

# Lab 6

*Pstat 174/274*

*Feb 13, 2020*

## Model Estimation

Summary of R commands for estimation:

1. To difference a time series at lag  $d$ :

```
diff(data, lag = d)
```

2. To estimate parameters of an AR model:

```
ar(data, aic = TRUE, order.max = NULL, method = c("..."))
```

If `aic = TRUE`, Akaike's Information Criterion is used to select the order of the AR model to fit. If `aic = FALSE`, the model of order equal to `order.max` is fitted. `method` could be set equal to "yule-walker" or "mle" for different estimation methods of AR parameters.

3. To estimate parameters of an ARMA model:

```
arima(data, order = c(p, 0, q), method = c("..."))
```

The middle number in `order` is the number of times  $d$  the data should be differenced before fitting an ARMA model. For example, `order = c(2, 1, 2)` would fit an ARMA(2,2) once the original time series has been differenced (e.g., to remove the trend component). For regular ARMA models, put 0 for this number. Refer to help file (type `?arima`) for different methods of estimation.

4. To compare models using AICC:

```
AICc(fittedModel)
```

The `qpcR` package needs to be downloaded and installed in R first before using this function. This function will give the Akaike's Information Criterion corrected for bias given a fitted model object from `arima()`. We want to select the model with the *lowest* AICC.

5. To predict future observations given a fitted ARMA model:

```
predict(fittedModel, number of future observations to forecast)
```

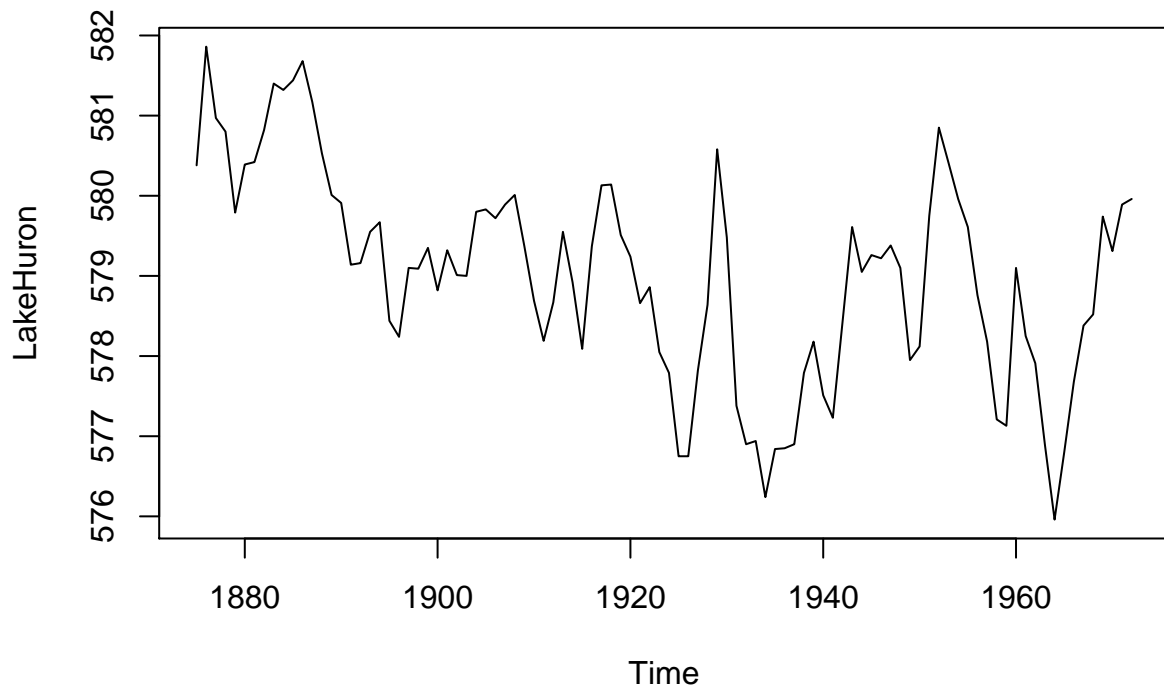
## Lake Huron data

1. Analyze the Lake Huron data (preloaded data set in R). Load the data set using the command

```
data(LakeHuron)
```

and plot the data set using

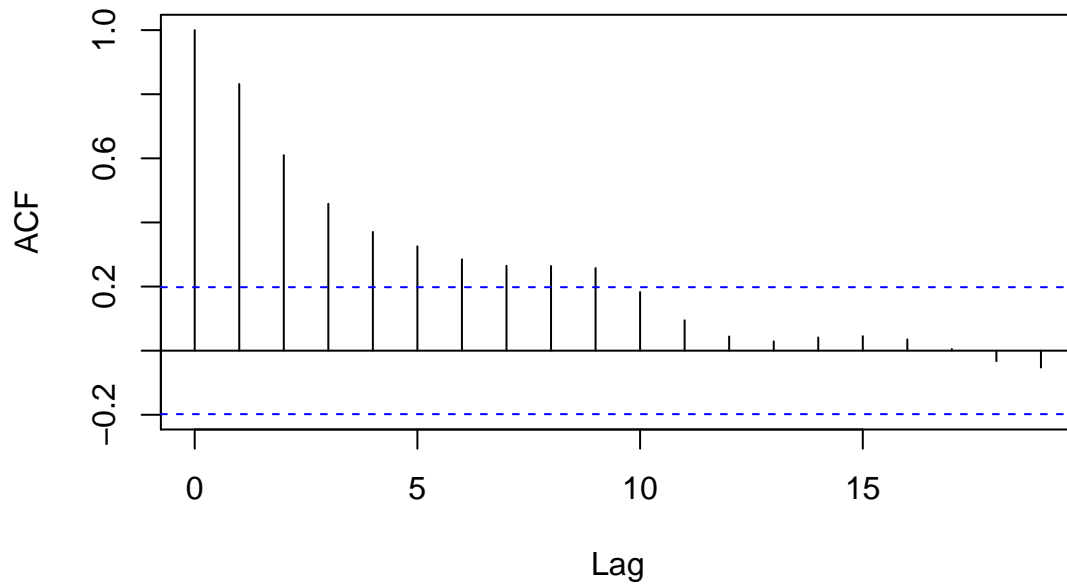
```
ts.plot(LakeHuron)
```



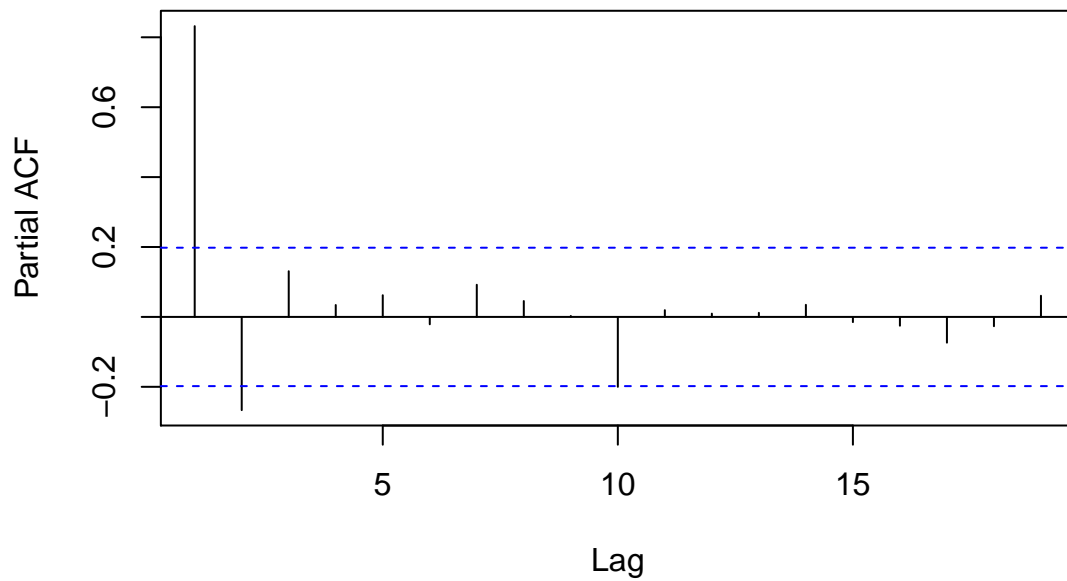
2. Plot the ACF and PACF. What kind of model do they suggest?

```
op <- par(mfrow=c(2,1))  
acf(LakeHuron)  
pacf(LakeHuron)  
par(op)
```

### Series LakeHuron



### Series LakeHuron



3. Consider some possible AR models that might fit the data and perform preliminary Yule-Walker estimation of the model parameters (Hint: use the function `ar()`).

```
# Preliminary estimation using Yule-Walker  
ar(LakeHuron, method="yule-walker")
```

```
##  
## Call:  
## ar(x = LakeHuron, method = "yule-walker")
```

```
##
## Coefficients:
##      1      2
## 1.0538 -0.2668
##
## Order selected 2  sigma^2 estimated as  0.5075
```

4. Consider some possible MA models that might fit the data and perform preliminary estimation of model parameters using the innovations algorithm (see lab6RCode.r for using the innovations algorithm using the function innovations.algorithm() in innovations.r file).

```
# Preliminary estimation using innovations algorithm
source("innovations.r")
?acf
acvf = acf(LakeHuron, plot=FALSE, lag.max = length(LakeHuron))$acf[,1,1] * var(LakeHuron)
m = length(acvf)
lh.ia = innovations.algorithm(m+1, acvf)
lh.ia$thetas[9,1:9] # Preliminary estimates of coefficients for MA(9)

## [1] 1.0810868 0.7691600 0.5218429 0.3315308 0.3085947 0.2479798 0.1932361
## [8] 0.1612645 0.2576989

lh.ia$thetas[10,1:10] # Preliminary estimates of coefficients for MA(10)

## [1] 1.0816255 0.7781248 0.5367164 0.3291559 0.3160040 0.2513755 0.2051537
## [8] 0.1441070 0.3431868 0.1827401

lh.ia$thetas[11,1:11] # Preliminary estimates of coefficients for MA(11)

## [1] 1.08549783 0.77865745 0.54561580 0.34486948 0.31306321 0.26042628
## [7] 0.20956724 0.16011877 0.31610604 0.33734448 0.09479822
```

5. Fit the different models under consideration using maximum likelihood estimation and compare the model fits using AICC (Hint: use arima() for estimation and AICc() in R library qpcR for model comparison - you will need to install this package into R first). Which model is preferred?

```
?arima
# Fit models using maximum likelihood estimation
fit_ar2 = arima(LakeHuron, order = c(2,0,0), method = "ML")
fit_ma9 = arima(LakeHuron, order = c(0,0,9), method = "ML")
fit_ma10 = arima(LakeHuron, order = c(0,0,10), method = "ML")
fit_ma11 = arima(LakeHuron, order = c(0,0,11), method = "ML")
```

Compare models using AICC

```
# install.packages("qpcR")
library(qpcR)
AICc(fit_ar2)
```

```
## [1] 215.5218
```

```
AICc(fit_ma9)
```

```
## [1] 228.0829
```

```
AICc(fit_ma10)
```

```
## [1] 228.9409
```

```
AICc(fit_ma11)
```

```
## [1] 230.2403
```

## Dow Jones Index

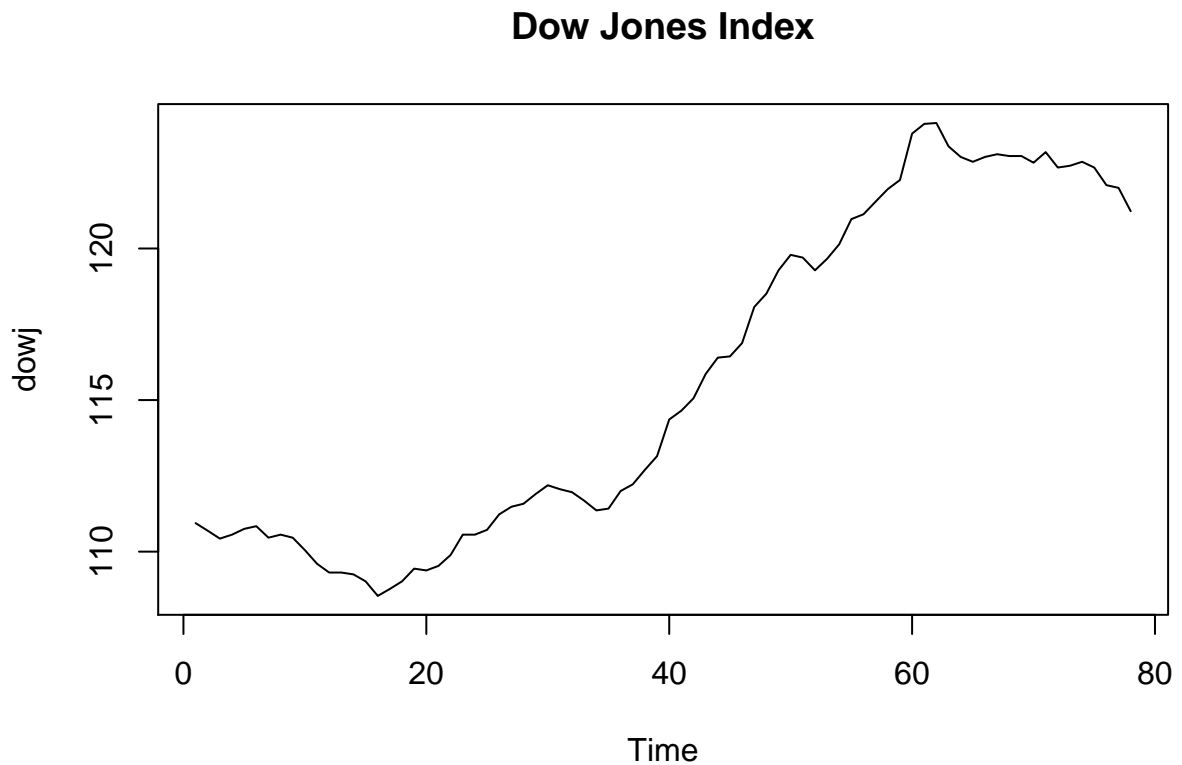
### Exploratory Data Analysis

1. Analyze the Dow Jones Index data by downloading `dowj.txt` from Gauchospace. Move the file into R's working directory and load the data set into R using the command `scan("dowj.txt")`.

```
# Load data
dowj_data <- scan("dowj.txt")
```

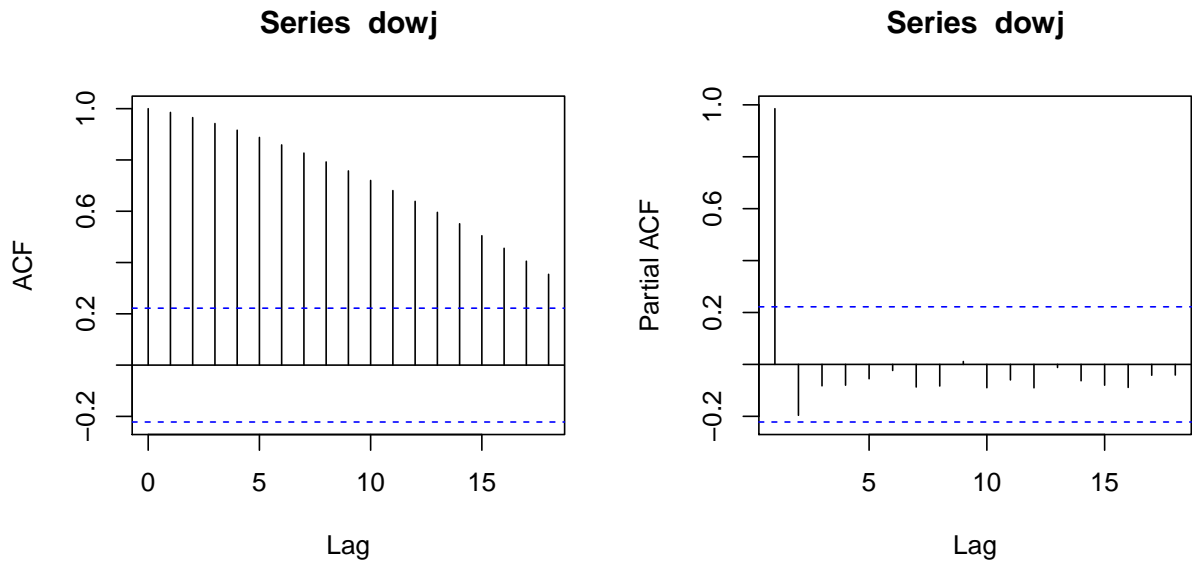
2. Plot the time series. What do you notice?

```
dowj <- ts(dowj_data)
# Plot data
ts.plot(dowj, main = "Dow Jones Index")
```



3. Make the data stationary. What procedures were used?

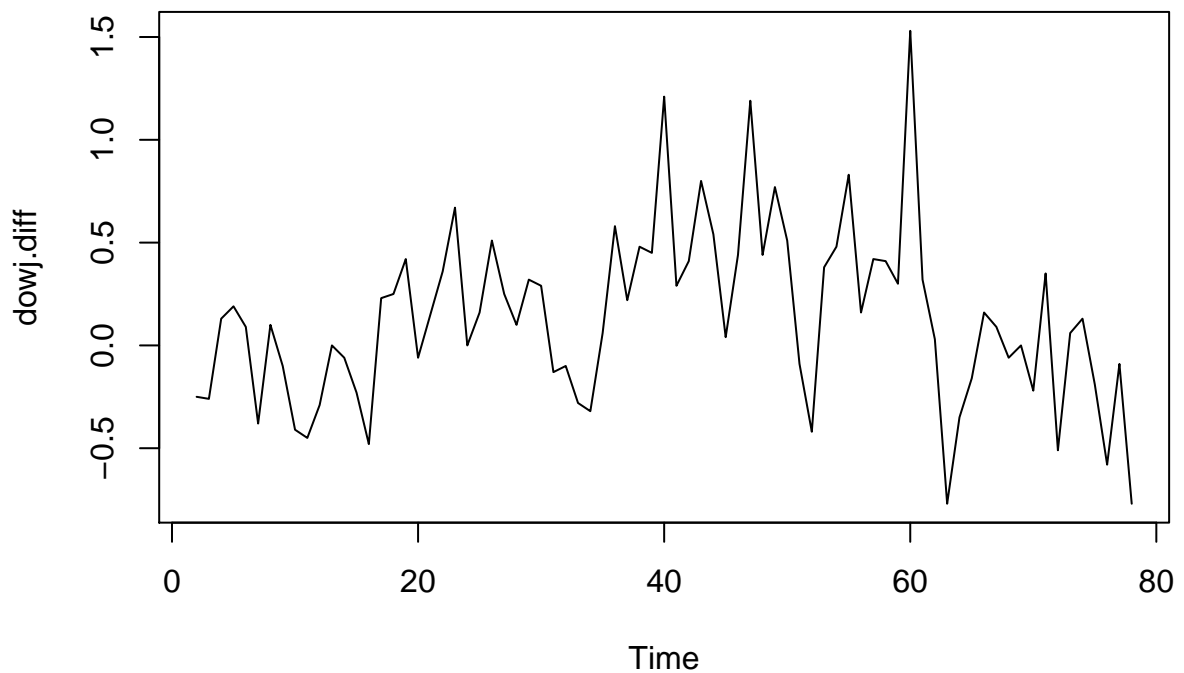
```
op <- par(mfrow=c(1,2))
acf(dowj)
pacf(dowj)
```



```
par(op)

dowj.diff <- diff(dowj,1)
ts.plot(dowj.diff, main = "De-trended data")
```

### De-trended data

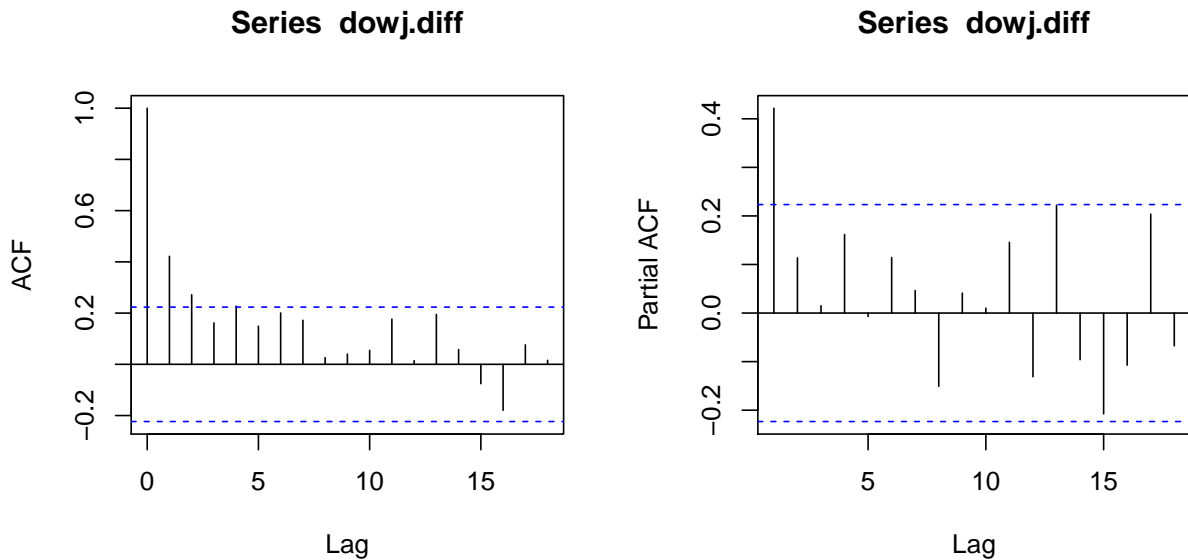


### Model Identification

4. Plot the ACF and PACF. What models do they suggest?

```
op <- par(mfrow=c(1,2))
acf(dowj.diff)
```

```
pacf(dowj.diff)
```



```
par(op)
```

## Model Estimation

5. Fit the AR model suggested by the sample PACF and estimate the coefficients using Yule-Walker estimation.

```
(fit.ar <- ar(dowj.diff, method="yw"))
```

```
##
## Call:
## ar(x = dowj.diff, method = "yw")
##
## Coefficients:
##      1
## 0.4219
##
## Order selected 1  sigma^2 estimated as  0.1518
```

6. Construct 95% confidence intervals for the estimated AR coefficients (Hint: obtain the asymptotic variance of the estimated coefficient from the fitted `ar()` object using `fittedObject$asy.var.coef`).

```
# 95% CI for phi1
ar1.se <- sqrt(fit.ar$asy.var.coef)
c(fit.ar$ar - 1.96*ar1.se, fit.ar$ar + 1.96*1.96*ar1.se)
```

```
## [1] 0.2166839 0.8240603
```

7. Fit different ARMA models using maximum likelihood estimation and compare the model fits using AICC (Hint: use `arima()` for estimation and `AICc()` in `library(qpcR)` for model comparison - you will need to install this package into R first). Which model is preferred?

```
library(qpcR)
# Calculate AICc for ARMA models with p and q running from 0 to 5
aiccs <- matrix(NA, nr = 6, nc = 6)
dimnames(aiccs) = list(p=0:5, q=0:5)
```

```
for(p in 0:5)
{
  for(q in 0:5)
  {
    aiccs[p+1,q+1] = AICc(arima(dowj.diff, order = c(p,0,q), method="ML"))
  }
}
```

```
## Warning in log(s2): NaNs produced
```

```
aiccs
```

```
##      q
## p      0      1      2      3      4      5
## 0 90.49584 81.14893 78.99601 81.17501 80.36057 82.70169
## 1 76.45411 77.04523 78.34399 80.21849 81.48857 83.13492
## 2 77.55328 79.49979 81.56072 81.16088 82.77818 85.09205
## 3 79.69841 81.76437 80.79646 82.25071 84.61907 76.69106
## 4 79.62134 81.83544 83.86231 84.60000 76.04349 86.21979
## 5 81.95313 84.24831 86.18115 87.16679 89.81553 88.75433
```

```
(aiccs==min(aiccs))
```

```
##      q
## p      0      1      2      3      4      5
## 0 FALSE FALSE FALSE FALSE FALSE FALSE
## 1 FALSE FALSE FALSE FALSE FALSE FALSE
## 2 FALSE FALSE FALSE FALSE FALSE FALSE
## 3 FALSE FALSE FALSE FALSE FALSE FALSE
## 4 FALSE FALSE FALSE FALSE  TRUE FALSE
## 5 FALSE FALSE FALSE FALSE FALSE FALSE
```

## Model Diagnostics

8. Perform diagnostics on the chosen model fit. Do the residuals appear to be white noise? Are they normally distributed?

```
# Pick AR(1) and perform residual analysis:
fit = arima(dowj, order=c(1,1,0), method="ML")
```

```
# Test for independence of residuals
Box.test(residuals(fit), type="Ljung")
```

```
##
## Box-Ljung test
##
## data: residuals(fit)
## X-squared = 0.87778, df = 1, p-value = 0.3488
```

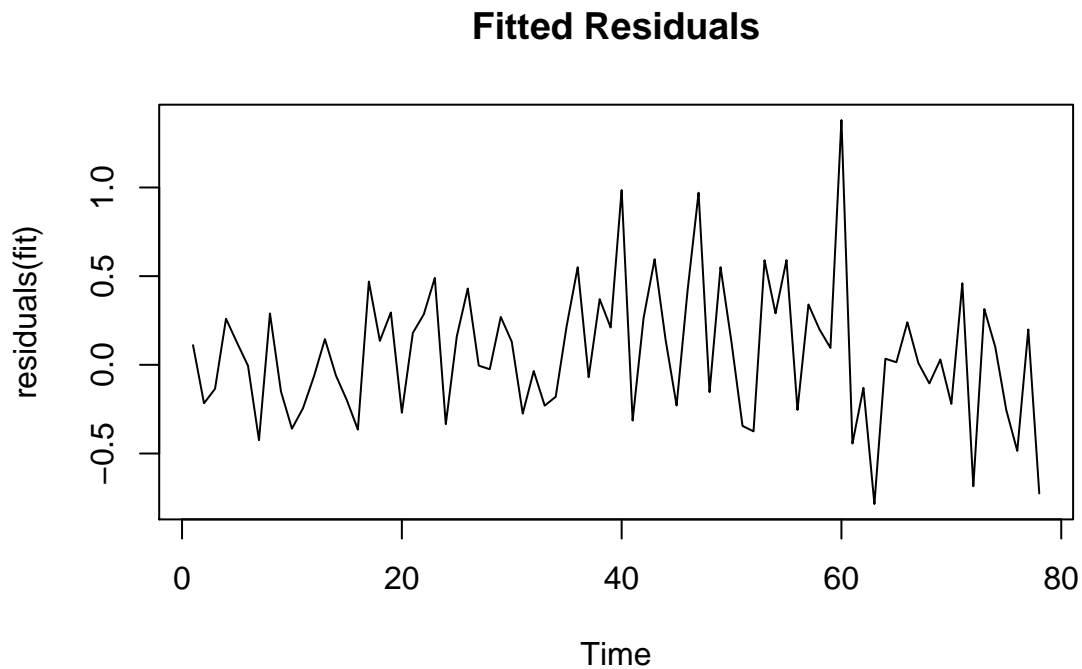
```
# Test for normality of residuals
shapiro.test(residuals(fit))
```

```
##
## Shapiro-Wilk normality test
##
## data: residuals(fit)
```



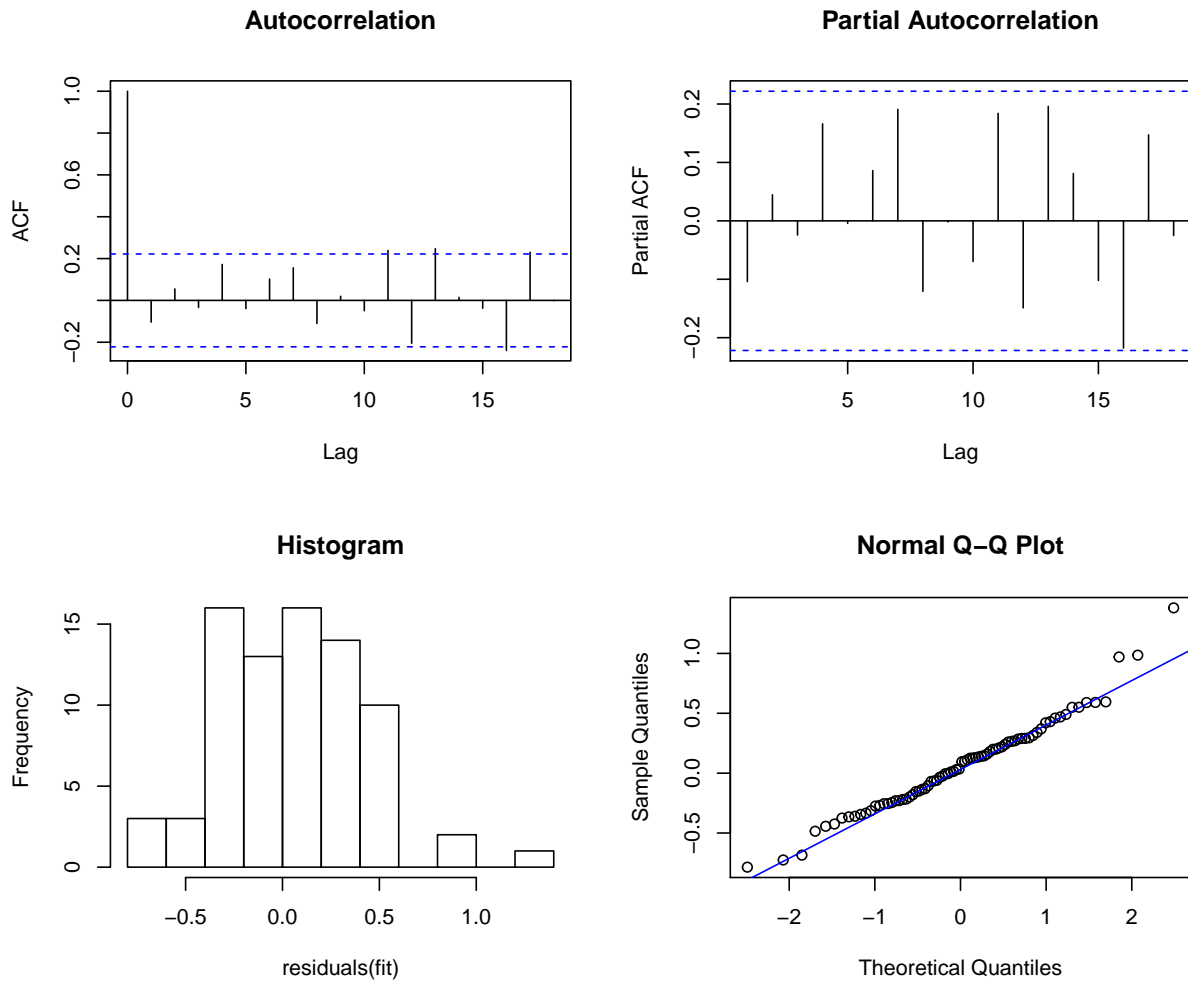
```
## W = 0.97422, p-value = 0.1144
```

```
ts.plot(residuals(fit),main = "Fitted Residuals")
```



```
par(mfrow=c(1,2),oma=c(0,0,2,0))  
# Plot diagnostics of residuals  
op <- par(mfrow=c(2,2))  
# acf  
acf(residuals(fit),main = "Autocorrelation")  
# pacf  
pacf(residuals(fit),main = "Partial Autocorrelation")  
# Histogram  
hist(residuals(fit),main = "Histogram")  
# q-q plot  
qqnorm(residuals(fit))  
qqline(residuals(fit),col = "blue")  
# Add overall title  
title("Fitted Residuals Diagnostics", outer=TRUE)
```

## Fitted Residuals Diagnostics



```
par(op)
```

## Data Forecasting

- Forecast the next 10 observations using your model.

```
# Predict 10 future observations and plot
mypred <- predict(fit, n.ahead=10)
ts.plot(dowj, xlim=c(0,89))
points(79:88,mypred$pred)
lines(79:88,mypred$pred+1.96*mypred$se,lty=2)
lines(79:88,mypred$pred-1.96*mypred$se,lty=2)
```

