

Chapter 6: Box-Jenkins Methodology

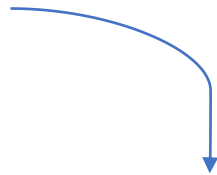
- The methodology consists several steps:

Identification: (see page 107).

Estimation: (see page 89).

Diagnosis: (see page 116).

Forecasting:



1. Stationarity analysis.

2. Invertibility analysis.

3. Residual analysis.

1. Randomness

H_0 : Residuals are random

H_1 : Residuals are not random

Runs test around zero.

2. Constant variance

3. Follow the normal distribution

H_0 : Residuals follow normal distribution

H_1 : Residuals do not follow normal distribution

Shapiro test

4. Test if the residual of the fitted model up to lag k are uncorrelated

H_0 : $\rho_1 = \rho_2 = \dots = \rho_k = 0$

H_1 : at least two $\neq 0$

the Ljung – Box test

4. Fitting the lower model.

5. Fitting the higher model.

The theoretical forms of ACF and PACF for the models: AR(p), MA(q) and ARMA(p, q)

Model	ACF (ρ_k)	PACF (ϕ_{kk})
AR(1)	Approach zero exponentially or in a sinusoidal manner	Cut off completely after the 1 st time lag
AR(2)	Approach zero exponentially or in a sinusoidal manner	Cut off completely after the 2 nd time lag
AR(p)	Approach zero exponentially or in a sinusoidal manner	Cut off completely after time lag p
MA(1)	Cut off completely after the 1 st time lag	Approach zero exponentially or in a sinusoidal manner
MA(2)	Cut off completely after the 2 nd time gap	Approach zero exponentially or in a sinusoidal manner
MA(q)	Cut off completely after a time gap q	Approach zero exponentially or in a sinusoidal manner
ARMA(p, q)	Gradually approaching zero after (q-p) lags exponentially or in a sinusoidal manner	Gradually approaching zero after (p-q) lags exponentially or in a sinusoidal manner

Steps of Time series analysis:

1. Checking stationarity. (Make an appropriate transformations if need)

Differencing can help stabilise the mean of a time series by removing changes in the level of a time series . Box-Cox can help make the variance constant.(R code)

```
(lambda <-BoxCox.lambda(x.D1))
x.B<-BoxCox(x.D1,lambda)
```

2. Checking ACF and PACF.

3. Checking the coefficients.

4. Diagnose the Residuals.

- a. Random, PAC, L-Jung Box and normality graphs
- b. Randomness test
- c. Normality test

5. If we have more than model, we use AIC or BIC to compare.

6. Forecasting.

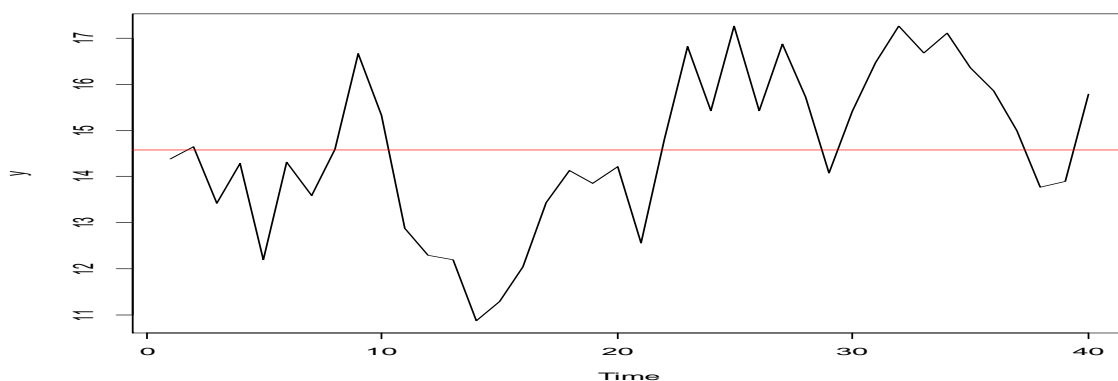
Exercise 1 using R:

the packages used in time series analysis

```
install.packages("forecast")
install.packages("tseries")
install.packages("randtests")
install.packages("astsa")
install.packages("lmtest")
library(forecast)
library(tseries)
library(randtests)
library(astsa)
library(lmtest)
```

1. Checking stationarity of the series:

```
> d=read.csv("ex.csv")
> d=c(14.383,14.649,13.416,14.288,12.201,14.307
,13.586,14.592,16.660,15.332,12.884
,12.296,12.201,10.873,11.290,12.049
,13.435,14.137,13.852,14.213,12.562
,14.801,16.812,15.427,17.268,15.427
,16.869,15.712,14.080,15.408,16.471
,17.268,16.679,17.116,16.357,15.863
,14.991,13.776,13.890,15.787)
> d=ts(d,frequency=1)
> plot(d)
> #plot.ts(d)
```



The data seems to be stationary in the mean.

```
> shapiro.test(d)
Shapiro-Wilk normality test

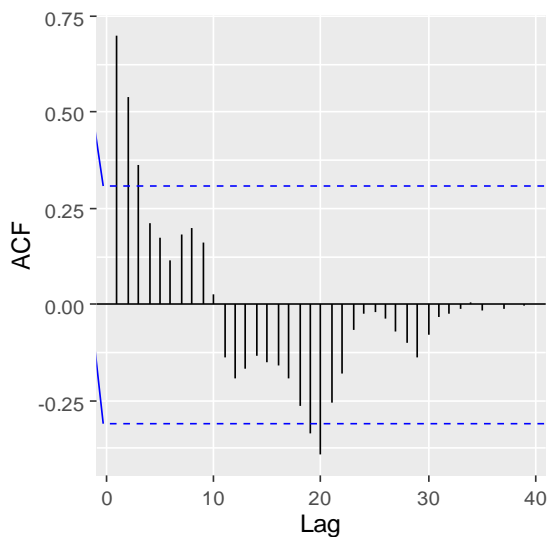
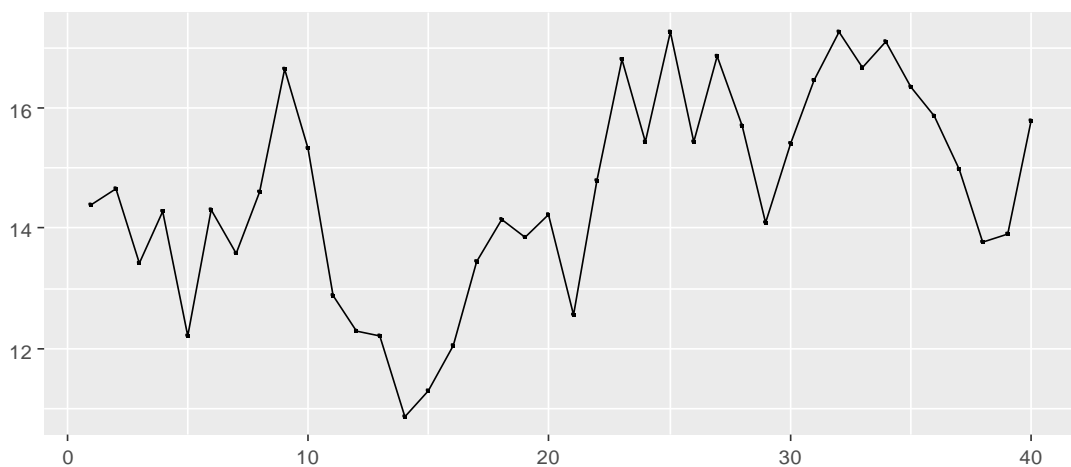
data: d
```

$W = 0.9688$, $p\text{-value} = 0.3296$

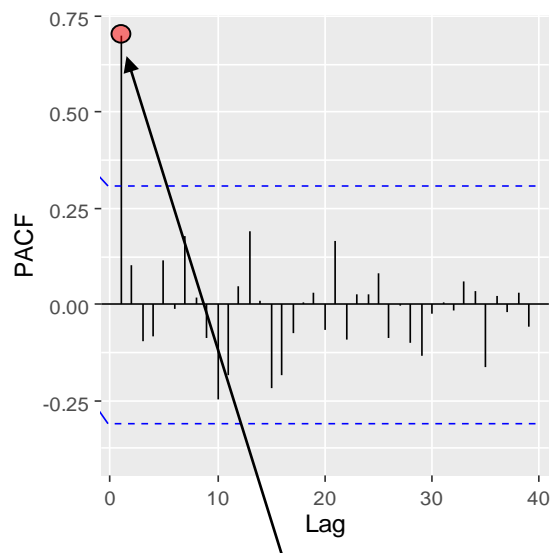
The data seems to be stationary in the variance.

2. Finding the appropriate model using ACF and PACF plot:

```
> ggtsdisplay(d,lag.max=20)
> #acf(d,lag.max=20)
> #pacf(d,lag.max=20)
```

 Or

Approach zero exponentially or in a sinusoidal manner



Cut off completely after the 1st time lag

ARIMA(1,0,0)

- Determine the model:

ARIMA(1,0,0):

```
> fit1=arima(d,order=c(1,0,0))
```

```
> fit1
```

Call:

```
arima(x = d, order = c(1, 0, 0))
```

Coefficients:

```
ar1 intercept
```

```
0.6909 14.6309
```

```
s.e. 0.1094 0.5840
```

```
sigma^2 estimated as 1.447: log likelihood = -64.47, aic = 134.94
```

3. Testing the coefficients for:

```
> coefstest(fit1)
```

z test of coefficients:

	Estimate	Std. Error	z value	Pr(> z)
ar1	0.69090	0.10945	6.3126	2.744e-10 ***
intercept	14.63095	0.58402	25.0523	< 2.2e-16 ***

$$H_0: \phi_1 = 0 \quad vs \quad H_1: \phi_1 \neq 0$$

p -value = $2.744e-10 < 0.05$, we reject H_0

ARIMA(0,0,1):

```
> fit2=arima(d,order=c(0,0,1))
```

```
> coefstest(fit2)
```

z test of coefficients:

	Estimate	Std. Error	z value	Pr(> z)
ma1	0.55701	0.12509	4.453	8.467e-06 ***
intercept	14.58814	0.33366	43.721	< 2.2e-16 ***

$$H_0: \theta_1 = 0 \quad vs \quad H_1: \theta_1 \neq 0$$

p -value = $8.467e-06 < 0.05$, means, we reject H_0

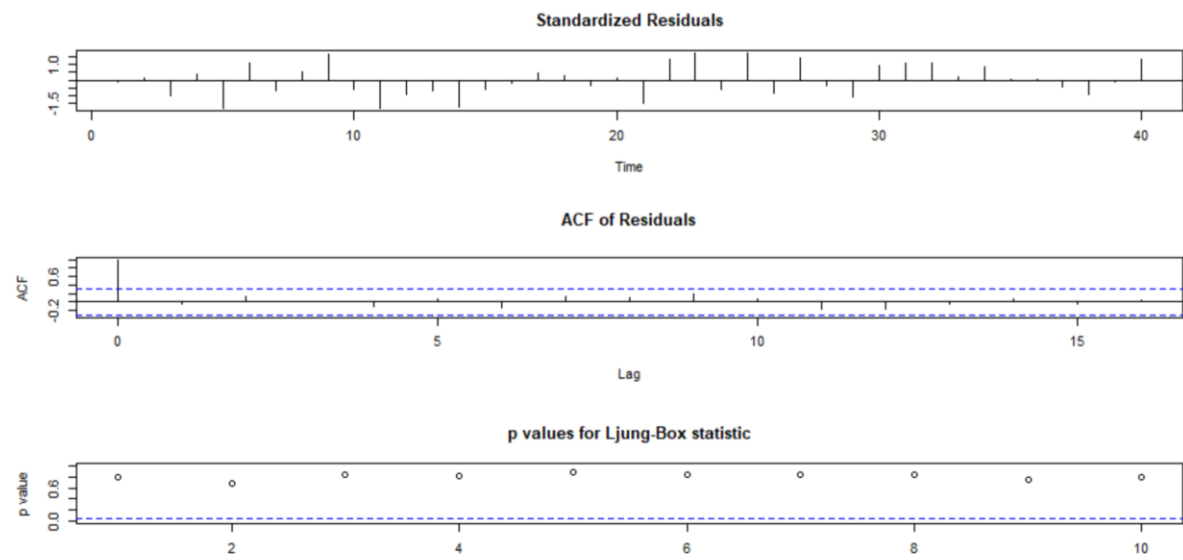
we have the models, $ARIMA(1,0,0)$ and $ARIMA(0,0,1)$

4. Diagnosing the Residuals.

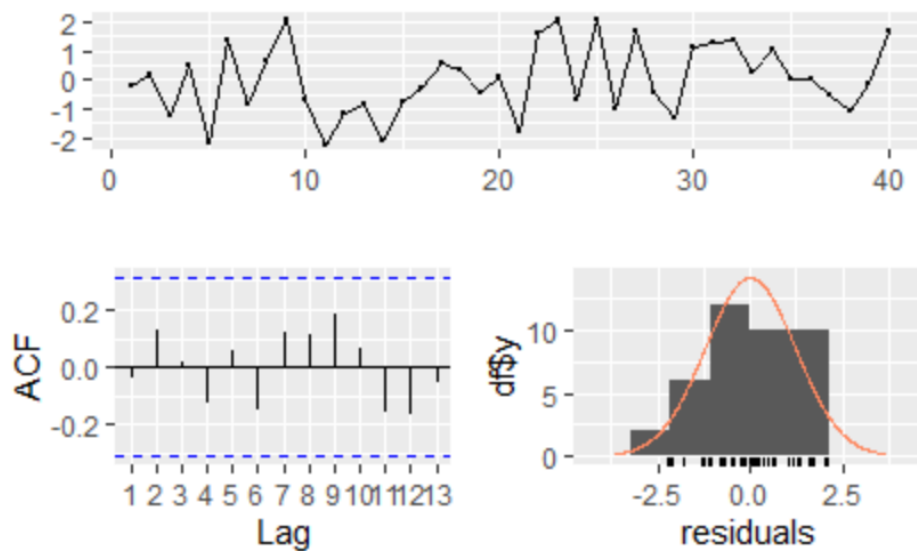
ARIMA(1,0,0)

a) graphs.

```
> tsdiag(fit1)
> checkresiduals(fit1)
```



Residuals from ARIMA(1,0,0) with non-zero mean



- The residuals are random around the zero
- All p-values of the Ljung Box test > 0.05
- The ACF of the Residuals are zeros
- The residuals seem to be normal

b) randomness test

H_0 : Residuals are random

H_1 : Residuals are *not* random

```
> fit1=arima(d,order=c(1,0,0))
> runs.test(fit1$r)
Runs Test
data: fit1$r
statistic = -1.6018, runs = 16, n1 = 20, n2 = 20, n = 40, p-value = 0.7487
alternative hypothesis: nonrandomness
```

$p\text{-value} = 0.7487 > 0.05$, means, we accept H_0 (the residuals are random)

$H_0: E(\varepsilon_t) = 0$ vs $H_1: E(\varepsilon_t) \neq 0$

c) normality test:

H_0 : Residuals follow normal

H_1 : Residuals *do not* follow normal

```
> shapiro.test(fit1$r)
```

```
Shapiro-Wilk normality test
data: fit1$r
W = 0.96633, p-value = 0.2737
```

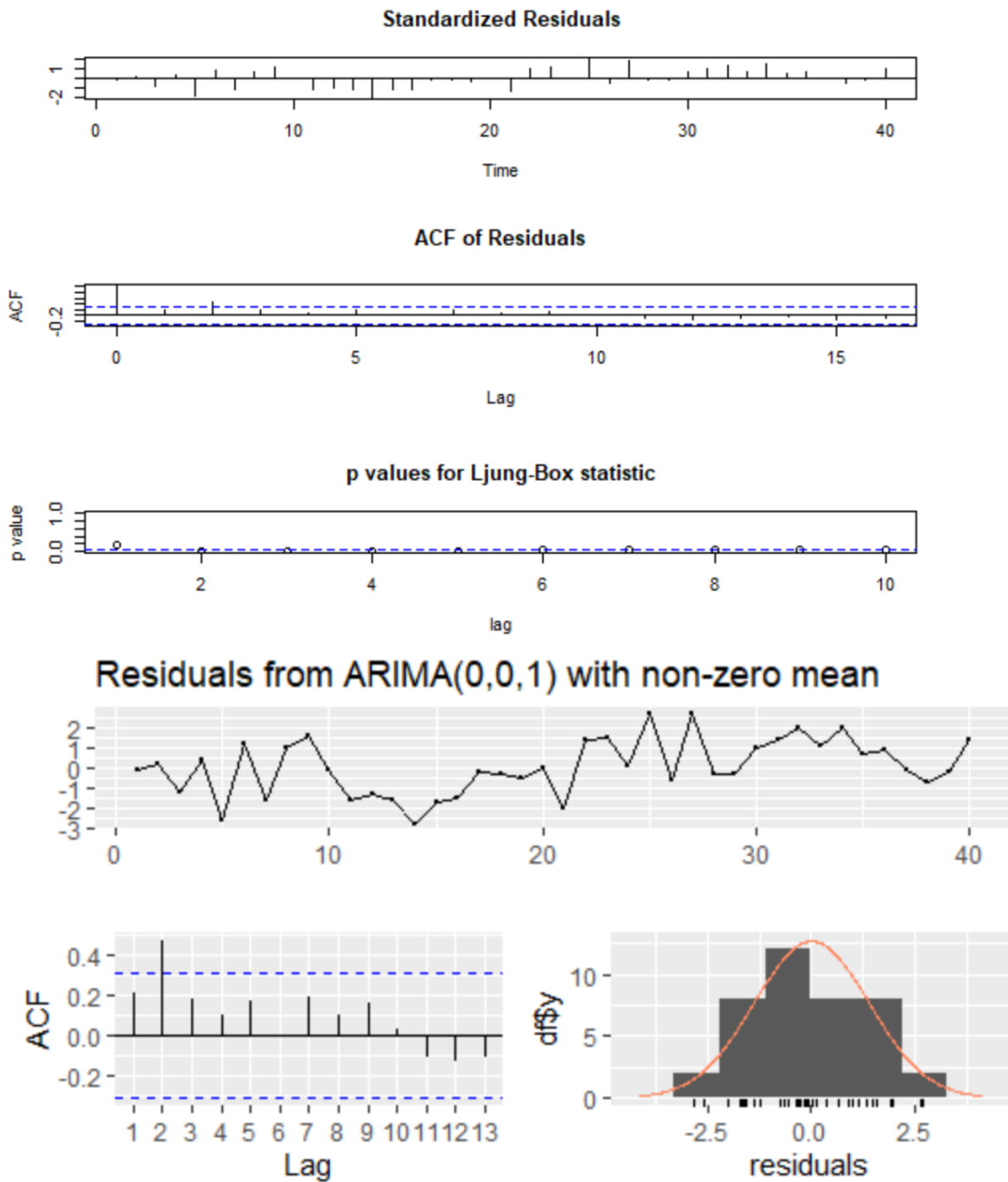
} Accept H_0 (Residuals follow normal)

ARIMA(1,0,0) Model :

$$\hat{y}_t = 14.6309 + 0.6909\hat{y}_{t-1} + \varepsilon_t$$

ARIMA(0,0,1)

```
> fit2=arima(d,order=c(0,0,1))
> tsdiag(fit2)
> checkresiduals(fit2)
```



- The residuals are random around the zero (Except for ρ_2 , it could be a random error)
- Almost all p-values of the Ljung Box test < 0.05
- The ACF of the Residuals are zeros
- The residuals seem to be normal

The fitted model is not adequate


```

> fit2=arima(d,order=c(0,0,1))
> runs.test(fit2$r)
Runs Test
data: fit2$r
statistic = -0.96108, runs = 18, n1 = 20, n2 = 20, n = 40, p-value = 0.3365
alternative hypothesis: nonrandomness

```

$p\text{-value} = 0.3365 > 0.05$, means, we accept H_0 (the residuals are random)

$$H_0: E(\varepsilon_t) = 0 \text{ vs } H_1: E(\varepsilon_t) \neq 0$$

Testing the normality of the residuals:

H_0 : Residuals follow normal

H_1 : Residuals **do not** follow normal

alternative hypothesis: two-sided

```

> shapiro.test(fit2$r)

```

```

Shapiro-Wilk normality test
data: fit2$r
W = 0.97718, p-value = 0.586

```

} Accept H_0 (Residuals follow normal)

ARIMA(0,0,1) Model :

$$\hat{y}_t = 14.58814 + \varepsilon_t - 0.55701 \hat{\varepsilon}_{t-1}$$

More codes for checking normality and Randomness:

```

> hist(fit1$r)
> qqnorm(fit1$r)
> qqline(fit1$r)
> acf(fit1$r)
> pacf(fit1$r)

```

5. Using AIC or BIC to choose between ARIMA(1,0,0) and ARIMA(0,0,1)

```

> fit1=arima(d,order=c(1,0,0))
> fit2=arima(d,order=c(0,0,1))

> fit1$aic
[1] 134.9385
> fit2$aic
[1] 144.9162

```

For forecasting we will use the model with lowest AIC. which is ARIMA(1,0,0)

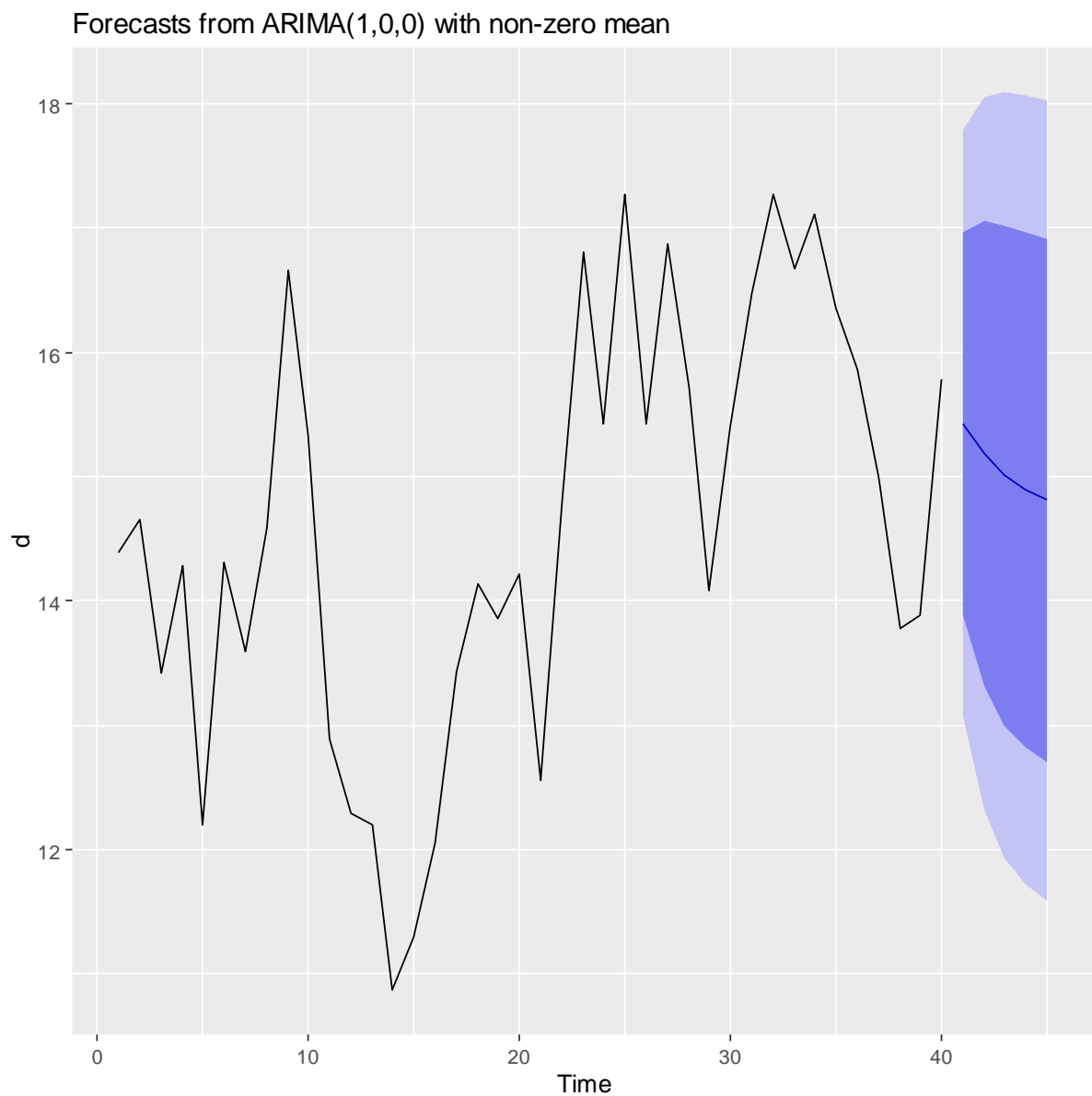
6. Forecasting using ARIMA(1,0,0):

```
> f=forecast(fit1, h=5)
```

```
> autoplot(f)
```

```
> f
```

Point	Forecast	Lo 80	Hi 80	Lo 95	Hi 95
41	15.42967	13.88817	16.97116	13.07215	17.78718
42	15.18278	13.30916	17.05641	12.31732	18.04825
43	15.01221	12.99927	17.02515	11.93369	18.09074
44	14.89436	12.81822	16.97051	11.71917	18.06956
45	14.81294	12.70729	16.91859	11.59263	18.03325



Exercise 2

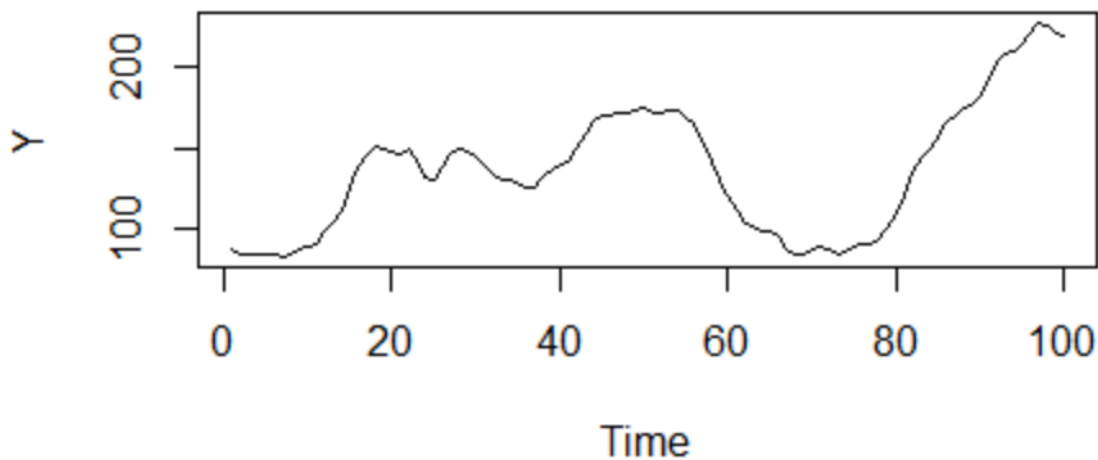
For WWWusage data, is a time series of the number of users on a server every minute for 100 minutes, do the following:

- 1- Plot the series and check its stationarity in mean and variance.
- 2- plot the ACF and PACF , suggest a preliminary model for the data.
- 3- Fit the suggested models and get acquainted with the R output.
- 4- Predict number of users for next 10 minutes.

Solution:

1. Checking stationarity of the series:

```
> data1 = read.csv("C:/ WWWusage.txt", sep="",header=TRUE)
> Y=ts(data1$Y,frequency=1)
> plot(Y)
```



The data seems to be not stationary in the mean.

```
> shapiro.test(Y)
```

Shapiro-Wilk normality test

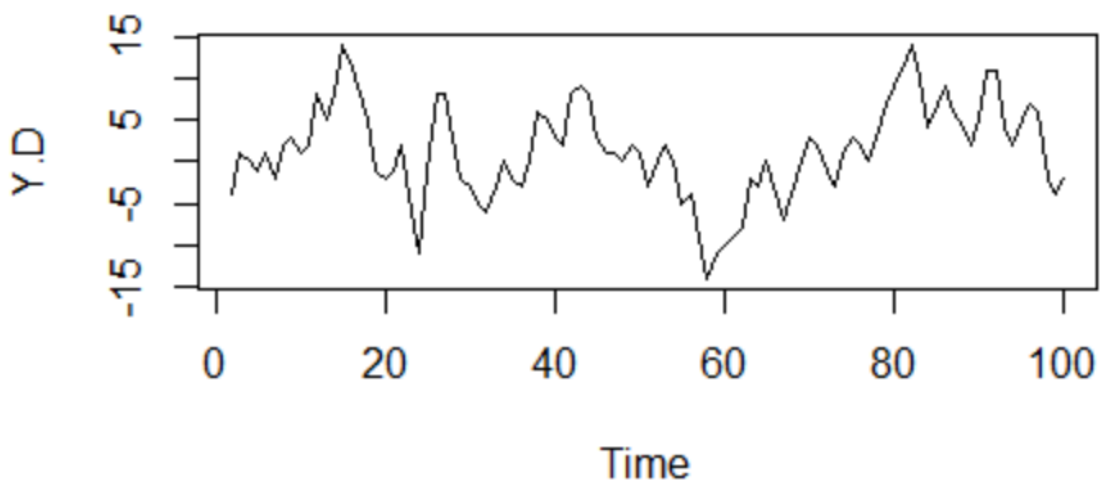
data: Y

W = 0.9373, p-value = 0.0001325

The data is not stationary in the variance.

- *First starting by taking the first difference:*

```
> Y.D<-diff(Y,difference=1)
> plot(Y.D)
```



The data now is seems to be stationary in the mean

```
Shapiro-Wilk normality test
```

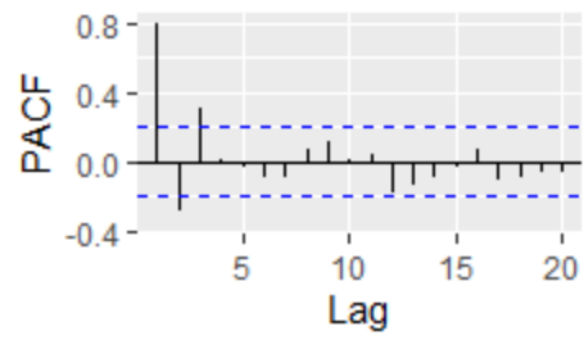
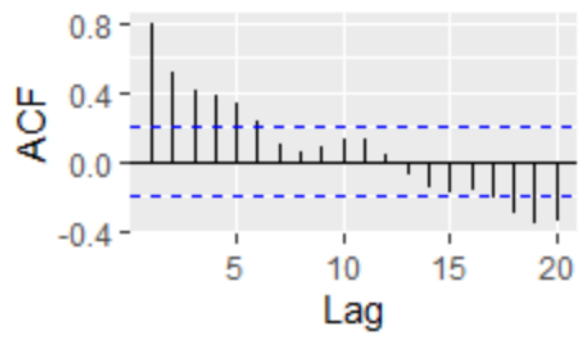
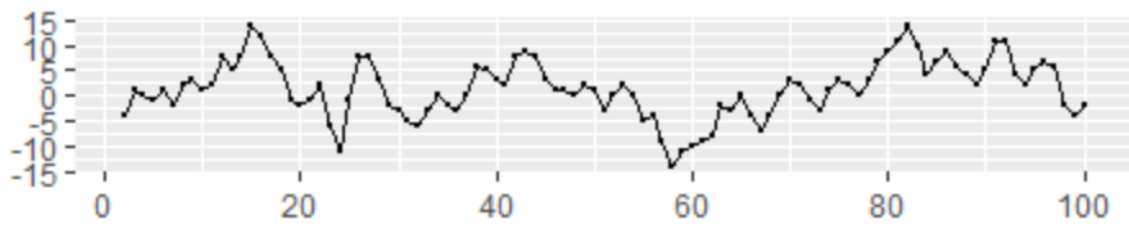
```
data: Y.D
```

```
W = 0.9891, p-value = 0.5997
```

The data now is stationary in the variance.

2. Finding the appropriate model using ACF and PACF plot:

```
> ggtsdisplay(Y.D,lag.max=20)
```



Approach zero exponentially or in a sinusoidal manner

Cut off completely after the 3rd time lag

ARIMA(3,1,0)

- Determine the model:

ARIMA(3,1,0):

```
> fit1=arima(Y,order=c(3,1,0))
```

```
> fit1
```

Call:

```
arima(x = Y, order = c(3, 1, 0))
```

Coefficients:

```

ar1  ar2  ar3
1.1513 -0.6612 0.3407
s.e. 0.0950 0.1353 0.0941

```

sigma² estimated as 9.363: log likelihood = -252, aic = 511.99

3. Testing the coefficients for:

```
> coeftest(fit1)
```

z test of coefficients:

```

Estimate Std. Error z value Pr(>|z|)
ar1 1.151340 0.094984 12.1214 < 2.2e-16 ***

```

```

ar2 -0.661227 0.135263 -4.8885 1.016e-06 ***
ar3 0.340713 0.094146 3.6190 0.0002957 ***
---
Signif. codes:
0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

1) For ϕ_1 :

$$H_0: \phi_1 = 0 \quad vs \quad H_1: \phi_1 \neq 0$$

$p\text{-value} = 2.2e-16 < 0.05$, we reject H_0

2) For ϕ_2 :

$$H_0: \phi_2 = 0 \quad vs \quad H_1: \phi_2 \neq 0$$

$p\text{-value} 1.016e-06 < 0.05$, we reject H_0

3) For ϕ_3 :

$$H_0: \phi_3 = 0 \quad vs \quad H_1: \phi_3 \neq 0$$

$p\text{-value} 0.0002957 < 0.05$, we reject H_0

4. Diagnosing the Residuals.

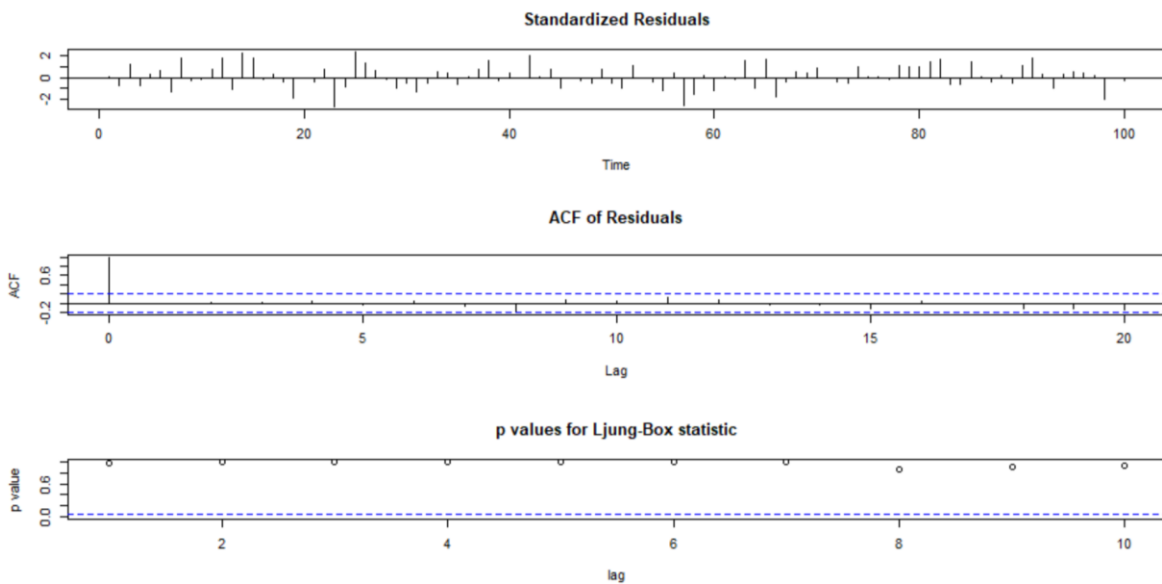
ARIMA(1,0,0)

a) graphs.

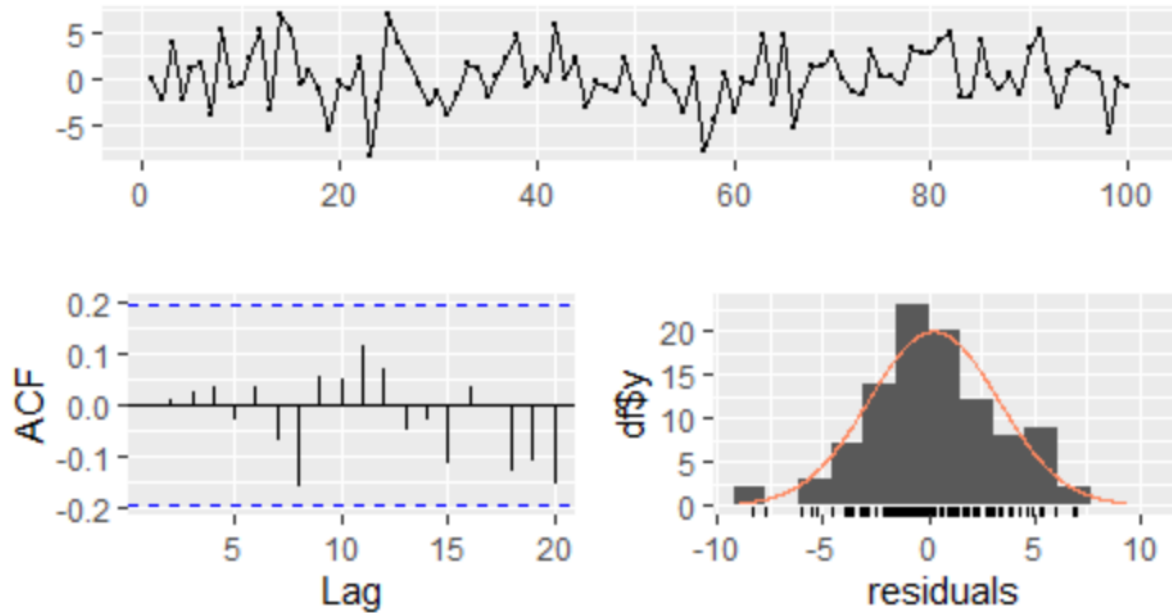
```

> tsdiaf(fit1)
> checkresiduals(fit1)

```



Residuals from ARIMA(3,1,0)



- The residuals are random around the zero
- All p-values of the Ljung Box test > 0.05
- The ACF of the Residuals are zeros
- The residuals seem to be normal

b) randomness test

H_0 : Residuals are random

H_1 : Residuals are **not** random

```
> runs.test(fit1$r)
```

Runs Test

data: fit1\$r

statistic = 0.20102, runs = 52, n1 = 50, n2 =
50, n = 100, p-value = 0.8407

alternative hypothesis: nonrandomness

$p\text{-value} = 0.8407 > 0.05$, means, we accept H_0 (the residuals are random)

$H_0: E(\varepsilon_t) = 0$ vs $H_1: E(\varepsilon_t) \neq 0$

c) normality test:

H_0 : Residuals follow normal
 H_1 : Residuals **do not** follow normal

```
> shapiro.test(fit1$r)
```

Shapiro-Wilk normality test

```
data: fit1$r
```

```
W = 0.98913, p-value = 0.5951
```

Accept H_0 (Residuals follow normal)

ARIMA(3,1,0) Model :

$$(1 - \phi_1 B - \phi_2 B^2 - \phi_3 B^3)(1 - B) y_t = \epsilon_t \gg$$
$$\gg (1 - 1.1513B + 0.6612B^2 - 0.3407B^3)(1 - B) y_t = \epsilon_t$$

6. Forecasting:

```
> f=forecast(fit1, h=10)
```

```
> autoplot(f)
```

```
> f
```

	Point Forecast	Lo 80	Hi 80	Lo 95	Hi 95
101	219.6608	215.7393	223.5823	213.6634	
102	219.2299	209.9265	228.5332	205.0016	
103	218.2766	203.8380	232.7151	196.1947	
104	217.3484	198.3212	236.3756	188.2489	
105	216.7633	193.2807	240.2458	180.8498	
106	216.3785	188.3324	244.4246	173.4858	
107	216.0062	183.3651	248.6473	166.0860	
108	215.6326	178.5027	252.7624	158.8474	
109	215.3175	173.8431	256.7919	151.8879	
110	215.0749	169.3780	260.7719	145.1874	

	Hi 95
101	225.6582
102	233.4581
103	240.3585
104	246.4479
105	252.6768
106	259.2713
107	265.9264
108	272.4178
109	278.7471
110	284.9625

Forecasts from ARIMA(3,1,0)

