

Residual Analysis

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Introduction

In time series models, the residuals are equal to the difference between the observations and the corresponding fitted values:

#Residuals are useful in checking whether a model has adequately captured the information in the data. A good forecasting method will yield residuals with the following properties:

#The residuals are uncorrelated. If there are correlations between residuals, then there is information left in the residuals which should be used in computing forecasts.

#The residuals have zero mean. If the residuals have a mean other than zero, then the forecasts are biased.

Data Description

#The dataset contains the number of earthquakes from the year 1916 to 2015 that occurred in Japan.

Objective

#1) Fit a suitable ARMA model

#2) Perform residual analysis for the fitted model

Analysis

```
library(tseries)
```

#1) Importing the dataset

```
EQ <- read.delim("C:/Users/Srikar/Desktop/SS/R/Sem 6/Practical  
8/earthquakes.txt")
```

```
head(EQ)
```

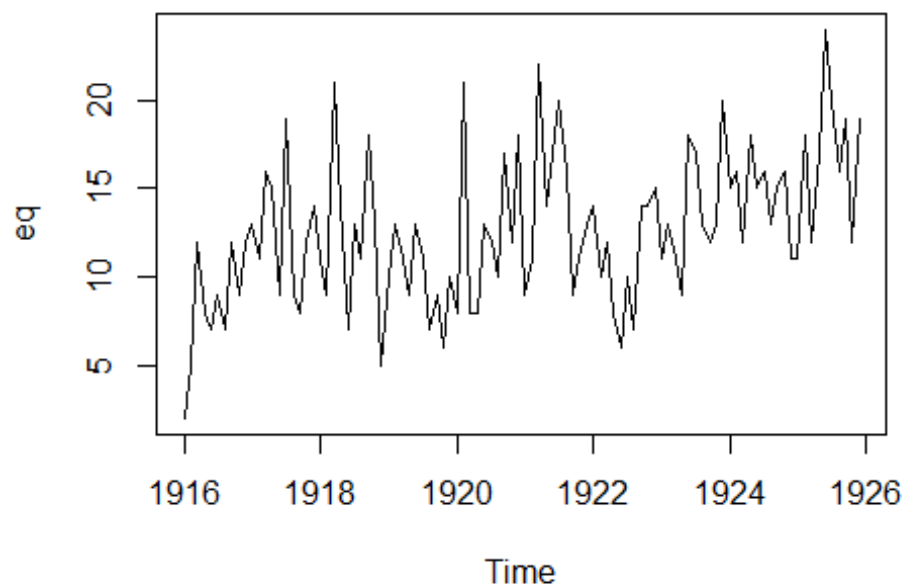
```
##   Year Quakes  
## 1 1916      2  
## 2 1917      5  
## 3 1918     12  
## 4 1919      8  
## 5 1920      7  
## 6 1921      9
```

#Making it a time series object

```
eq=ts(EQ$Quakes,start = 1916,frequency = 10)
```

#2) Plotting the dataset

```
ts.plot(eq)
```



```
adf.test(eq)
```

```
##  
## Augmented Dickey-Fuller Test  
##  
## data: eq
```

```
## Dickey-Fuller = -3.452, Lag order = 4, p-value = 0.04991
## alternative hypothesis: stationary

#We observe that it is stationary

#3) Fitting an ARMA model

library(forecast)

## Warning: package 'forecast' was built under R version 3.6.3

fit=auto.arima(eq,seasonal=FALSE)
fit

## Series: eq
## ARIMA(0,1,1)
##
## Coefficients:
##          ma1
##        -0.8092
## s.e.    0.0710
##
## sigma^2 estimated as 15.53: log likelihood=-276.26
## AIC=556.53   AICc=556.65   BIC=561.72

#The best fitted model is ARMA(0,1). The equation can be given as -0.8092a

#4) Fitting the model

arim=arima(order=c(0,1,1),eq)
```

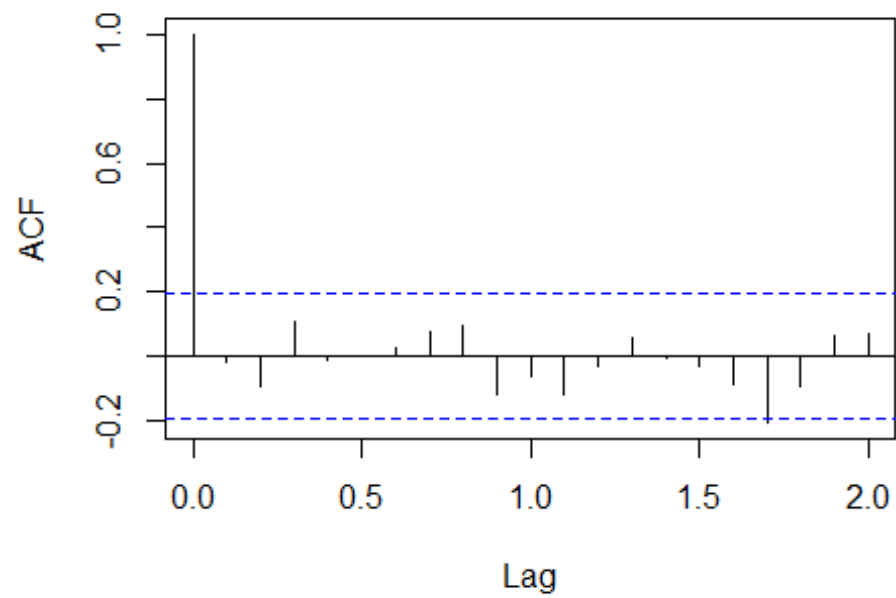
Residual Analysis

```
res=resid(fit)

#1.Checking for uncorrelated errors

acf(res)
```

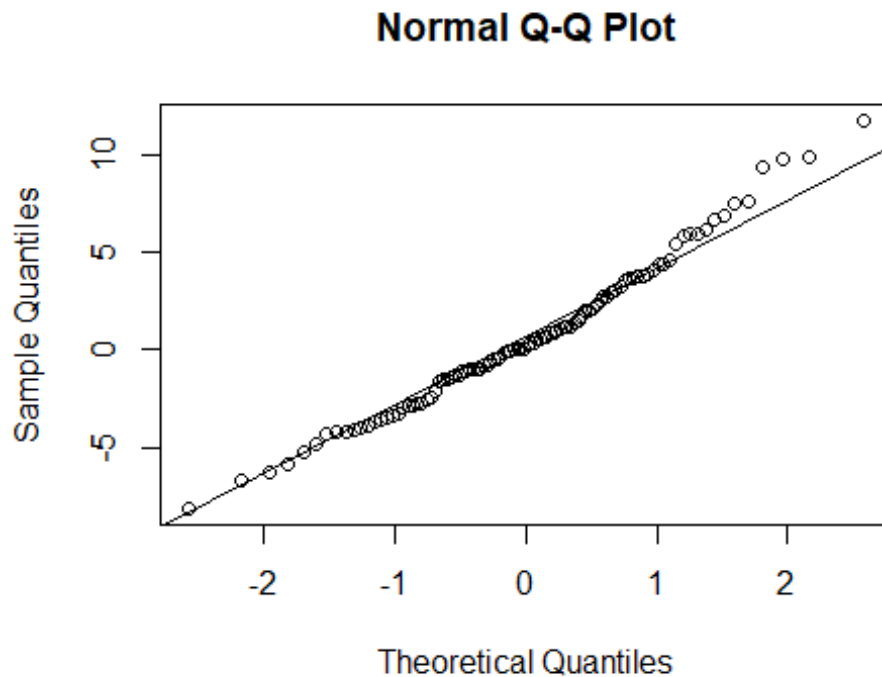
Series res



#From the ACF plot, it is evident that none of the errors are autocorrelated.

#2.Checking if residuals are normally distributed

```
qqnorm(res)  
qqline(res)
```



#We observe that some observations do not fall on the line. To confirm normality, we can use the Shapiro-wilk test.

```
shapiro.test(res)
```

```
##  
##  Shapiro-Wilk normality test  
##  
## data:  res  
## W = 0.98273, p-value = 0.2156
```

#Since p-value is greater than 0.05, we accept the NULL Hypothesis and say that the errors are normally distributed.

Conclusion

The best suited ARMA model is an ARMA(0,1) and from the residual analysis, the assumption of no autocorrelation and normally distributed are satisfied.