

# Statistical Analysis (Lab Sheet 10.1)

Index No: - 18001149

1. Load the data set “cats” in the package “MASS”. This data set contains Body weight (Bwt) and Heart weight (Hwt) of 144 domestic cats.

- Check whether there is any relationship between Bwt and Hwt, by using a suitable graphical method. Comment on your plot.
- Fit a simple linear regression to model the Heart weight of the cats using Body weight as the independent variable. Write the equation of the fitted model.
- Test the goodness of fit of the fitted model and justify your answer.
- Test the significance of the relationship between the variables Bwt and Hwt. (Mention all the steps clearly).
- Predict the heart weight of a cat if its body weight is 2.52 kg.

beav1	Body Temperature Series of Beaver 1
beav2	Body Temperature Series of Beaver 2
biopsy	Biopsy Data on Breast Cancer Patients
birthwt	Risk Factors Associated with Low Infant Birth Weight
cabbages	Data from a cabbage field trial
caith	Colours of Eyes and Hair of People in Caithness
cats	Anatomical Data from Domestic Cats
cement	Heat Evolved by Setting Cements
chem	Copper in Wholemeal Flour
coop	Co-operative Trial in Analytical Chemistry
cpus	Performance of Computer CPUs
crabs	Morphological Measurements on Leptograpsus Crabs
deaths	Monthly Deaths from Lung Diseases in the UK
drivers	Deaths of Car Drivers in Great Britain 1969-84
eagles	Foraging Ecology of Bald Eagles
epil	Seizure Counts for Epileptics
farms	Ecological Factors in Farm Management
fgl	Measurements of Forensic Glass Fragments
forbes	Forbes' Data on Boiling Points in the Alps
galaxies	Velocities for 82 Galaxies
gehan	Remission Times of Leukaemia Patients
genotype	Rat Genotype Data
geyser	Old Faithful Geyser Data
gilgais	Line Transect of Soil in Gilgai Territory
hills	Record Times in Scottish Hill Races
housing	Frequency Table from a Copenhagen Housing

```
Console Terminal x Jobs x
C:/Local Disk D/Ucsc/Second Year/2nd sem/SCS2211 Laboratory II/tutorial/R tute5/

R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.

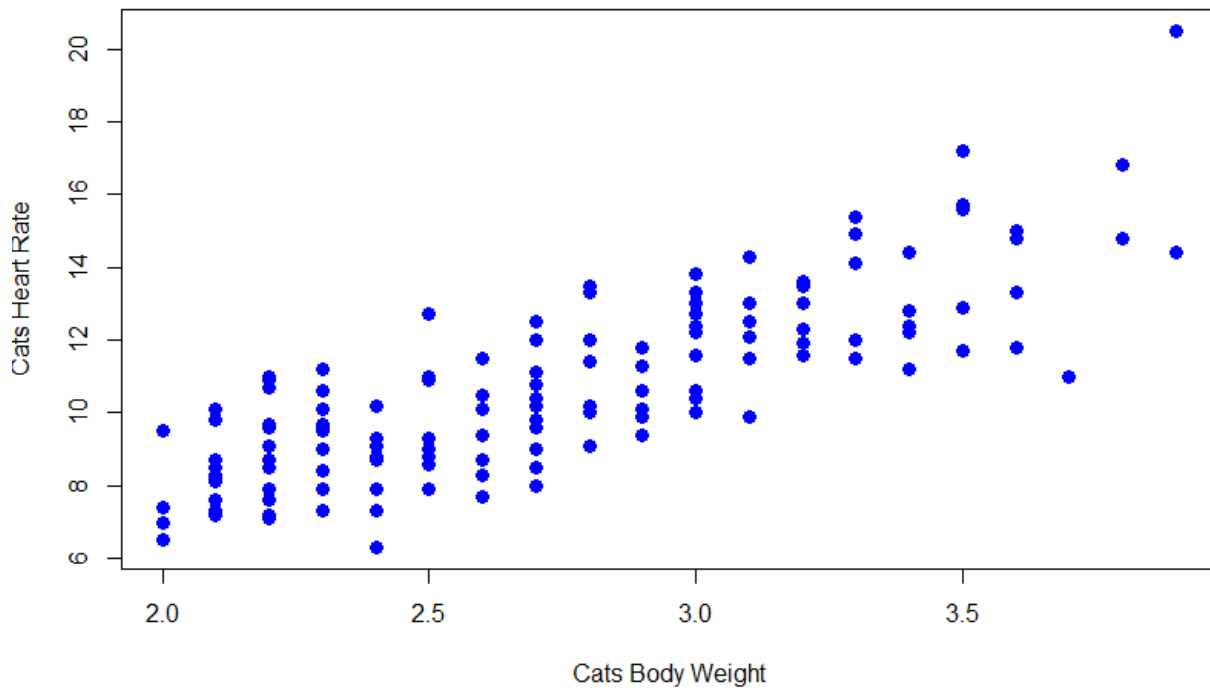
Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

[workspace loaded from ~/.RData]

> setwd("C:/Local Disk D/Ucsc/Second Year/2nd sem/SCS2211 Laboratory II/tutorial
R tute5")
> library('datasets')
> library(datasets)
> library(MASS)
warning message:
package 'MASS' was built under R version 4.0.4
> data()
> |
```

```
a)
> bwt=cats[,2]
> hwt<-cats[,3]
> catone<-data.frame(bwt,hwt)
> plot(catone,col="blue",pch=16,cex =1.3, main = "Cats weight vs Heart Rate",xla
b = "Cats Body Weight",ylab = "Cats Heart Rate")
> |
```

Cats Weight vs Heart Rate



b) `> lm(hwt~bwt,data=cats)`

Call:  
`lm(formula = hwt ~ bwt, data = cats)`

Coefficients:  
 (Intercept)            bwt  
       -0.3567        4.0341

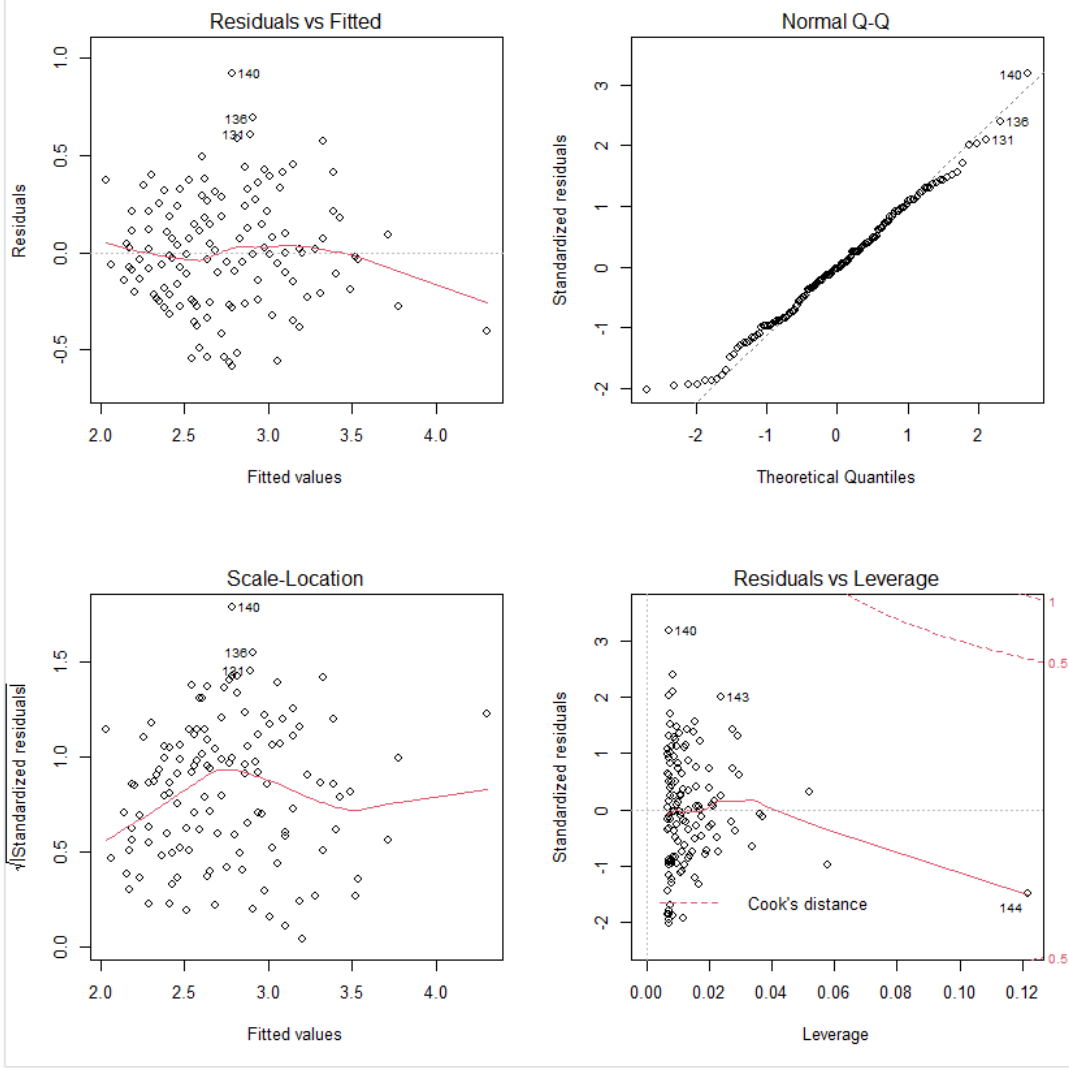
**So, the Formula of This Plot is:**

**Hwt = 4.0341(Bwt) – 0.3567;**

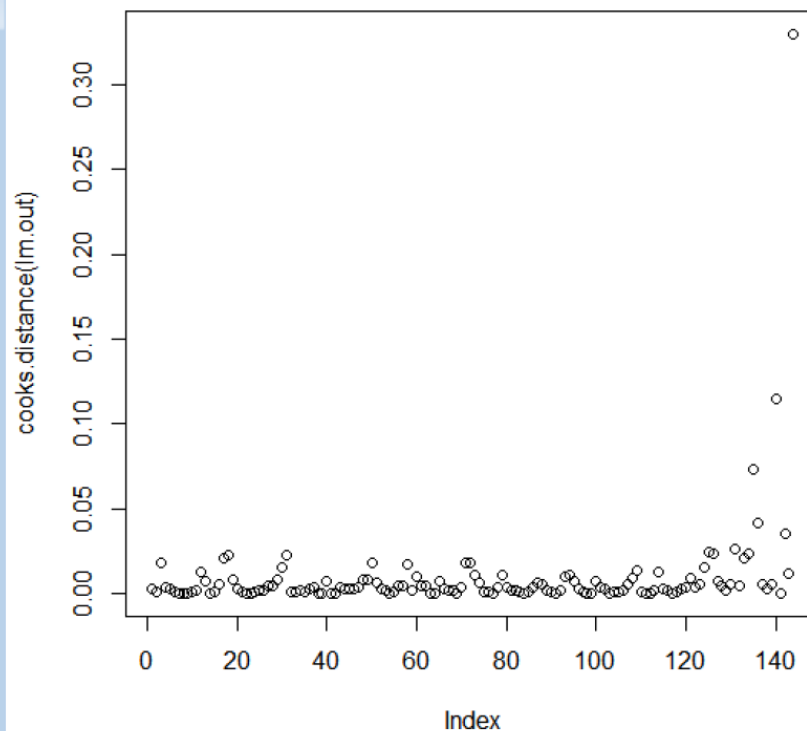
c)

```
> abline(lm(bwt~hwt,data=cats))
> par(mfrow=c(2,2))
> plot(lm(bwt~hwt,data=cats))
> |
```

---



According to this diagram most fitted diagram can be identified as the **“cook’s Distance”**.



d)

e) Predict the heart weight of a cat if its body weight is 2.52 kg.

```
> predict.lm(lm.out,x0,interval = "prediction",cof.level=0.95)
      fit      lwr      upr
1  2.141668 1.562965 2.720372
```

2) Load the data set “iris” in the package “datasets”. This data set gives the measurements in centimeters of the variable’s sepal length and width and petal length and width and the species for 50 flowers from each of 3 species of iris.

Construct a matrix scatter plot for the data and interpret the relationships among variables.

```
> library(datasets)
> str(iris)
'data.frame':  150 obs. of  5 variables:
 $ Sepal.Length: num  5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
 $ Sepal.Width : num  3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
 $ Petal.Length: num  1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
 $ Petal.Width : num  0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
 $ Species      : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1 1 1 1 1 1 ...
```

### # Correlation panel

```
> panel.cor<-function(x,y)
+ {
+   user<-par("usr");
+   on.exit(par(user))
+   par(usr=c(0,1,0,1))
+
+   r<-round(cor(x,y),digits = 2)
+   txt<-paste0("R =",r)
+   cex.cor<-0.8/strwidth(txt)
+   text(0.5,0.5,txt,cex=cex.cor*r)
+ }
> |
```

### # Customize upper panel

```
> upper.panel<-function(x,y)
+ {
+   points(x,y,pch=19,col=c("blue","green","red"))[iris$Species]
+   r<-round(cor(x,y),digits = 2)
+   txt<-paste0("R= ",r)
+   usr<-par("usr");
+   on.exit(par(usr))
+   par(usr=c(0,1,0,1))
+   text(0.5,0.9,txt)
+ }
```

## # Create the Plots

```
pairs(iris[,1:4], lower.panel = panel.cor, upper.panel = upper.panel)
```

