Homework1

陳凱騫(H24101222)

2024-09-19

Table of contents

Import data															1
Data Analysis .															2
Type of varia	ıbles(Q1) .														2
Visualize the	data(Q2)														2
Summarize o	data(Q3) .														6
Symmetric o	or not ?(Q4)														7
Outliers(Q5)															

Import data

```
data <- read.csv(file = "iris.csv")
head(data)</pre>
```

```
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
         5.1
                                1.4
1
                     3.5
                                          0.2 setosa
2
         4.9
                     3.0
                                1.4
                                           0.2 setosa
         4.7
                    3.2
                                1.3
3
                                           0.2 setosa
4
         4.6
                     3.1
                                1.5
                                           0.2 setosa
5
         5.0
                     3.6
                                1.4
                                           0.2 setosa
6
          5.4
                     3.9
                                1.7
                                           0.4 setosa
```

Data Analysis

Type of variables(Q1)

```
for (x in 1:length(data)) {
   print(class(data[,x]))
}

[1] "numeric"
[1] "numeric"
[1] "numeric"
[1] "numeric"
[1] "character"

data$Species <- factor(data$Species)
# change the class of the variable of the species
class(data$Species)</pre>
```

[1] "factor"

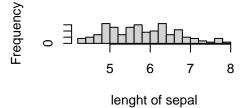
從上方的程式和table可以知道其各個variables如下表:

Variables	Type
Sepal.Length	Cardinal
Sepal.Width	Cardinal
Petal.Length	Cardinal
Petal.Width	Cardinal
Species	Nominal

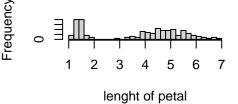
Visualize the data(Q2)

```
par(mfcol=c(2,2))
hist(data$Sepal.Length,main = "Histogram of lenght of sepal"
    ,xlab = "lenght of sepal",breaks = 25)
hist(data$Sepal.Width,main = "Histogram of width of sepal",
    xlab = "width of sepal",breaks = 25)
```

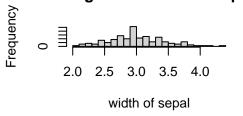
Histogram of lenght of sepal



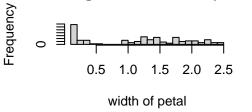
Histogram of lenght of petal



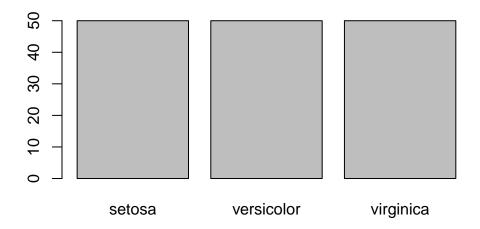
Histogram of width of sepal



Histogram of width of petal

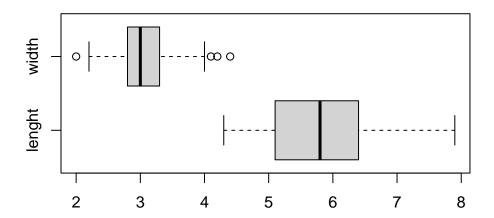


par(mfcol=c(1,1))
plot(data\$Species)



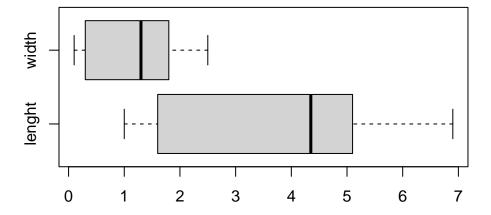
```
# change columns name
sepal <- data[,1:2]
names(sepal) <- c("lenght","width")
petal <- data[,3:4]
names(petal) <- c("lenght","width")
# boxplot
boxplot(sepal,horizontal = T,main="Sepal boxplot")</pre>
```

Sepal boxplot



boxplot(petal,horizontal = T,main="Petal boxplot")

Petal boxplot



Summarize data(Q3)

• 只做cardinal資料

```
# library
library(datasets)
library(e1071) # calculate skewness and kurtosis
# only cardinal data be considered
numeric_vars <- data[, 1:4]</pre>
# create table
summary_table <- data.frame(</pre>
  Mean = apply(numeric_vars, 2, mean),
  Median = apply(numeric_vars, 2, median),
  Variance = apply(numeric_vars, 2, var),
  Std_Dev = apply(numeric_vars, 2, sd),
  Range = apply(numeric_vars, 2, function(x) diff(range(x))),
  IQR = apply(numeric_vars, 2, IQR),
  Skewness = apply(numeric_vars, 2, skewness),
  Kurtosis = apply(numeric_vars, 2, kurtosis)
)
# print result
print(summary_table)
```

```
Mean MedianVarianceStd_DevRangeIQRSkewnessSepal.Length5.8433335.800.68569350.82806613.61.30.3086407Sepal.Width3.0573333.000.18997940.43586632.40.50.3126147Petal.Length3.7580004.353.11627791.76529825.93.5-0.2694109Petal.Width1.1993331.300.58100630.76223772.41.5-0.1009166KurtosisSepal.Length-0.6058125Sepal.Width0.1387047Petal.Length-1.4168574Petal.Width-1.3581792
```

 nominal data only need to know mode and we know that all of three species are the same, 50.

```
summary(data$Species)
```

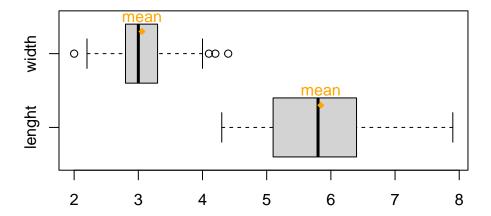
```
setosa versicolor virginica
50 50 50
```

Symmetric or not ?(Q4)

使用boxplot來決定是否對稱

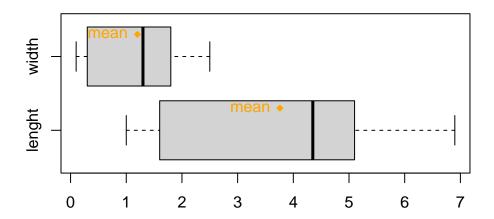
```
# boxplot
means <- apply(sepal, 2, mean)
meanp <- apply(petal, 2, mean)
bs <- boxplot(sepal,horizontal = T,main="Sepal boxplot")
xi <- 0.3 + seq(bs$n)
points(means,xi , col = "orange", pch = 18)
text(means,xi,"mean",pos=3,col = "orange")</pre>
```

Sepal boxplot



```
bp <- boxplot(petal,horizontal = T,main="Petal boxplot")
xi <- 0.3 + seq(bp$n)
points(meanp,xi , col = "orange", pch = 18)
text(meanp,xi,"mean",pos=2,col = "orange")</pre>
```

Petal boxplot



從上方幾張**boxplot**可知,**sepal**的lenght和width基本上為**對稱**,至於**petal**的lenght和width不**對稱**,原因為mean和median之間有差距,且median看起來似乎也比較不在資料的中心。而從visualize data 那裡的histogram也可發現資料(petal)似乎有一點雙峰的感覺。

Outliers(Q5)

從上方Sepal boxplot中可以發現sepal.width有4個極端值。

```
outliers <- c()
outliers <- c(outliers, sort(data$Sepal.Width)[1])
for (i in 0:2) {
  outliers <- c(outliers, sort(data$Sepal.Width)[length(data$Sepal.Width)-i])
}
outliers</pre>
```

[1] 2.0 4.4 4.2 4.1

從中可知其outliers為:

Outliers 2 4.4 4.2 4.1