi^2)(1 + \phi\theta) & -(1 - \phi^2)(1 - \theta^2) \\ -(1 - \phi^2)(1 - \theta^2) & 1 - \theta_2^2 \end{bmatrix}$$

- More generally, for "small" n , the covariance matrix of $(\hat{\phi}, \hat{\theta})$ can be approximated using the second derivatives of the log-likelihood function, known as the **Hessian matrix**

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Notes

MLE for Trend and Temporal Correlation in One Step

```
```{r}
(MLE_est4 <- arima(LakeHuron, order = c(2, 0, 0), xreg = yr))
```
```

Call:
arima(x = LakeHuron, order = c(2, 0, 0), xreg = yr)

Coefficients:

| | ar1 | ar2 | intercept | xreg |
|------|--------|---------|-----------|---------|
| | 1.0048 | -0.2913 | 620.5115 | -0.0216 |
| s.e. | 0.0976 | 0.1004 | 15.5771 | 0.0081 |

sigma^2 estimated as 0.4566: log likelihood = -101.2, aic = 210.4

Fitted model:

$$Y_t = 620.51 - 0.022\text{Year} + \eta_t,$$

where

$$\eta_t = 1.00\eta_{t-1} - 0.29\eta_{t-2} + Z_t, \quad Z_t \sim N(0, \sigma^2 = 0.46^2).$$

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Notes

What About Non-Gaussian Processes?

It is more challenging to express the joint distribution of X_t for non-Gaussian processes. Instead, we often rely on the [Gaussian likelihood](#) as an [approximate likelihood](#)

- In practice:
 - [Transform](#) the data to make the series as close to Gaussian as possible (e.g., using a log, square-root, or Box-Cox transformation)
 - Then use the [Gaussian likelihood](#) to estimate parameters, assuming the transformed series follows a near-Gaussian structure
 - For many real-world applications, this approximation works well and simplifies estimation. However, [residual diagnostics](#) are needed to ensure the model fits the data adequately

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Notes

Assessing Fit / Comparing Different Time Series Models

- We can use diagnostic plots for the “residuals” of the fitted time series, along with [Box tests](#) to assess whether an i.i.d. process is reasonable

```
> Box.test(YW_est$resid[-(1:2)], type = "Ljung-Box")
```

Box-Ljung test

```
data: YW_est$resid[-(1:2)]
X-squared = 0.56352, df = 1, p-value = 0.4528
```

- Use [confidence intervals](#) for the parameters. Intervals that contain zero may indicate that we can simplify the model
- We can also use model selection criteria, such as [AIC](#), to compare between different models

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Notes

Diagnostics via the Time Series Residuals

- Recall the innovations are given by

$$U_t = X_t - \hat{X}_t$$

- Under a **Gaussian** model, $\{U_t : t = 1, \dots, T\}$ is an independent set of RVs with

$$U_t \sim N(0, \nu_{t-1}) \stackrel{d}{=} \sigma N(0, r_{t-1}).$$

- Define the **residuals** $\{R_t\}$ by

$$R_t = \frac{U_t}{\sqrt{r_{t-1}}} = \frac{X_t - \hat{X}_t}{\sqrt{r_{t-1}}}$$

Under Gaussian model $R_t \stackrel{i.i.d.}{\sim} N(0, \sigma^2)$

Notes

ARMA Order Selection

- We would prefer to use models that compromise between a small residual error $\hat{\sigma}^2$ and a small number of parameters $(p + q + 1)$
- To choose the order $(p$ and $q)$ of ARMA model it makes sense to penalize models with a large number of parameters
- Here we consider an information based criteria to compare models

Notes

Akaike Information Criterion (AIC)

- The **Akaike information criterion** (AIC) is defined by

$$\text{AIC} = -2\ell_n(\hat{\phi}, \hat{\theta}, \hat{\sigma}^2) + 2(p + q + 1)$$

- We choose the values of p and q that **minimizes** the AIC value
- For $\text{AR}(p)$ models, AIC tends to overestimate p . The bias corrected version is

$$\text{AIC}_c = 2\ell_n(\hat{\phi}, \hat{\theta}, \hat{\sigma}^2) + \frac{2n(p + q + 1)}{(n - 1) - (p + q + 1)}$$

Notes

Lake Huron Example: AIC and AICc

```
m1 <- arima(LakeHuron, order = c(1, 0, 0), xreg = yr)
m2 <- arima(LakeHuron, order = c(1, 0, 1), xreg = yr)
m3 <- arima(LakeHuron, order = c(2, 0, 0), xreg = yr)
m4 <- arima(LakeHuron, order = c(2, 0, 1), xreg = yr)
AIC(m1); AIC(m2); AIC(m3); AIC(m4)
library(MuMIn)
AICc(m1); AICc(m2); AICc(m3); AICc(m4)
```

```

```
[1] 218.4501
[1] 212.3954
[1] 212.3965
[1] 214.0638
[1] 218.8803
[1] 213.0476
[1] 213.0487
[1] 214.9868
```

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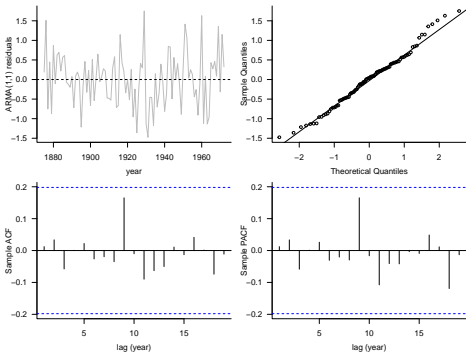
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### Lake Huron Model Diagnostics



```
> Box.test(resids, lag = 10, type = "Ljung-Box")
```

Box-Ljung test

```
data: resids
X-squared = 3.7882, df = 10, p-value = 0.9564
```

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