

# Introduction to R: basic programming

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```
## Warning: package 'mvtnorm' was built under R version 3.6.2
```

# Introduction

## Slides, code and tutorials

This part of the interactive book contains all R code that was used to produce the results and output presented in chapter 3 (modeling: simple linear regression) in the course's slides. We include YouTube tutorials as a part of the book and links to the relevant tutorials are provided in different chapters. Note that these tutorials were not developed especially for this book, they cover the same topics using different examples.

## R ?

No previous knowledge about R is required. We use the R function `lm()` to fit a simple linear regression model and the cars dataset `cars` is used for illustration. The model can be fitted using the function `glm()` as well.

## Slides

Slides for this part of the course are available online in the [eR-BioStat](#) website. See `RcourseModeling`.

## The cars data

The cars data gives the speed of cars (the response) and the distances taken to stop (the predictor). Note that the data were recorded in the 1920s.

```
x<-cars$speed
y<-cars$dist
head(cars)
```

```
##   speed dist
## 1     4    2
## 2     4   10
## 3     7    4
## 4     7   22
## 5     8   16
## 6     9   10
```

## Fitting Simple Linear regression Model in R

### YouTube tutorials: Simple linear regression in R

#### R - Simple Linear Regression (part 1)

For a YouTube tutorial about simple linear regression in R by Jalayer Academy see [YTREgression1a](#).

#### R - Simple Linear Regression (part 2)

For a YouTube tutorial about simple linear regression by Jalayer Academy in R see [YTREgression1b](#).

## Simple linear regression in R | R Tutorial 5.1

For a YouTube tutorial about simple linear regression in R by MarinStatsLectures see [YTREgression2](#).

### The `lm()` Function

The R function which we use to fit a linear regression model is `lm()`. A General call of the function has the form of `lm(dependent variable~predictor)`.

### Scatterplot

Figure @ref(fig:fig1) shows the scatterplot of Y versus X.

```
plot(x, y)
title("Y vs. X")
```

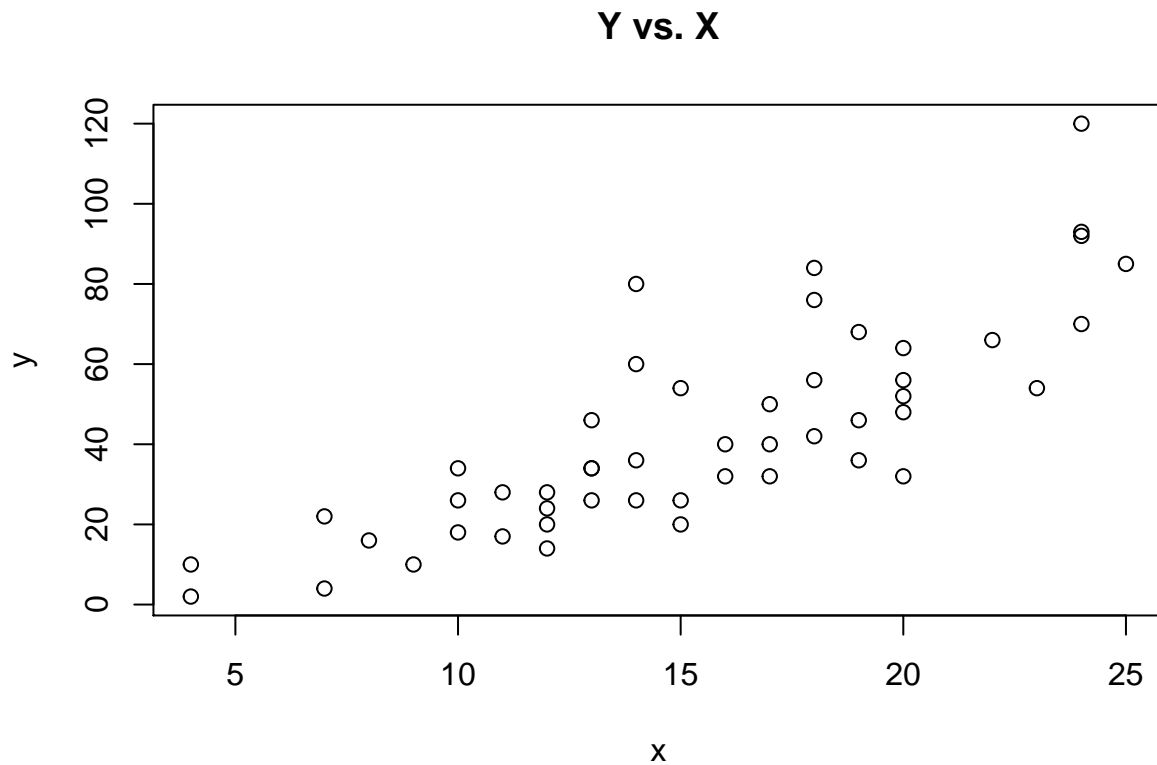


Figure 1: Speed versus stopping distance.

### Fitting the regression model in R

In order to fit a simple linear regression model,

$$Y_i = \beta_0 + \beta_1 X_i + \varepsilon_i, \quad i = 1, \dots, n,$$

we use the `lm()` function in the following way:

```
fit.LM <- lm(y ~ x)
```

The object `fit.LM` contains the results of the estimated model.

```
fit.LM

##
## Call:
## lm(formula = y ~ x)
##
## Coefficients:
## (Intercept)          x
##      -17.579         3.932
```

## Parameter estimates: inference

Parameter estimates, standard errors, t-tests (and p-values) are obtained using the function `summary()`

```
summary(fit.LM)

##
## Call:
## lm(formula = y ~ x)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -29.069  -9.525  -2.272   9.215  43.201
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -17.5791     6.7584  -2.601  0.0123 *
## x              3.9324     0.4155   9.464 1.49e-12 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 15.38 on 48 degrees of freedom
## Multiple R-squared:  0.6511, Adjusted R-squared:  0.6438
## F-statistic: 89.57 on 1 and 48 DF,  p-value: 1.49e-12
```

We use the function `anova()` in order to obtain the ANOVA Table and F-test.

```
anova(fit.LM)

## Analysis of Variance Table
##
## Response: y
##           Df Sum Sq Mean Sq F value    Pr(>F)
## x             1  21186  21185.5   89.567 1.49e-12 ***
## Residuals    48  11354    236.5
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# Residuals and model diagnostic

## Data and fitted model

The object `fit.LM$fit` contains the fitted values.

```
fit.LM$fit

##          1          2          3          4          5          6          7          8
## -1.849460 -1.849460  9.947766  9.947766 13.880175 17.812584 21.744993 21.744993
##          9          10         11         12         13         14         15         16
## 21.744993 25.677401 25.677401 29.609810 29.609810 29.609810 29.609810 33.542219
##         17         18         19         20         21         22         23         24
## 33.542219 33.542219 33.542219 37.474628 37.474628 37.474628 37.474628 41.407036
##         25         26         27         28         29         30         31         32
## 41.407036 41.407036 45.339445 45.339445 49.271854 49.271854 49.271854 53.204263
##         33         34         35         36         37         38         39         40
## 53.204263 53.204263 53.204263 57.136672 57.136672 57.136672 61.069080 61.069080
##         41         42         43         44         45         46         47         48
## 61.069080 61.069080 61.069080 68.933898 72.866307 76.798715 76.798715 76.798715
##         49         50
## 76.798715 80.731124
```

Figure @ref(fig:fig2) shows the scatterplot of Y versus X and the fitted values.

```
plot(x, y)
lines(x, fit.LM$fit)
title("data and fitted model")
```

Note that the straight (the fitted model) line is

$$\hat{Y}_i = \hat{\beta}_0 + \hat{\beta}_1 X_i.$$

## Distribution of the residuals

The object `fit.LM$resid` contains the residuals:  $e_i = Y_i - \hat{Y}_i$ .

```
fit.LM$resid

##          1          2          3          4          5          6          7
##  3.849460 11.849460 -5.947766 12.052234  2.119825 -7.812584 -3.744993
##          8          9         10         11         12         13         14
##  4.255007 12.255007 -8.677401  2.322599 -15.609810 -9.609810 -5.609810
##         15         16         17         18         19         20         21
## -1.609810 -7.542219  0.457781  0.457781 12.457781 -11.474628 -1.474628
##         22         23         24         25         26         27         28
## 22.525372 42.525372 -21.407036 -15.407036 12.592964 -13.339445 -5.339445
##         29         30         31         32         33         34         35
## -17.271854 -9.271854  0.728146 -11.204263  2.795737 22.795737 30.795737
##         36         37         38         39         40         41         42
## -21.136672 -11.136672 10.863328 -29.069080 -13.069080 -9.069080 -5.069080
##         43         44         45         46         47         48         49
##  2.930920 -2.933898 -18.866307 -6.798715 15.201285 16.201285 43.201285
##         50
##  4.268876
```

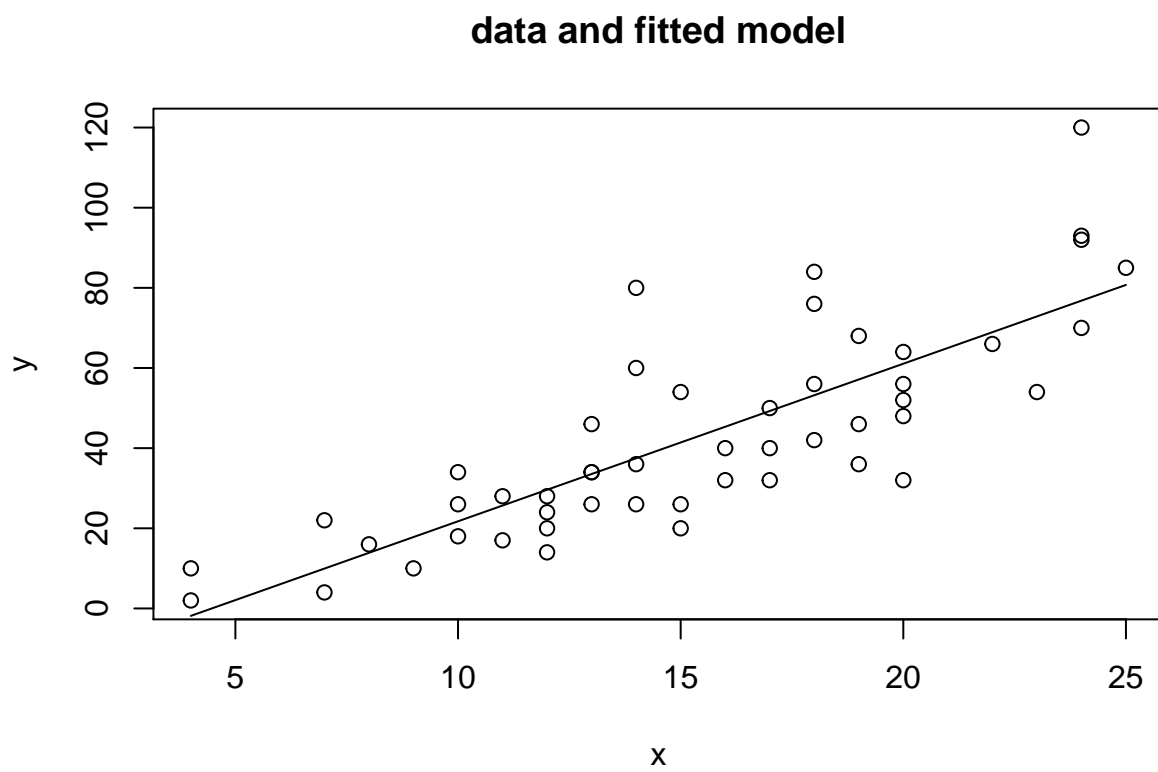


Figure 2: Data and fitted model.

Histogram and normal probability plot for the residuals are shown in Figure @ref(fig:fig3).

```
par(mfrow = c(1, 2))  
hist(fit.LM$resid)  
qqnorm(fit.LM$resid)
```

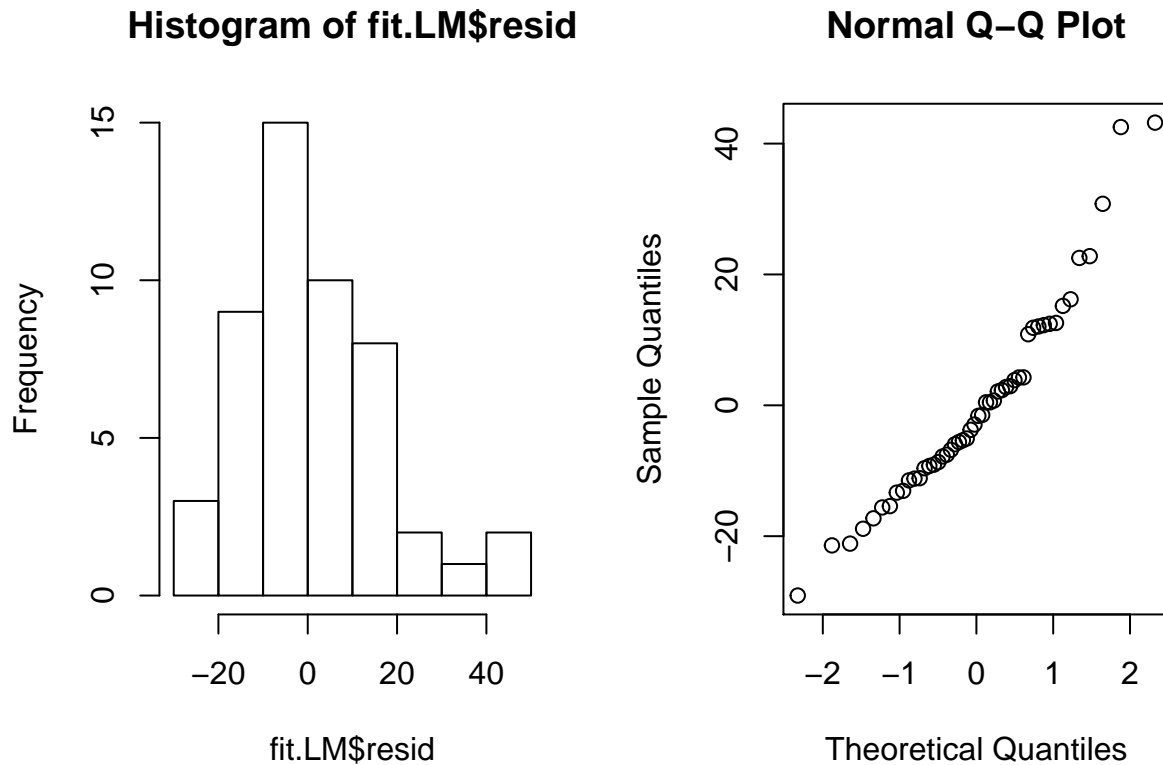


Figure 3: Distribution of the residuals.

Note that the set of diagnostic plots can be produced using the function `plot()` with the object `fit.LM`.

```
plot(fit.LM)
```

