CSSS 512: Lab 3

 ${\sf Modeling\ Stationary\ Time\ Series}$

2018-4-20

Agenda

- 1. Homework 2
- 2. Box-Jenkins method
- 3. Estimation and interpretation of ARMA models
- 4. Cross-validation and model selection
- 5. Counterfactual forecasting

Homework 2

What to know:

- 1. ACF, PACF, plotting, demeaning
- 2. Unit root tests
- 3. Estimation and interpretation of ARMA models
- 4. Cross-validation and model selection
- 5. Counterfactual forecasting

Box-Jenkins Method

Steps:

- 1. Study generic forms and properties
- 2. Study these realizations for an indication of which possibly applies to your data
- 3. Assess your guess-diagnose and iterate
- 4. Perform a meta-analysis at the end to determine which specification is best

The Box-Jenkins method assumes that time series are composed by multiple temporal processes. It then performs diagnostics to compare the observed series with generic forms to decide what processes occur in the data

```
rm(list=ls())
# Load Libraries
library(forecast) # For auto.arima and cross-validation
library(tseries) # For unit root tests
library(lmtest) # For Breusch-Godfrey LM test of serial correlation
library(RColorBrewer) # For nice colors
library(MASS)
library(simcf)
# ARIMA Cross-validation by rolling windows
# Adapted from Rob J Hyndman's code:
# http://robihundman.com/hundsight/tscvexample/
# Could use further generalization, e.g. to seasonality
# Careful! This can produce singularities using categorical covariates
arimaCV <- function(x, order, xreg, include, mean, forward=1, minper=50) {
 require(forecast)
 if (!anv(class(x)=="ts")) x \leftarrow ts(x)
 n <- length(x)
 mae <- matrix(NA, nrow=n-minper, ncol=forward)
 st \leftarrow tsp(x)[1]+(minper-2)
 for(i in 1:(n-minper)) {
    xshort <- window(x, start=st+(i-minper+1), end=st+i)</pre>
    xnext <- window(x, start=st+(i+1), end=min(n, st+(i+forward)))</pre>
    xregshort <- window(xreg, start=st+(i-minper+1), end=st+i)</pre>
    xregnext <- window(xreg, start=st+(i+1), end=min(n, st+(i+forward)))</pre>
    fit <- Arima(xshort, order=order, xreg=xregshort, include.mean=include.mean)
    fcast <- forecast(fit, h=length(xnext), xreg=xregnext)</pre>
    mae[i,1:length(xnext)] <- abs(fcast[['mean']]-xnext)
  colMeans(mae, na.rm=TRUE)
```

[1] "death" "law" "vear" "mt"

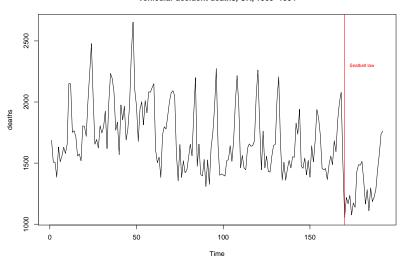
```
# Load data
# Number of deaths and serious injuries in UK road accidents each month.
# Jan 1969 - Dec 1984. Seatbelt law introduced in Feb 1983
# (indicator in second column). Source: Harvey, 1989, p.519ff.
# http://www.staff.city.ac.uk/~sc397/courses/3ts/datasets.html
#
# Variable names: death law
ukdata <- read.csv("ukdeaths.csv",header=TRUE)
attach(ukdata)
colnames(ukdata)</pre>
```

```
# Look at the time series
```

"month"

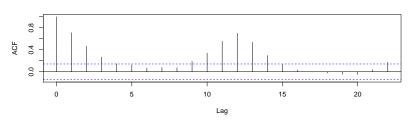
```
plot(death,type="1",ylab="deaths",xlab="Time", main = "Vehicular accident deaths, UK, 1969-1984")
lines(x=c(170,170),y=c(0,5000),col="red"); text("Seatbelt law",x = 180, y = 2300, col="red",cex=0.7)
```

Vehicular accident deaths, UK, 1969-1984

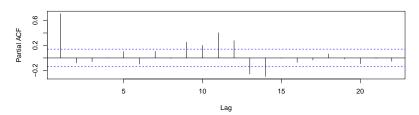


Look at the ACF and PACF
par(mfrow=c(2,1))
acf(death); pacf(death)

Series death

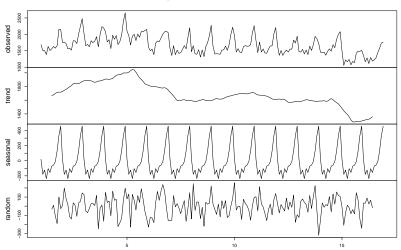


Series death



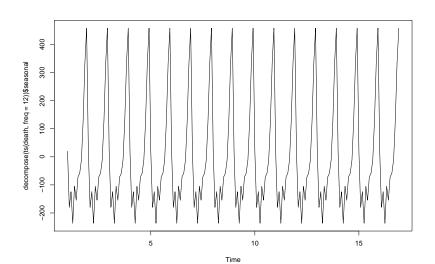
Look at the decomposed time series
plot(decompose(ts(death,freq=12)))

Decomposition of additive time series



Time

Look at the monthly cycle
plot(decompose(ts(death,freq=12))\$seasonal)



```
#Look at the monthly cycle by plotting each year separately

# Make some colors
col <- brewer.pal(8, "Blues")

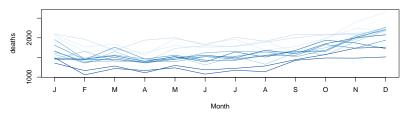
# Gather the data (sort the number of deaths by month and year in a matrix)
deathmat <- matrix(death,nrow=12,ncol=length(death)/12, byrow=FALSE)

# Repeat them as many times as needed
col <- as.vector(t(matrix(col, nrow=length(col),
ncol=ceiling(ncol(deathmat)/length(col)))))

# Plot each year over the months</pre>
```

```
par(mfrow=c(2,1))
matplot(deathmat, type="1", col=col, lty=1, xaxt="n", ylab="deaths", xlab="Month",
    main=expression(paste("Monthly view of accident deaths, UK, 1969-1984")))
axis(1, at=1:12, labels=c("J", "F", "M", "A", "M", "J", "J", "A", "S", "O", "N", "D"))
abline(a=0,b=0,lty="dashed")
```

Monthly view of accident deaths, UK, 1969-1984



Unit Root Tests

Intuition: if time series is stationary, then regressing $y_t - y_{t-1}$ on y_{t-1} should produce a negative coefficient. Why?

In a stationary series, knowing the past value of the series helps to predict the next period's change. Positive shifts should be followed by negative shifts (mean reversion).

$$\begin{aligned} y_t &= \rho y_{t-1} + \epsilon_t \\ y_t - y_{t-1} &= \rho y_{t-1} - y_{t-1} + \epsilon_t \\ \Delta y_t &= \gamma y_{t-1} + \epsilon_t \text{, where } \gamma = (\rho - 1) \end{aligned}$$

Augmented Dickey-Fuller test: null hypothesis of unit root.

Same with Phillips-Perron test, but differs in how the AR(p) time series is modeled: lags, serial correlation, heteroskedasticity.

Unit Root Tests

```
# Check for a unit root
PP.test(death)
##
## Phillips-Perron Unit Root Test
##
## data: death
## Dickey-Fuller = -6.4349, Truncation lag parameter = 4, p-value =
## 0.01
adf.test(death)
## Warning in adf.test(death): p-value smaller than printed p-value
##
## Augmented Dickey-Fuller Test
##
## data: death
## Dickey-Fuller = -6.5366, Lag order = 5, p-value = 0.01
## alternative hypothesis: stationary
```

#Do we have evidence that the time series stationary?

```
#It looks like there is seasonality in the time series, so let's try to control for each month or Q4
# Make some month variables (there are easier ways!)
jan <- as.numeric(month=="January")</pre>
feb <- as.numeric(month=="February")</pre>
mar <- as.numeric(month=="March")
apr <- as.numeric(month=="April")
may <- as.numeric(month=="May")
jun <- as.numeric(month=="June")</pre>
jul <- as.numeric(month=="Julv")</pre>
aug <- as.numeric(month=="August")</pre>
sep <- as.numeric(month=="September")</pre>
oct <- as.numeric(month=="October")
nov <- as.numeric(month=="November")
dec <- as.numeric(month=="December")
# Make a fourth quarter indicator
q4 <- as.numeric(oct nov dec)
```

```
# Store all these variables in the dataframe
ukdata$jan <- jan
ukdata$feb <- feb
ukdata$mar <- mar
ukdata$apr <- apr
ukdata$may <- may
ukdata$iun <- iun
ukdata$jul <- jul
ukdata$aug <- aug
ukdata$sep <- sep
ukdata$oct <- oct
ukdata$nov <- nov
ukdata$dec <- dec
ukdata$q4 <- q4
# Set rolling window length and look ahead period for cross-validation
minper <- 170
forward <- 12
```

Recall that we use ML to estimate the parameters of an ARMA model (although we also discussed using LS).

This should be familiar:

- 1. Express the joint probability of the data using the chosen probability distribution
- 2. Convert the joint probability to the likelihood
- 3. Simplify the likelihood for easy maximization
- 4. Substitute the systematic component

For an AR(1) process, we are left with the log likelihood function:

$$\mathcal{L}(\boldsymbol{\beta}, \phi_1 | \mathbf{y}, \mathbf{X}) = -\frac{1}{2} \log \left(\frac{\sigma^2}{1 - \phi_1^2} \right) - \frac{\left(y_1 - \frac{\mathbf{x_1} \boldsymbol{\beta}}{1 - \phi_1} \right)^2}{\frac{2\sigma^2}{1 - \phi_1^2}} - \frac{T - 1}{2} \log \sigma^2 - \sum_{T=2}^{T} \frac{(y_t - \mathbf{x_t} \boldsymbol{\beta} - \phi_1 y_{t-1})^2}{2\sigma^2}$$

We can estimate the parameters, β and ϕ_1 that make the data most likely using numerical methods.

The arima function in R will compute these for us.

```
## Model 1a: AR(1) model of death as function of law
##
## Estimate an AR(1) using arima
xcovariates <- law
arima.res1a \leftarrow arima(death, order = c(1.0.0).
                     xreg = xcovariates, include.mean = TRUE
print(arima.res1a)
##
## Call:
## arima(x = death, order = c(1, 0, 0), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
            ar1 intercept xcovariates
         0.6439 1719.193 -377.4542
##
## s.e. 0.0553 42.078 107.6520
##
## sigma^2 estimated as 39289: log likelihood = -1288.26, aic = 2584.52
#How do we interpret these parameter estimates?
# Extract estimation results from arima.res1a
pe.1a <- arima.res1a$coef
                                             # parameter estimates (betas)
se.1a <- sqrt(diag(arima.res1a$var.coef)) # standard errors
ll.1a <- arima.res1a$loglik
                                             # log likelihood at its maximum
sigma2hat.1a <- arima.res1a$sigma2
                                             # standard error of the regression
aic.1a <- arima.res1a$aic
                                             # Akaike Information Criterion
resid 1a <- arima res1a$resid
                                             # residuals
```

How do we interpret these parameter estimates?

Introducing the seatbelt law is associated on average with a 378 decrease in the number of road accidents in the next period, all else constant.

```
#Recall that the AIC is equal to the deviance (-2*log likelihood at its maximum) of the model #plus 2 * the dimension of the model (number of free parameters of the model) -2*ll.1a + 2*length(pe.1a)
```

[1] 2582.52

#And the standard error of the regression is just the expected value of the squared residuals $mean(resid.1a^2)$

[1] 39289.43

#With a fixed mean, where does y_t converge?

We know in an AR(1) process the effect accumulates over time with a fixed mean.

$$\mathbb{E}(y_t) = \frac{x_t \beta}{1 - \phi_1}$$

We expect the mean level of accidents to converge to

$$\mathbb{E}(y_t) = \frac{x_t \beta}{1 - \phi_1} = \frac{1719.193 - 377.454}{1 - 0.644} = 3768$$

$$\mathbb{E}(y_t) = \frac{x_t \beta}{1 - \phi_1} = \frac{1719.193}{1 - 0.644} = 4828$$

In general, we will forecast these expected values and also predicted values using simulation.

Cross Validation and Model Selection

```
## [1] 119.6679 136.2173 175.0493 182.9675 185.3571 187.4022 198.2450
## [8] 188.4625 183.4294 165.0588 164.3636 161.9931
```

Recall that we use a rolling forecast window to perform cross validation.

Describe the steps in your own words.

```
## Model 1b: AR(1) model of death as function of law & q4
## Estimate an AR(1) using arima
xcovariates <- cbind(law, q4)
arima.res1b <- arima(death, order = c(1,0,0),
                    xreg = xcovariates, include.mean = TRUE
print(arima.res1b)
##
## Call:
## arima(x = death, order = c(1, 0, 0), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
                                 law
##
            ar1 intercept
##
        0.5352 1638.0301 -395.6701 324.5653
## s.e. 0.0636 28.1199 72.3030 34.5033
##
## sigma^2 estimated as 26669: log likelihood = -1250.97, aic = 2511.93
# Extract estimation results from arima.res1b
pe.1b <- arima.res1b$coef
                                            # parameter estimates (betas)
se.1b <- sqrt(diag(arima.res1b$var.coef))
                                            # standard errors
ll.1b <- arima.res1b$loglik
                                            # log likelihood at its maximum
sigma2hat.1b <- arima.res1b$sigma2
                                            # standard error of the regression
aic.1b <- arima.res1b$aic
                                            # Akaike Information Criterion
                                            # residuals
resid.1b <- arima.res1b$resid
```

Cross Validation and Model Selection

```
## [1] 108.51792 81.04685 99.33850 83.35743 100.56609 95.20655 110.10528
## [8] 91.06577 104.57805 106.20068 123.39408 115.11284
```

resid_1c <- arima_res1c@resid_

```
## Model 1c: AR(1) model of death as function of law & months
## Estimate an AR(1) using arima
xcovariates <- cbind(law, jan, feb, mar, apr, may, jun, aug, sep, oct, nov, dec)
arima.res1c <- arima(death, order = c(1.0.0).
                   xreg = xcovariates, include.mean = TRUE
print(arima.res1c)
##
## Call:
## arima(x = death, order = c(1, 0, 0), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
##
           ar1 intercept law jan feb mar
                                                                    apr
   0.6442 1638.6270 -370.0694 81.3021 -95.1350 -44.3298 -157.3445
## s.e. 0.0550 42.9093 70.2727 54.8127 54.5036 53.0792
                                                                50.2149
                     iun
                              aug
                                      sep
                                               oct
                                                                  dec
##
            mav
                                                         nov
##
      -19.9428 -75.6674 14.7670 67.4890 206.6686 405.9134 522.0696
## s.e. 45.0247 35.1890 35.1882 45.0184 50.1913 53.0074 54.3054
##
## sigma^2 estimated as 16333: log likelihood = -1204. aic = 2437.99
# Extract estimation results from arima.res1c
pe.1c <- arima.res1c$coef
                                         # parameter estimates (betas)
se.1c <- sqrt(diag(arima.res1c$var.coef)) # standard errors
ll.1c <- arima.res1c$loglik
                                         # log likelihood at its maximum
sigma2hat.1c <- arima.res1c$sigma2 # standard error of the regression
aic.1c <- arima.res1c$aic
                                         # Akaike Information Criterion
```

residuals

Cross Validation and Model Selection

```
## [1] 83.90928 101.80611 98.30016 92.09344 82.96249 80.83424 77.65451
## [8] 79.87631 95.52618 77.39619 50.74492 43.74215
```

resid_1d_<- arima_res1d\$resid_

```
## Model 1d: AR(1) model of death as function of law & select months
## Estimate an AR(1) using arima
xcovariates <- cbind(law, jan, sep, oct, nov, dec)
arima.res1d <- arima(death, order = c(1.0.0).
                   xreg = xcovariates, include.mean = TRUE
print(arima.res1d)
##
## Call:
## arima(x = death, order = c(1, 0, 0), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
           ar1 intercept law
                                         ian
                                                sep oct
                                                                    nov
   0.6045 1589.4405 -377.7457 154.7288 80.7422 238.3880 451.3567
## s.e. 0.0575
                 29.4161 69.7719 35.7336 35.8534 42.6836 44.3474
             dec
##
##
      579 9770
## s.e. 42.6108
##
## sigma^2 estimated as 18989: log likelihood = -1218.42. aic = 2454.83
# Extract estimation results from arima.res1d
pe.1d <- arima.res1d$coef
                                          # parameter estimates (betas)
se.1d <- sqrt(diag(arima.res1d$var.coef)) # standard errors
                                          # log likelihood at its maximum
ll.1d <- arima.res1d$loglik
sigma2hat.1d <- arima.res1d$sigma2
                                         # standard error of the regression
aic.1d <- arima.res1d$aic
                                          # Akaike Information Criterion
```

residuals

Cross Validation and Model Selection

```
## [1] 99.18077 92.74232 107.56732 95.79555 92.29416 93.66586 102.52082 ## [8] 99.48355 109.96694 90.24875 74.12182 65.34772
```

```
## Model 1e: AR(1)AR(1)_12 model of death as function of law
##
## Estimate an AR(1)AR(1)_12 using arima
xcovariates <- cbind(law)
arima.res1e \leftarrow arima(death, order = c(1,0,0).
                     seasonal = list(order = c(1,0,0), period = 12),
                     xreg = xcovariates, include.mean = TRUE
print(arima.res1e)
##
## Call:
## arima(x = death, order = c(1, 0, 0), seasonal = list(order = c(1, 0, 0), period = 12),
##
       xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
            ar1 sar1 intercept
##
                                         ไลน
        0.4446 0.6511 1710.1531 -347.6812
##
## s.e. 0.0695 0.0564 53.3648
                                      73.0634
##
## sigma^2 estimated as 23693: log likelihood = -1242.86. aic = 2495.71
# Extract estimation results from arima.res1e
pe.1e <- arima.res1e$coef
                                             # parameter estimates (betas)
se.1e <- sqrt(diag(arima.res1e$var.coef)) # standard errors
ll.1e <- arima.res1e$loglik
                                             # log likelihood at its maximum
sigma2hat.1e <- arima.res1e$sigma2
                                             # standard error of the regression
aic.1e <- arima.res1e$aic
                                             # Akaike Information Criterion
resid le <- arima reslegresid
                                             # residuals
```

Cross Validation and Model Selection

Attempt at rolling window cross-validation (see caveats)
cv.1e <- arimaCV(ts(death), order=c(1,0,0), forward=forward,</pre>

```
## [1] 119.6679 136.2173 175.0493 182.9675 185.3571 187.4022 198.2450
## [8] 188.4625 183.4294 165.0588 164.3636 161.9931

# So far, an AR(1) with additive seasonality looks best according to AIC
# But maybe a different ARMA(p,q) would fit better?
# Let's keep the additive seasonality and try various ARMA models manually
```

xreg=xcovariates, include.mean=TRUE, minper=minper)

```
## Model 2a: AR(2) model of death as function of law & months
## Estimate an AR(2) using arima
xcovariates <- cbind(law. jan. feb. mar. apr. mav. jun. aug. sep. oct. nov. dec)
arima.res2a \leftarrow arima(death, order = c(2.0.0).
                   xreg = xcovariates, include.mean = TRUE
print(arima.res2a)
##
## Call:
## arima(x = death, order = c(2, 0, 0), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
##
                  ar2 intercept law jan feb
           ar1
                                                                 mar
## 0.4696 0.2711 1635.0869 -347.9213 83.7469 -94.9882 -44.0442
## s.e. 0.0692 0.0694 45.6076
                                   80.5683 46.9299 46.5145 45.0452
##
              apr
                      may jun
                                        aug
                                                sep
                                                          oct
                                                                   nov
## -157.2316 -19.8376 -75.5957 14.8059 67.5047 206.7362 406.0569
## s.e. 42.8448 37.9719 35.0631 35.0623 37.9640 42.8242 44.9760
            dec
##
## 522,4596
## s.e. 46.4368
##
## sigma^2 estimated as 15118: log likelihood = -1196.65, aic = 2425.3
# Extract estimation results from arima.res2a
pe.2a <- arima.res2a$coef
                                         # parameter estimates (betas)
se.2a <- sqrt(diag(arima.res2a$var.coef)) # standard errors
11.2a <- arima.res2a$loglik
                                         # log likelihood at its maximum
sigma2hat.2a <- arima.res2a$sigma2
                                        # standard error of the regression
aic.2a <- arima.res2a$aic
                                         # Akaike Information Criterion
resid 2a <- arima res2a$resid
                                         # residuals
```

Cross Validation and Model Selection

```
## [1] 92.57935 109.47089 105.70520 91.33796 78.80066 78.69454 76.70600
## [8] 82.33373 94.98572 78.40537 50.54681 37.00711
```

```
## Model 2b: MA(1) model of death as function of law & months
## Estimate an MA(1) using arima
xcovariates <- cbind(law, jan, feb, mar, apr, may, jun, aug, sep, oct, nov, dec)
arima.res2b \leftarrow arima(death, order = c(0.0.1).
                   xreg = xcovariates, include.mean = TRUE
print(arima.res2b)
##
## Call:
## arima(x = death, order = c(0, 0, 1), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
##
          ma1 intercept law jan feb mar
                                                                     apr
        0.4539 1641.4834 -391.7280 79.9732 -94.6320 -44.0097 -157.2155
## s.e. 0.0538 39.7814 45.5288 55.5797 55.6807 55.6807 55.6807
                     iun
                              aug
                                       sep
                                                oct
                                                                   dec
##
             mav
                                                         nov
##
      -19.8754 -75.6604 14.8400 67.6897 207.0297 406.5988 522.4457
## s.e. 55.6807 43.9719 43.9719 55.6807 55.6807 55.6807 55.5411
##
## sigma^2 estimated as 20566: log likelihood = -1225.97, aic = 2481.93
# Extract estimation results from arima.res2b
pe.2b <- arima.res2b$coef
                                          # parameter estimates (betas)
se.2b <- sqrt(diag(arima.res2b$var.coef)) # standard errors
11.2b <- arima.res2b$loglik
                                          # log likelihood at its maximum
sigma2hat.2b <- arima.res2b$sigma2 # standard error of the regression
aic.2b <- arima.res2b$aic
                                          # Akaike Information Criterion
resid 2b <- arima res2b$resid
                                          # residuals
```

Cross Validation and Model Selection

```
## [1] 79.86555 93.67862 91.43308 90.51553 87.20853 90.10537 92.36365
## [8] 87.97704 89.02266 82.18906 58.03778 61.96120
```

```
## Model 2c: ARMA(1.1) model of death as function of law & months
## Estimate an ARMA(1,1) using arima
xcovariates <- cbind(law. jan. feb. mar. apr. mav. jun. aug. sep. oct. nov. dec)
arima.res2c \leftarrow arima(death, order = c(1.0.1).
                   xreg = xcovariates, include.mean = TRUE
print(arima.res2c)
##
## Call:
## arima(x = death, order = c(1, 0, 1), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
##
           ar1 ma1 intercept law jan
                                                        feb
                                                                 mar
## 0.9349 -0.5994 1629.5549 -323.4929 85.7471 -94.0923 -43.6000
## s.e. 0.0383 0.1076 58.6795 83.2081 40.4544 40.2349 39.7247
##
              apr
                      may jun
                                        aug
                                                 sep
                                                          oct
                                                                   nov
## -156.8606 -19.6467 -75.5028 14.7339 67.3872 206.5916 405.9572
## s.e. 38.8954 37.7225 36.1673 36.1671 37.7207 38.8896 39.7111
            dec
##
## 522,3735
## s.e. 40.2083
##
## sigma^2 estimated as 14568: log likelihood = -1193.18. aic = 2418.37
# Extract estimation results from arima.res2c
pe.2c <- arima.res2c$coef
                                         # parameter estimates (betas)
se.2c <- sqrt(diag(arima.res2c$var.coef)) # standard errors
11.2c <- arima.res2c$loglik
                                         # log likelihood at its maximum
sigma2hat.2c <- arima.res2c$sigma2
                                        # standard error of the regression
aic.2c <- arima.res2c$aic
                                         # Akaike Information Criterion
resid 2c <- arima res2c$resid
                                         # residuals
```

```
## [1] 89.53537 104.43751 95.71992 79.82402 72.93142 79.67158 88.12675
## [8] 87.42871 88.24699 71.12715 47.74176 48.65514
```

```
## Model 2d: ARMA(2.1) model of death as function of law & months
## Estimate an ARMA(2,1) using arima
xcovariates <- cbind(law. jan. feb. mar. apr. mav. jun. aug. sep. oct. nov. dec)
arima.res2d \leftarrow arima(death, order = c(2.0.1).
                   xreg = xcovariates, include.mean = TRUE
print(arima.res2d)
##
## Call:
## arima(x = death, order = c(2, 0, 1), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
##
                   ar2 ma1 intercept law
           ar1
                                                        ian
                                                                 feb
## 1.1899 -0.2157 -0.7950 1626.1862 -321.2201 84.8843 -94.5311
## s.e. 0.1071 0.0976 0.0724 68.6982 78.8301 41.3869 41.3010
##
             mar
                       apr
                                mav iun
                                                  aug
                                                          sep
                                                                    oct
## -43.8782 -157.0544 -19.7871 -75.5646 14.8208 67.5749 206.8634
## s.e. 41.0435
                40.5352 39.3222 35.1484 35.1483 39.3216 40.5327
                      dec
##
             nov
## 406.3691 522.9159
## s.e. 41.0341 41.2487
##
## sigma^2 estimated as 14284: log likelihood = -1191.33, aic = 2416.66
# Extract estimation results from arima.res2d
pe.2d <- arima.res2d$coef
                                          # parameter estimates (betas)
se.2d <- sqrt(diag(arima.res2d$var.coef)) # standard errors
11.2d <- arima.res2d$loglik
                                         # log likelihood at its maximum
sigma2hat.2d <- arima.res2d$sigma2
                                         # standard error of the regression
aic.2d <- arima.res2d$aic
                                         # Akaike Information Criterion
resid 2d <- arima res2d$resid
                                          # residuals
```

```
## [1] 83.46358 99.21901 91.91804 81.74041 77.82364 83.17353 91.88016
## [8] 86.66464 85.28058 72.42659 53.34436 56.62732
```

```
## Model 2e: ARMA(1.2) model of death as function of law & months
## Estimate an ARMA(1,2) using arima
xcovariates <- cbind(law. jan. feb. mar. apr. mav. jun. aug. sep. oct. nov. dec)
arima.res2e \leftarrow arima(death, order = c(1.0.2).
                   xreg = xcovariates, include.mean = TRUE
print(arima.res2e)
##
## Call:
## arima(x = death, order = c(1, 0, 2), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
##
           ar1 ma1 ma2 intercept law
                                                        ian
                                                                 feh
## 0.9620 -0.5892 -0.1228 1627.146 -322.6854 85.1562 -94.1511
## s.e. 0.0253 0.0752 0.0705 66.814 79.2449 40.7504 40.6400
##
            mar
                       apr
                                mav jun
                                                  aug
                                                          sep
                                                                   oct
## -43,6591 -156,9126 -19,6915 -75,5237 14,7645 67,4691 206,7084
## s.e. 40.3701 39.9498 39.3736 35.5453 35.5453 39.3730 39.9476
                      dec
##
            nov
## 406,1477 522,6613
## s.e. 40.3650 40.5994
##
## sigma^2 estimated as 14356: log likelihood = -1191.82, aic = 2417.63
# Extract estimation results from arima.res2e
pe.2e <- arima.res2e$coef
                                         # parameter estimates (betas)
se.2e <- sqrt(diag(arima.res2e$var.coef)) # standard errors
11.2e <- arima.res2e$loglik
                                         # log likelihood at its maximum
sigma2hat.2e <- arima.res2e$sigma2
                                         # standard error of the regression
aic.2e <- arima.res2e$aic
                                         # Akaike Information Criterion
resid 2e <- arima res2e$resid
                                         # residuals
```

```
## [1] 84.82861 99.53238 91.96193 80.66522 76.50626 83.03813 92.32656
## [8] 87.44648 85.38180 71.74271 52.74060 55.51375
```

```
## Model 2f: ARMA(2.2) model of death as function of law & months
## Estimate an ARMA(2,2) using arima
xcovariates <- cbind(law. jan. feb. mar. apr. mav. jun. aug. sep. oct. nov. dec)
arima.res2f \leftarrow arima(death, order = c(2.0.2).
                   xreg = xcovariates, include.mean = TRUE
print(arima.res2f)
##
## Call:
## arima(x = death, order = c(2, 0, 2), xreg = xcovariates, include.mean = TRUE)
##
## Coefficients:
##
                  ar2 ma1 ma2 intercept law
           ar1
                                                              ian
## 0.0526 0.8449 0.3497 -0.6503 1625.7793 -312.2308 86.0931
## s.e. 0.0538 0.0413 0.1006 0.0998 61.5565
                                                  81.8335 40.9421
##
            feb
                     mar
                                apr
                                         may
                                                  iun
                                                           aug
                                                                   sep
## -91.7482 -43.7677 -154.3960 -19.6984 -72.8430 17.6629 67.3856
## s.e. 38.1258 40.4084 36.9053 38.9443 34.4385 34.4299 38.9431
                               dec
##
            oct.
                     nov
## 209.8757 405.8869 526.1152
## s.e. 36.8765 40.3991 38.0647
##
## sigma^2 estimated as 13794: log likelihood = -1189.2, aic = 2414.39
# Extract estimation results from arima.res2f
pe.2f <- arima.res2f$coef
                                         # parameter estimates (betas)
se.2f <- sqrt(diag(arima.res2f$var.coef)) # standard errors
11.2f <- arima.res2f$loglik
                                         # log likelihood at its maximum
sigma2hat.2f <- arima.res2f$sigma2
                                        # standard error of the regression
aic.2f <- arima.res2f$aic
                                         # Akaike Information Criterion
resid 2f <- arima res2f$resid
                                         # residuals
```

```
## [1] 85.54471 102.70679 92.51201 81.72152 71.43657 81.67783 84.97804 ## [8] 87.64424 89.59192 70.33841 39.72134 48.94544
```

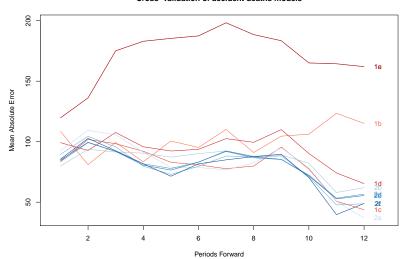
```
## Series: death
## Regression with ARIMA(2,0,1) errors
##
## Coefficients:
##
           ar1
                    ar2
                            ma1 intercept
                                                 law
                                                          jan
                                                                    feb
        1.1899 -0.2157 -0.7950 1626.1862 -321.2201 84.8843 -94.5311
##
## s.e. 0.1071
                 0.0976 0.0724
                                   68 6982
                                              78 8301 41 3869
                                                                41 3010
##
                                           jun
             mar
                        apr
                                 may
                                                   aug
                                                            sep
                                                                      oct
        -43.8782
                 -157.0544 -19.7871 -75.5646 14.8208 67.5749 206.8634
##
         41 0435
                    40.5352
                             39.3222 35.1484 35.1483 39.3216 40.5327
## s.e.
##
                      dec
             nov
##
        406.3691 522.9159
## s.e. 41 0341 41 2487
##
## sigma^2 estimated as 15582: log likelihood=-1191.33
## ATC=2416.66 ATCc=2420.18 BTC=2472.04
```

```
# Extract estimation results from arima.res3a
pe.3a <- arima.res3a$coef
                                             # parameter estimates (betas)
se.3a <- sqrt(diag(arima.res3a$var.coef))
                                          # standard errors
11.3a <- arima.res3a$loglik
                                             # log likelihood at its maximum
sigma2hat.3a <- arima.res3a$sigma2
                                             # standard error of the regression
aic.3a <- arima.res3a$aic
                                             # Akaike Information Criterion
resid.3a <- arima.res3a$resid
                                             # residuals
# Attempt at rolling window cross-validation (see caveats)
cv.3a <- arimaCV(ts(death), order=c(2,0,1), forward=forward,
                 xreg=xcovariates, include.mean=TRUE, minper=minper)
cv.3a
```

```
## [1] 83.46358 99.21901 91.91804 81.74041 77.82364 83.17353 91.88016
## [8] 86.66464 85.28058 72.42659 53.34436 56.62732
```

```
matplot(allCV, type="1", col=col, lty=1, ylab="Mean Absolute Error", xlab="Periods Forward",
    main="Cross-validation of accident deaths models", xlim=c(0.75,12.75))
text(labs, x=rep(12.5,length(labs)), y=allCV[nrow(allCV),], col=col)
```

Cross-validation of accident deaths models



```
# Average cross-validation results
avgCV12 <- sort(apply(allCV, 2, mean))
avgCV8 <- sort(apply(allCV[1:8,], 2, mean))

# Based on AIC & cross-validation,
# let's select Model 2f to be our final model;
# there are other plausible models
arima.resF <- arima.resF <- arima.res2f
```

print(summary(lm.res1f))

```
##
## Call:
## lm(formula = death ~ lagdeath + jan + feb + mar + apr + may +
      jun + aug + sep + oct + nov + dec + law)
##
## Residuals:
##
      Min
               1Q Median
                                     Max
## -323.58 -84.45 -3.80 80.97 404.88
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 635.11393 96.64706 6.571 5.38e-10 ***
## lagdeath
                 0.64313 0.05787 11.114 < 2e-16 ***
              -302.58936 59.33982 -5.099 8.71e-07 ***
## jan
## feb
              -211.00947
                          48.46926 -4.353 2.26e-05 ***
## mar
             -31.82070
                          47.33602 -0.672 0.502314
            -177.52653
                          47.35870 -3.749 0.000241 ***
## apr
## may
                32.58040
                          47.55810 0.685 0.494199
## jun
              -111.47957
                          47.43316 -2.350 0.019863 *
## aug
              -33.76181 47.52523 -0.710 0.478393
                 9.48411
                          47.61220 0.199 0.842339
## sep
## oct
              114.89374
                          48.04444 2.391 0.017832 *
## nov
               224.81981
                          50.07068 4.490 1.28e-05 ***
               213.09991
                          54.93824 3.879 0.000148 ***
## dec
## law
              -145.31036
                          37.36477 -3.889 0.000142 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 133.9 on 177 degrees of freedom
    (1 observation deleted due to missingness)
## Multiple R-squared: 0.802, Adjusted R-squared: 0.7875
## F-statistic: 55.17 on 13 and 177 DF. p-value: < 2.2e-16
```

```
# Check LS result for serial correlation in the first or second order
bgtest(lm.res1f.1)
##
## Breusch-Godfrey test for serial correlation of order up to 1
##
## data: lm.res1f
## LM test = 11.546, df = 1, p-value = 0.000679
bgtest(lm.res1f,2)
##
   Breusch-Godfrey test for serial correlation of order up to 2
##
## data: lm.res1f
## LM test = 11.984, df = 2, p-value = 0.002498
# Evidence of residual serial correlation is strong
```

print(summary(lm.res1g))

```
##
## Call:
## lm(formula = death ~ lagdeath + lag2death + jan + feb + mar +
      apr + may + jun + aug + sep + oct + nov + dec + law)
##
## Residuals:
      Min
              1Q Median
                                    Max
## -378.22 -88.29 -5.04 89.71 308.44
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 475.12645 103.68324 4.582 8.71e-06 ***
## lagdeath
                0.47250
                          0.07332 6.445 1.09e-09 ***
                ## lag2death
## ian
             -311.45937 57.62112 -5.405 2.09e-07 ***
## feb
             -329.58156 57.96856 -5.686 5.37e-08 ***
            -68.08737 46.99905 -1.449 0.149212
## mar
## apr
             -152.44095
                         46.46031 -3.281 0.001248 **
## mav
               25.02334 46.18114 0.542 0.588610
              -65.76811 47.71466 -1.378 0.169851
## jun
              -6.16090 46.72852 -0.132 0.895259
## aug
## sep
              19.68658
                         46.27238 0.425 0.671032
## oct
              130.18618 46.79714 2.782 0.005997 **
              249.97112 49.06743 5.094 9.00e-07 ***
## nov
## dec
              235.55993 53.65766 4.390 1.96e-05 ***
             -111.47166
                        37.45979 -2.976 0.003336 **
## law
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 129.8 on 175 degrees of freedom
    (2 observations deleted due to missingness)
## Multiple R-squared: 0.8155, Adjusted R-squared: 0.8008
```

Describe in your own words the steps of the Breusch-Godfrey test.

```
# Check LS result for serial correlation in the first or second order
bgtest(lm.res1g.1)
##
## Breusch-Godfrey test for serial correlation of order up to 1
##
## data: lm.res1g
## LM test = 0.6961, df = 1, p-value = 0.4041
bgtest(lm.res1g,2)
##
   Breusch-Godfrev test for serial correlation of order up to 2
##
## data: lm.res1g
## LM test = 3.2256, df = 2, p-value = 0.1993
# Borderline evidence of serial correlation, but substantively different result.
# (Even small time series assumptions can have big implications for substance.)
# MA terms in ARMA(2,2) seems justified;
# we reject the LS model with lags of the DV
```

Recall that we covered two approaches: forecasting predicted values and forecasting expected values.

How are these different?

Describe the steps in your own words.

```
## Now that we've selected a model, let's interpret it
## using counterfactuals iterated over time
##
## Predict out five years (60 periods) assuming law is kept
# Make newdata dataframe for prediction
xcovariates <- cbind(law, jan, feb, mar, apr, may,
                     jun, aug, sep, oct, nov, dec)
n ahead <-60
lawhyp0 <- rep(1,n,ahead)
janhyp0 \leftarrow rep(c(1,0,0,0,0,0,0,0,0,0,0,0),5)
febhyp0 \leftarrow rep( c( 0,1,0, 0,0,0, 0,0,0, 0,0,0), 5)
marhvp0 \leftarrow rep(c(0.0.1, 0.0.0, 0.0.0, 0.0.0), 5)
aprhyp0 \leftarrow rep(c(0,0,0,1,0,0,0,0,0,0,0,0),5)
mayhyp0 \leftarrow rep( c( 0,0,0, 0,1,0, 0,0,0, 0,0,0), 5)
junhyp0 \leftarrow rep(c(0,0,0,0,0,1,0,0,0,0,0,0),5)
aughyp0 \leftarrow rep(c(0,0,0,0,0,0,0,1,0,0,0,0),5)
sephyp0 \leftarrow rep(c(0,0,0,0,0,0,0,1,0,0,0),5)
octhyp0 \leftarrow rep(c(0.0.0, 0.0.0, 0.0.0, 1.0.0), 5)
novhyp0 <- rep( c(0,0,0,0,0,0,0,0,0,0,0,1,0), 5)
dechyp0 \leftarrow rep(c(0,0,0,0,0,0,0,0,0,0,0,1), 5)
newdata0 <- cbind(lawhyp0, janhyp0, febhyp0, marhyp0,
                  aprhyp0, mayhyp0, junhyp0, aughyp0,
                  sephyp0, octhyp0, novhyp0, dechyp0)
# Must be in same order as model!
newdata0 <- as.data.frame(newdata0)
names(newdata0) <- c("law", "jan", "feb", "mar", "apr",
                     "may", "jun", "aug", "sep", "oct",
                     "nov", "dec")
```

```
# Run predict
vpred0 <- predict(arima.resF.
                  n.ahead = n.ahead,
                  newxreg = newdata0)
# Simulate predicted values
sims <- 10000
simparam <- myrnorm(sims, pe.2f, arima.res2f$var.coef)
xhvp <- newdata0
simphi <- simparam[,1:2]
simrho <- simparam[,3:4]
simbetas <- simparam[,5:ncol(simparam)]</pre>
lagY <- c(death[length(death)],death[length(death)-1])</pre>
lagY <- as.vector(lagY)</pre>
lagEps <- c(arima.res2f$resid[length(death)], arima.res2f$resid[length(death)-1])
lagEps <- as.vector(lagEps)
sigma <- sqrt(arima.res2f$sigma)
sim.ev2f <- ldvsimpv(xhyp,
                     simbetas,
                     ci=0.95,
                     constant=1.
                     phi=simphi,
                    lagY=lagY,
                     rho=simrho.
                     lagEps=lagEps,
                     sigma=sigma
```

```
# Predict out five years (60 periods) assuming law is repealed
# Make newdata dataframe for prediction
xcovariates <- cbind(law, jan, feb, mar, apr, may,
                      jun, aug, sep, oct, nov, dec)
n.ahead <-60
lawhyp <- rep(0,n.ahead)
ianhvp \leftarrow rep(c(1.0.0, 0.0.0, 0.0.0, 0.0.0), 5)
febhyp \leftarrow rep( c( 0,1,0, 0,0,0, 0,0,0, 0,0,0 ), 5)
marhyp \leftarrow rep( c( 0,0,1, 0,0,0, 0,0,0, 0,0,0), 5)
aprhyp \leftarrow rep( c(0,0,0, 1,0,0, 0,0,0, 0,0,0), 5)
mayhyp \leftarrow rep( c(0,0,0,0,1,0,0,0,0,0,0,0), 5)
junhyp \leftarrow rep( c( 0,0,0, 0,0,1, 0,0,0, 0,0,0 ), 5)
aughyp \leftarrow rep( c(0,0,0,0,0,0,0,1,0,0,0,0), 5)
sephyp \leftarrow rep( c(0,0,0,0,0,0,0,1,0,0,0), 5)
octhyp \leftarrow rep( c( 0,0,0, 0,0,0, 0,0,0, 1,0,0 ), 5)
novhyp \leftarrow rep( c( 0,0,0, 0,0,0, 0,0,0, 0,1,0 ), 5)
dechyp \leftarrow rep( c(0,0,0,0,0,0,0,0,0,0,1), 5)
newdata <- cbind(lawhyp, janhyp, febhyp, marhyp,
                  aprhyp, mayhyp, junhyp, aughyp,
                  sephyp, octhyp, novhyp, dechyp)
# Must be in same order as model!
newdata <- as.data.frame(newdata)
names(newdata) <- c("law", "jan", "feb", "mar", "apr",</pre>
                     "may", "jun", "aug", "sep", "oct",
                     "nov", "dec")
```

```
# Run predict
vpred <- predict(arima.resF.</pre>
                  n.ahead = n.ahead,
                  newxreg = newdata)
# Simulate predicted values
sims <- 10000
simparam <- myrnorm(sims, pe.2f, arima.res2f$var.coef)
xhvp <- newdata
simphi <- simparam[,1:2]
simrho <- simparam[,3:4]
simbetas <- simparam[,5:ncol(simparam)]</pre>
lagY <- c(death[length(death)],death[length(death)-1])</pre>
lagY <- as.vector(lagY)</pre>
lagEps <- c(arima.res2f$resid[length(death)], arima.res2f$resid[length(death)-1])
lagEps <- as.vector(lagEps)
sigma <- sqrt(arima.res2f$sigma)
sim.ev2f <- ldvsimpv(xhyp,
                     simbetas,
                     ci=0.95,
                     constant=1.
                     phi=simphi,
                     lagY=lagY,
                     rho=simrho.
                     lagEps=lagEps,
                     sigma=sigma
```

```
# Make a plot
pdf("prediction1.pdf", width=6, height=3.25)
plot.new()
par(usr = c(0, length(death) + n.ahead, 1000, 3000))
# make the x-axis
axis(1.
     at = seq(from = 10, to = 252, by = 12),
    labels = 1969:1989
axis(2)
title(xlab = "Time",
      vlab = "Deaths".
      main="Predicted effect of reversing seat belt law")
# Polygon of predictive interval for no law (optional)
x0 <- (length(death)+1):(length(death) + n.ahead)
y0 <- c(ypred$pred - 2*ypred$se, rev(ypred$pred + 2*ypred$se), (ypred$pred - 2*ypred$se)[1])
polygon(x = c(x0, rev(x0), x0[1]),
        y = y0,
       border=NA.
        col="#FFBFBFFF"
# Plot the actual data
lines(x = 1:length(death),
      v = death
# Add the predictions for no law
lines(x = length(death):(length(death)+n.ahead),
      y = c(death[length(death)], ypred$pred), # link up the actual data to the prediction
      col = "red"
```