Lab 4 - Learning Via Computer Simulation

Benjamin M. Taylor, Kevin Hayes January 20, 2023

1 AR(1) Process

The following R code simulates and plots an AR(1) time series process of length n=1001 having parameter $\phi=0.7$ with $\sigma_e^2=1$, plots the empirical / estimated ACF (correlogram) and PACF (partial correlogram) and their theoretical counterparts. Write the code into an R script file and execute the code from there.

```
AR1.acf = ARMAacf(ar = alpha, lag.max = 25)

AR1.pacf = ARMAacf(ar = alpha, lag.max = 25, pacf = TRUE)

names(AR1.pacf) = names(AR1.acf)[-1]

layout(mat= matrix(c(1,1,2:5), byrow=TRUE, ncol=2))

plot(Yt, main= "Simulated AR(1) Process")

acf(Yt, lag.max = 25, main= "Estimated ACF")

pacf(Yt, lag.max = 25, main= "Estimated PACF")

barplot(AR1.acf, col = "blue", main= "Theoretical ACF")

barplot(AR1.pacf, col = "red", main= "Theoretical PACF")
```

Execute the code 3 times to make sure the results are consistent.

- Repeat the previous exercise using the value $\phi = -0.7$.
- Investigate $\phi = 0.7$ and then $\phi = -0.7$ when n = 101.
- Investigate $\phi = 0.99$ and n = 1001.
- Investigate $\phi = -0.99$ and n = 1001.

2 AR(2) Process

The following R command simulates the AR(2) process $Y_t = 0.7Y_{t-1} - 0.4Y_{t-2} + e_t$ with $\sigma_e^2 = 1$. Plot the data and investigate the shape of the ACF and PACF. Repeat three times.

```
[]: Yt = arima.sim(list(ar = c(0.7, -0.4)), n=1001)
```

[]:

Repeat the previous exercise using a shorter time series. Try the AR(2) model with $\phi_1 = 0.7$ and $\phi_2 = 0.4$.

What went wrong?

[]:

3 Stationarity of the AR(2) Process

The AR(2) model $(1 - aB)(1 - bB)Y_t = e_t$ expands to

$$Y_t = \overbrace{(a+b)}^{\phi_1} Y_{t-1} + \overbrace{(-ab)}^{\phi_2} Y_{t-2} + e_t.$$

For this model to be stationary we require that the roots (both real and complex) of the generating polynomial $\phi(B) = (1-aB)(1-bB) = 0$ lie outside the unit circle. That is, we require |a| < 1 and |b| < 1. Also, as there are only two roots, if complex roots arise then $b = \bar{a}$. We can investigate this algebraically elsewhere. Here can use computer simulation to map and colour the admissible regions in the plane (ϕ_1, ϕ_2) .

Specifying admissible real roots is straightforward. Specifying admissible complex roots we need to write a = x + iy and $b = \bar{a} = x - iy$, so that $\phi_1 = 2x$ and $\phi_2 = -(x^2 + y^2)$. The following R code will do this:

```
[]: n = 5000
a = runif(n, min= -1, max = 1)
b = runif(n, min= -1, max = 1)
Phi.RootsReal = cbind( a+b , -a*b )
```

```
[]: x = runif(n, min= -1, max = 1)
y = numeric()
for(i in 1:n){
    y[i] = runif(1, -1*sqrt(1-(x[i])^2), sqrt(1-(x[i])^2))
}

Phi.RootsComplex = cbind( 2*x , -(x^2 +y^2) )
```

```
curve(-x^2/4, add = TRUE, lwd = 3) # condition (phi1)^2 + 4phi2 < 0</pre>
```

```
[]: CheckComplexRoots = (Phi.RootsComplex[,1])^2 + 4* Phi.RootsComplex[,2] all(CheckComplexRoots < 0)
```

4 ARMA(p,q) Process

Investigate the simulated models in the following R code:

5 Yule-Walker Equations

Consider again the time series 1h measuring the Luteinizing hormone level in blood samples at 10 mins intervals from a human female, 48 samples. The R command ar(1h) automatically selects and fits an AR(3) model to these data. Try it. The default fitting algorithm of ar is based on the Yule-Walker equations. The theoretical Yule-Walker equations for an AR(3) model can be written

$$\begin{bmatrix} \gamma_1 \\ \gamma_2 \\ \gamma_3 \end{bmatrix} = \begin{bmatrix} \gamma_0 & \gamma_1 & \gamma_2 \\ \gamma_1 & \gamma_0 & \gamma_1 \\ \gamma_2 & \gamma_1 & \gamma_0 \end{bmatrix} \begin{bmatrix} \phi_1 \\ \phi_2 \\ \phi_3 \end{bmatrix}$$

with

$$\gamma_0 = \sum_{j=1}^3 \phi_j \gamma_j$$

If we replace the theoretical covariances with their sample-statistics counterparts and solve, we should obtain estimate $\hat{\phi}_1$, $\hat{\phi}_2$, and $\hat{\phi}_3$ for the AR(3) model.

The following R code does this manually to show how the magic-box answers from ar(1h) are obtained.

```
[]: ar(lh)
```

```
[]: lh.acf = acf(lh, lag.max=3)
    lh.acf
[]: names(lh.acf)
[]: gamma.vec = matrix(data = lh.acf$acf[-1], ncol=1)

    Gamma.mtx = diag(3)
    diag(Gamma.mtx) = lh.acf$acf[1]
    Gamma.mtx[abs(row(Gamma.mtx)-col(Gamma.mtx)) == 1] = lh.acf$acf[2]
    Gamma.mtx[abs(row(Gamma.mtx)-col(Gamma.mtx)) == 2] = lh.acf$acf[3]
[]: gamma.vec
[]: Gamma.mtx
[]: Gamma.mtx
[]: (sunspot.ar <- ar(sunspot.year))
    sunspot.lead = predict(sunspot.ar, n.ahead = 25)

    ts.plot(sunspot.year, sunspot.lead$pred, col=c(5,2))</pre>
```

$6 \quad ARIMA(p,d,q) \text{ Models}$

The R code below simulates the time series model:

$$\begin{array}{rcl} (1-1.7B+0.7B^2)Y_t & = & e_t, \\ \\ (1-0.7B)(1-B)Y_t & = & e_t, \\ \\ (1-B)Y_t & = & 0.7(1-B)Y_{t-1}+e_t, \\ \\ (Y_t-Y_{t-1}) & = & 0.7(Y_{t-1}-Y_{t-2})+e_t, \\ \\ Y_t & = & 1.7Y_{t-1}-0.7Y_{t-2}+e_t \end{array}$$

Note that the coefficients of Y on the LHS sum 1.7 - 0.7 = 1. This is a quick check to spot that B = 1 is a root of the AR characteristic polynomial $\phi(B) = 0$.

```
[]: ts.sim = arima.sim(list(order = c(1,1,0), ar = 0.7), n = 500)
d = diff(ts.sim)
ts.plot(ts.sim)
```

```
[]: ts.plot(d)
```

```
[]: par(mfrow=c(2,2))
    acf(ts.sim)
    pacf(ts.sim)
    acf(d)
    pacf(d)
```

Note: The easiest way to get the roots of the polynomial (1-1.7B+ 0.7B^2)=0 is the R command

```
[]: polyroot(z = c(0.7,-1.7,1))
```

7 Simulating Time Series in R

Use arima.sim() to simulate from the model

$$Y_t = 2.7Y_{t-1} - 2.4Y_{t-2} + 0.7Y_{t-3} + e_t - 0.5e_{t-1}.$$

You should produce time series plots, ACF plots, PACF plots as necessary. It is a good idea to use the command set.seed() so that your simulation results can be reproduced. Use your own student ID number as the seed number. Compare the ACF (and PACF) plots to their theoretical counterparts.

Upload a "boilerplate" plot of the time series, the ACF and the PACF.