

Lecture 10

ARMA Models: Estimation, Diagnostics, and Model Selection

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

MATH 4070: Regression and Time-Series Analysis

Whitney Huang
Clemson University

Agenda

ARMA Models:
Estimation,
Diagnostics, and
Model Selection



Parameter Estimation

Model Diagnostics and
Selection

1 Parameter Estimation

2 Model Diagnostics and Selection

Estimation of the ARMA Process Parameters

Suppose we choose an $\text{ARMA}(p, q)$ 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

Suppose η_t is a **causal** $\text{AR}(p)$ process

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

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

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

- Taking expectations:

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

we get $\boxed{\gamma(h) - \phi_1\gamma(h-1) - \cdots - \phi_p\gamma(h-p) = \mathbb{E}(Z_t\eta_{t-h})}$

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 $\hat{\gamma}(h)$

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

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

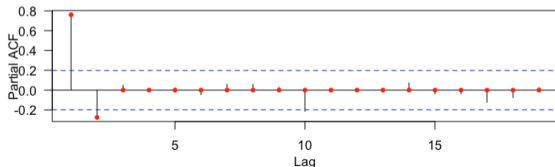
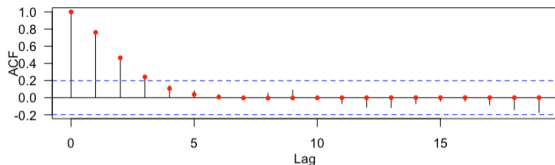
and

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

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

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)
``
```



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

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

- 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)

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

- 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).

Maximum Likelihood Estimation

- The setup:

ARMA Models:
Estimation,
Diagnostics, and
Model Selection



Parameter Estimation

Model Diagnostics and
Selection

Maximum Likelihood Estimation

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

Maximum Likelihood Estimation

ARMA Models:
Estimation,
Diagnostics, and
Model Selection



Parameter Estimation

Model Diagnostics and
Selection

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

- The setup:
 - Model: $\mathbf{X} = (X_1, X_2, \dots, X_n)$ has joint probability density function $f(\mathbf{x}; \boldsymbol{\omega})$ where $\boldsymbol{\omega} = (\omega_1, \omega_2, \dots, \omega_p)$ is a vector of p parameters
 - Data: $\mathbf{x} = (x_1, x_2, \dots, x_n)$
- The **likelihood function** is defined as the the “likelihood” of the data, \mathbf{x} , given the parameters, $\boldsymbol{\omega}$

$$L_n(\boldsymbol{\omega}) = f(\mathbf{x}; \boldsymbol{\omega})$$

- The setup:
 - Model: $\mathbf{X} = (X_1, X_2, \dots, X_n)$ has joint probability density function $f(\mathbf{x}; \boldsymbol{\omega})$ where $\boldsymbol{\omega} = (\omega_1, \omega_2, \dots, \omega_p)$ is a vector of p parameters
 - Data: $\mathbf{x} = (x_1, x_2, \dots, x_n)$
- The **likelihood function** is defined as the the “likelihood” of the data, \mathbf{x} , given the parameters, $\boldsymbol{\omega}$

$$L_n(\boldsymbol{\omega}) = f(\mathbf{x}; \boldsymbol{\omega})$$

- The **maximum likelihood estimate** (MLE) is the value of $\boldsymbol{\omega}$ which maximizes the likelihood, $L_n(\boldsymbol{\omega})$, of the data \mathbf{x} :

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

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

$$\ell_n(\boldsymbol{\omega}) = \log L_n(\boldsymbol{\omega})$$

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 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

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

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

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

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

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

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

Likelihood for Stationary Gaussian Time Series Models

ARMA Models:
Estimation,
Diagnostics, and
Model Selection



Parameter Estimation

Model Diagnostics and
Selection

Suppose $\{X_t\}$ be a mean zero **stationary Gaussian** time series with ACVF $\gamma(h)$. If $\gamma(h)$ depends on p parameters, $\omega = (\omega_1, \dots, \omega_p)$

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

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

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

- The log-likelihood is

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

Typically need to solve it numerically

Decomposing Joint Density into Conditional Densities

ARMA Models:
Estimation,
Diagnostics, and
Model Selection

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



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

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

Parameter Estimation

Model Diagnostics and
Selection

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

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

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 $\boldsymbol{\theta} = (\phi, \sigma^2)$

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

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 $[\eta_t | \eta_{t-1}] \sim N(\phi\eta_{t-1}, \sigma^2) \Rightarrow \ell_{n,2} =$

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

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

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

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

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

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

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

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

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

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

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** $(\sum_{t=2}^n (\eta_t - \phi\eta_{t-1})^2)$

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

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**: $\text{Var}(\hat{\boldsymbol{\theta}}) = H(\hat{\boldsymbol{\theta}})^{-1}$, where $H(\boldsymbol{\theta}) = \frac{\partial^2 \ell_n(\boldsymbol{\theta})}{\partial \boldsymbol{\theta} \partial \boldsymbol{\theta}^T}$

arima in R with the Lake Huron Example

ARMA Models:
Estimation,
Diagnostics, and
Model Selection



arima: ARIMA Modelling of Time Series

Description

Fit an ARIMA model to a univariate time series.

Usage

```
arima(x, order = c(0L, 0L, 0L),
      seasonal = list(order = c(0L, 0L, 0L), period = NA),
      xreg = NULL, include.mean = TRUE,
      transform.pars = TRUE,
      fixed = NULL, init = NULL,
      method = c("CSS-ML", "ML", "CSS"), n.cond,
      SSinit = c("Gardner1980", "Rossignol2011"),
      optim.method = "BFGS",
      optim.control = list(), kappa = 1e6)
```

Parameter Estimation

Model Diagnostics and
Selection

arima in R with the Lake Huron Example

ARMA Models:
Estimation,
Diagnostics, and
Model Selection



Parameter Estimation

Model Diagnostics and
Selection

arima: ARIMA Modelling of Time Series

Description

Fit an ARIMA model to a univariate time series.

Usage

```
arima(x, order = c(0L, 0L, 0L),
      seasonal = list(order = c(0L, 0L, 0L), period = NA),
      xreg = NULL, include.mean = TRUE,
      transform.pars = TRUE,
      fixed = NULL, init = NULL,
      method = c("CSS-ML", "ML", "CSS"), n.cond,
      SSinit = c("Gardner1980", "Rossignol2011"),
      optim.method = "BFGS",
      optim.control = list(), kappa = 1e6)
```

```
```{r}
(MLE_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

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:

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

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

- 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:

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

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**

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

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

Parameter Estimation

Model Diagnostics and
Selection

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

- 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
```

- 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 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

Diagnostics via the Time Series Residuals

ARMA Models:
Estimation,
Diagnostics, and
Model Selection

- Recall the innovations are given by

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



Parameter Estimation

Model Diagnostics and
Selection

- 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

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

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

- We would prefer to use models that compromise between a small residual error $\hat{\sigma}^2$ and a small number of parameters $(p + q + 1)$

- 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

- 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

- 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

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

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



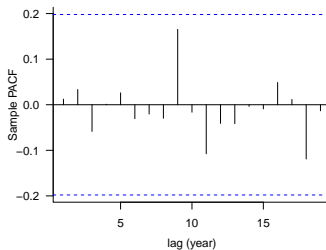
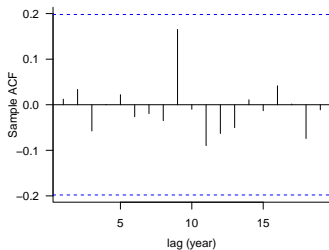
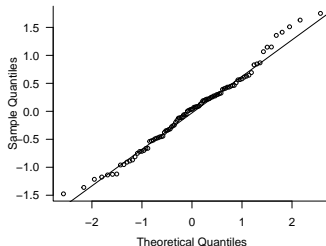
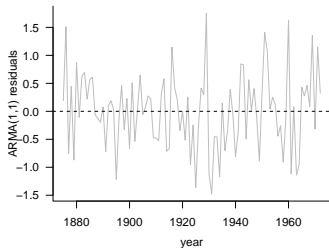
# Lake Huron Model Diagnostics

ARMA Models:  
Estimation,  
Diagnostics, and  
Model Selection



Parameter Estimation

Model Diagnostics and  
Selection



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

Box-Ljung test

```
data: resids
```

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