# MATH 8090: Stationary processes and Model-Free Estimation of Stationary Means and Covariances

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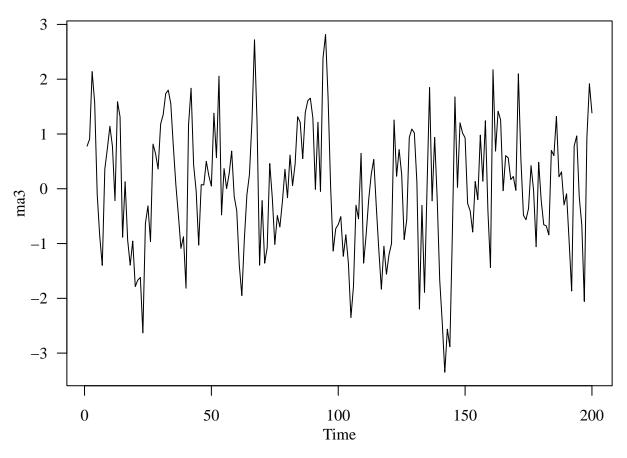
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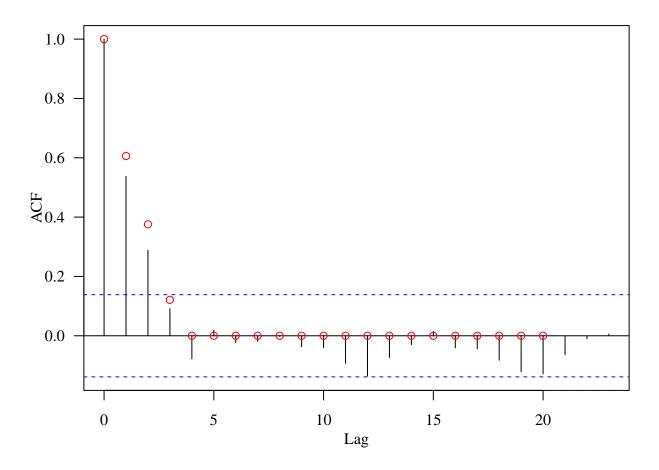
# **Stationary Processes**

## Examples of MA(q) processes

```
set.seed(123)
ma3 <- arima.sim(n = 200, list(ma = c(0.6, 0.5, 0.2)))
par(mar = c(3, 3.5, 0.5, 0.6), mgp = c(2, 1, 0), las = 1, family = "serif")
ts.plot(ma3)</pre>
```

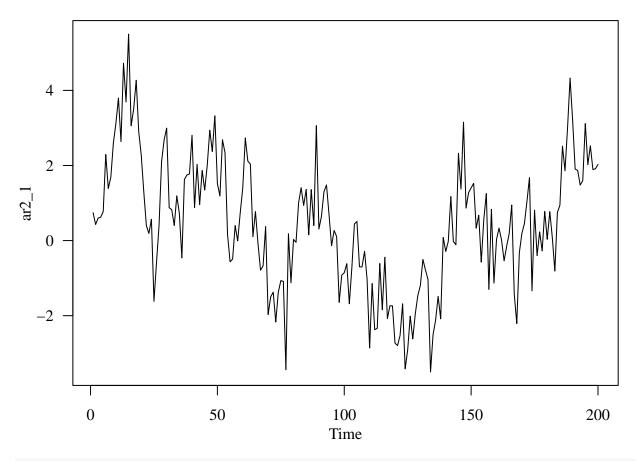


```
acf(ma3)
acf_true <- ARMAacf(ma = c(0.6, 0.5, 0.2), lag.max = 20)
points(0:20, acf_true, col = "red")</pre>
```

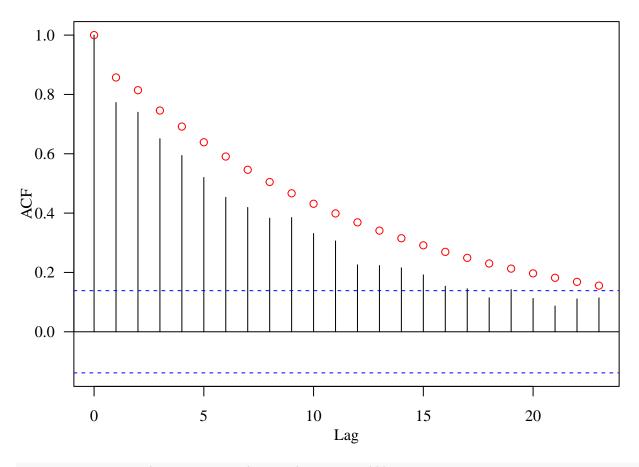


# Examples of AR(2) processes

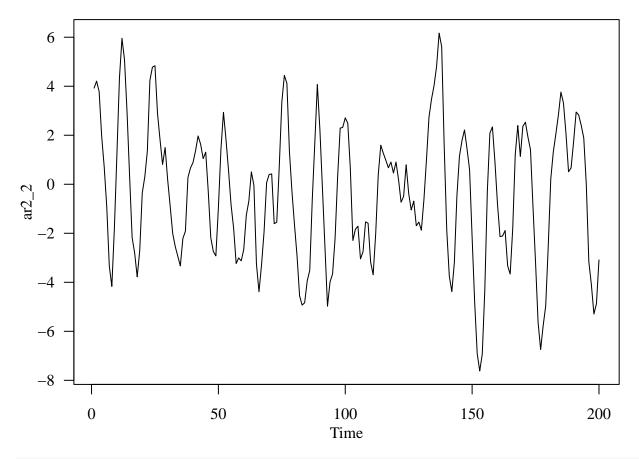
```
par(mar = c(3, 3.5, 0.5, 0.6), mgp = c(2, 1, 0), las = 1, family = "serif")
ar2_1 <- arima.sim(n = 200, list(ar = c(0.6, 0.3)))
ts.plot(ar2_1)</pre>
```



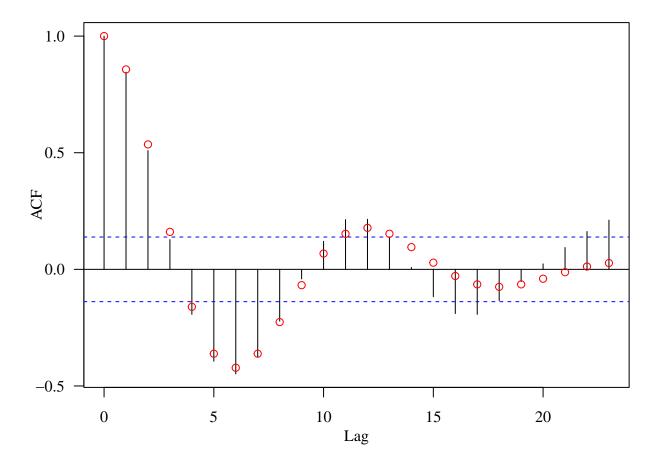
acf(ar2\_1)
points(0:23, ARMAacf(ar = c(0.6, 0.3), lag.max = 23), col = "red")



 $ar2_2 \leftarrow arima.sim(n = 200, list(ar = c(1.5, -0.75)))$ ts.plot(ar2\_2)



```
acf(ar2_2)
points(0:23, ARMAacf(ar = c(1.5, -0.75), lag.max = 23), col = "red")
```



#### Model-Free Estimation of Stationary Means and Covariances

#### Mean Estimation

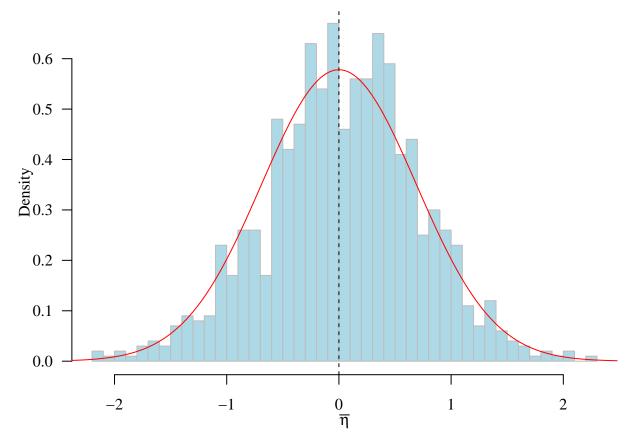
Given a stationary process  $\{\eta_t\}_{t=1}^T$ , the point estimator is  $\bar{\eta} = \frac{1}{T} \sum_{t=1}^T \eta_t$ . The varaince of this estimator is

$$\nu_T = \operatorname{Var}(\bar{\eta}) = \operatorname{Var}\left(\frac{1}{T} \sum_{t=1}^T \eta_t\right) = \frac{1}{T} \sum_{h=-(T-1)}^{T-1} \left(1 - \frac{|h|}{T}\right) \gamma(h)$$

#### ## [1] 0.4659671

```
# Theoretical sampling dist
mu = 0
h <- -(T-1):(T-1)</pre>
```

```
nu_T <- (1 / T) * sum((1 - (abs(h) / T)) * (phi^(abs(h)) / (1 - phi^2)))
## Superimpose the true density curve
xg <- seq(-2.5, 2.5, 0.01)
abline(v = mu, lty = 2)
lines(xg, dnorm(xg, sd = sqrt(nu_T)), col = "red")</pre>
```



```
## Compare nu_T and nu
(nu_T <- (1 / T) * sum((1 - (abs(h) / T)) * (phi^(abs(h)) / (1 - phi^2))))
```

## [1] 0.4763158

```
(nu <- (1 / T) * (1 / (1 - phi)^2))
```

## [1] 0.5

#### Autocovariance Function (ACVF)

```
gamma <- apply(sim, 2, function(x) acf(x, plot = F, type = "covariance")$acf)
(est <- apply(gamma, 1, mean))

## [1] 4.762396047 4.210822098 3.718189717 3.276955489 2.883257698 2.529033768
## [7] 2.215391725 1.934657288 1.684410252 1.460895692 1.261443145 1.085257775
## [13] 0.927192703 0.791189571 0.671906177 0.563884323 0.468728429 0.380609541</pre>
```

## [19] 0.300366800 0.227258194 0.161799558 0.106047046 0.056009807 0.007812902

```
(true \leftarrow ARMAacf(ar = c(0.9), lag.max = 23) * (1 / (1 - phi^2)))
                               2
##
           0
                     1
                                         3
                                                              5
                                                                        6
## 5.2631579 4.7368421 4.2631579 3.8368421 3.4531579 3.1078421 2.7970579 2.5173521
                     9
                              10
                                        11
                                                  12
                                                             13
## 2.2656169 2.0390552 1.8351497 1.6516347 1.4864712 1.3378241 1.2040417 1.0836375
          16
                    17
                              18
                                        19
                                                  20
                                                             21
## 0.9752738 0.8777464 0.7899718 0.7109746 0.6398771 0.5758894 0.5183005 0.4664704
# The sample ACVF is a biased estimator
rg <- range(true, est)
par(mar = c(3, 3.5, 0.5, 0.6), mgp = c(2, 1, 0), las = 1, family = "serif")
plot(true, est, pch = 16, xlim = rg, ylim = rg, las = 1)
abline(0, 1, col = "blue")
   5 -
   4
   3
est
   2 -
    1 -
   0
         0
                                      2
                                                    3
                                                                  4
                                                                                5
                        1
                                             true
# ACVF is non-negative definite
h <- outer(1:24, 1:24, "-")
Sigma <- phi^abs(h) / (1 - phi^2)
eigen(Sigma)$values
   [1] 64.7315984 26.5011371 11.7018213 6.1924112 3.7670568
                                                                 2.5262589
        1.8165664 1.3759195 1.0849122 0.8833909 0.7385536
##
   [7]
                                                                 0.6313427
```

**##** [13] 0.5500994 0.4873821 0.4382698 0.3994115 0.3684691

## [19] 0.3241390 0.3086742 0.2967447 0.2878873 0.2817747 0.2781898

#### Autocorrelation Function (ACF)

 $Population\ ACF$ 

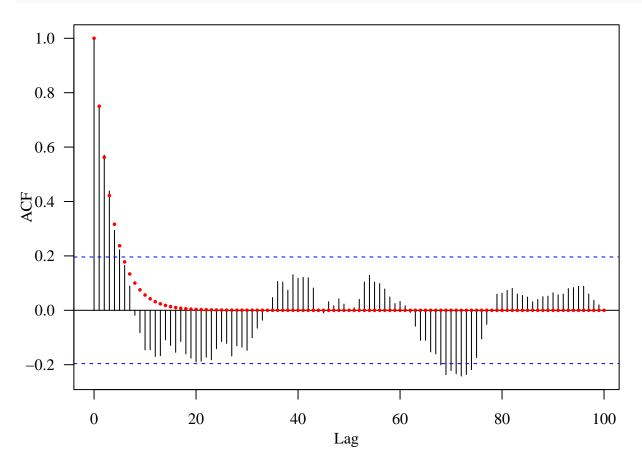
$$\rho(h) = \operatorname{Cor}(\eta_t, \eta_{t+h}) = \frac{\operatorname{E}\left[(\eta_t - \mu)(\eta_{t+h} - \mu)\right]}{\sqrt{\operatorname{Var}(\eta_t)\operatorname{Var}(\eta_{t+h})}}$$

 $Sample\ ACF$ 

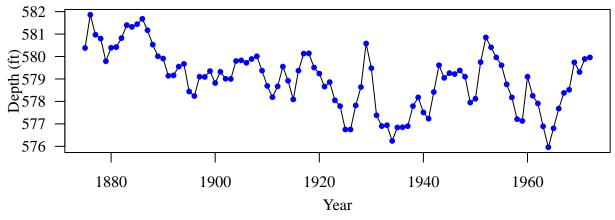
$$\hat{\rho}(h) = \frac{\hat{\gamma}(h)}{\hat{\gamma}(0)},$$

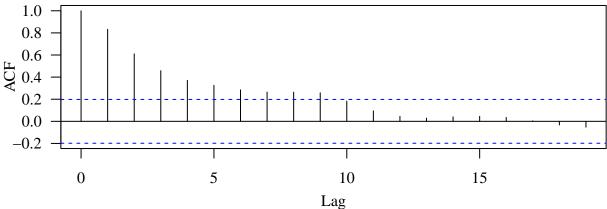
where  $\hat{\gamma}(h) = \frac{1}{T} \sum_{t=1}^{T-|h|} (\eta_t - \bar{\eta})(\eta_{t+h} - \bar{\eta}).$ 

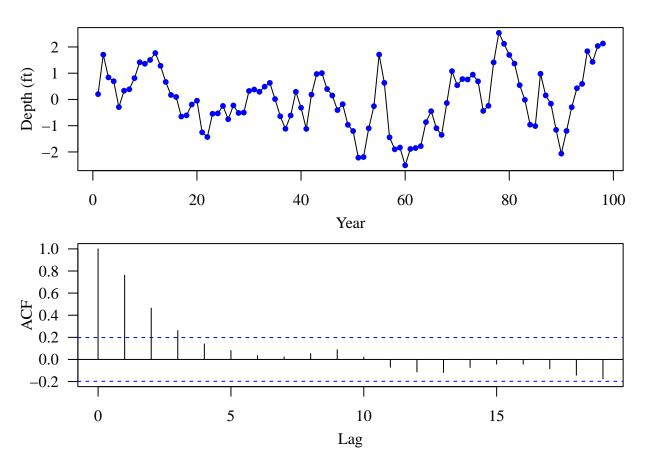
```
set.seed(123)
AR1 <- arima.sim(n = 100, list(ar = c(0.75)))
par(mar = c(3, 3.5, 0.5, 0.6), mgp = c(2, 1, 0), las = 1, family = "serif")
acf(AR1, lag = 100)
acf_true <- ARMAacf(ar = c(0.75), lag.max = 100)
points(0:100, acf_true, pch = 16, cex = 0.5, col = "red")</pre>
```

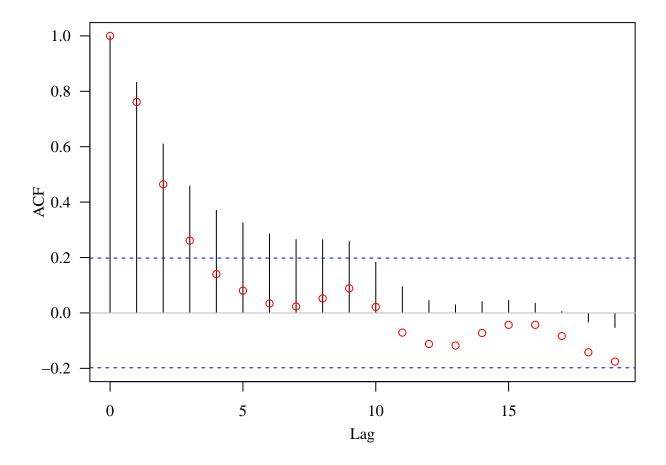


Lake Huron Example









## Box test for temporal independence

# Box and Pierce test Box and Pierce (1970)

We wish to test:

 $H_0: \{\eta_1, \eta_2, \cdots, \eta_T\}$  is an i.i.d. noise sequence

 $H_1: H_0$  is false

1. Under  $H_0$ ,

$$\hat{\rho}(h) \stackrel{.}{\sim} \mathrm{N}(0, \frac{1}{T}) \stackrel{d}{=} \frac{1}{\sqrt{n}} \mathrm{N}(0, 1)$$

2. Hence

$$Q = T \sum_{i=1}^{k} \hat{\rho}^2(h) \stackrel{\cdot}{\sim} \chi_{df=k}^2$$

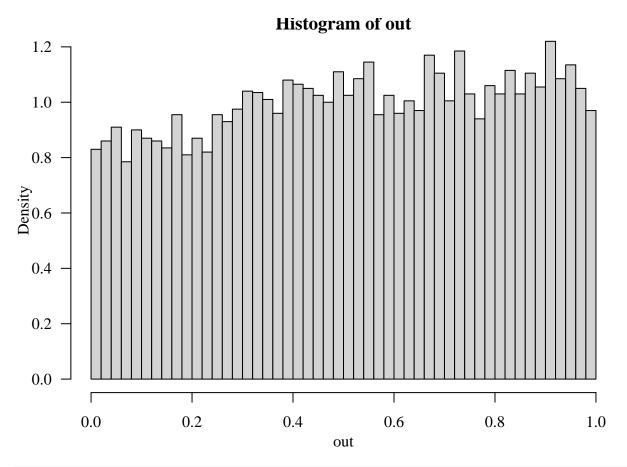
3. We reject  $H_0$  if  $Q > \chi_k^2(1-\alpha)$ , the  $1-\alpha$  quatile of the chi-squared distribution with k degrees of freedom

#### Ljung-Box Test

$$Q_{LB} = T(T-2) \sum_{h=1}^{k} \frac{\hat{\rho}^2(h)}{n-h} \stackrel{.}{\sim} \chi_k^2.$$

The Ljung-Box test Ljung and Box (1978) can be more powerful than the Box and Pierce test

```
out <- numeric(10000)
for (i in 1:10000) out[i] <- Box.test(rnorm(100), 5)$p.value
par(mar = c(3, 3.5, 0.5, 0.6), mgp = c(2, 1, 0), las = 1, family = "serif")
hist(out, 50, prob = T)</pre>
```



```
Box.test(lm$residuals, 10)
```

```
##
## Box-Pierce test
##
## data: lm$residuals
## X-squared = 88.469, df = 10, p-value = 1.077e-14

Box.test(lm$residuals, 10, type = "Ljung")

##
## Box-Ljung test
##
## data: lm$residuals
## X-squared = 91.776, df = 10, p-value = 2.331e-15

library(WeightedPortTest)
Weighted.Box.test(lm$residuals, lag = 10)
```

```
##
## Weighted Box-Pierce test (Gamma Approximation)
##
## data: lm$residuals
## Weighted X-squared on Residuals for fitted ARMA process = 83.252, Shape
## = 3.9286, Scale = 1.4000, p-value < 2.2e-16</pre>
```

#### References

Box, George EP, and David A Pierce. 1970. "Distribution of Residual Autocorrelations in Autoregressive-Integrated Moving Average Time Series Models." *Journal of the American Statistical Association* 65 (332): 1509–26.

Ljung, Greta M, and George EP Box. 1978. "On a Measure of Lack of Fit in Time Series Models." *Biometrika* 65 (2): 297–303.