

# Data Exploration and Visualisation with R <sup>1</sup>

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<sup>1</sup>Chapter 3: Data Exploration, in *R and Data Mining: Examples and Case Studies*.

## Introduction

Have a Look at Data

Explore Individual Variables

Explore Multiple Variables

More Explorations

Save Charts to Files

Further Readings and Online Resources

## Data Exploration and Visualisation

- ▶ Summary and stats
- ▶ Various charts like pie charts and histograms
- ▶ Exploration of multiple variables
- ▶ Level plot, contour plot and 3D plot
- ▶ Saving charts into files

# Quiz: What's the Name of This Flower?




Oleg Yunakov [CC BY-SA 3.0 (<https://creativecommons.org/licenses/by-sa/3.0>)], from Wikimedia Commons.

The iris dataset [Frank and Asuncion, 2010] consists of 50 samples from each of three classes of iris flowers. There are five attributes in the dataset:

- ▶ sepal length in cm,
- ▶ sepal width in cm,
- ▶ petal length in cm,
- ▶ petal width in cm, and
- ▶ class: Iris Setosa, Iris Versicolour, and Iris Virginica.

Detailed description of the dataset can be found at the UCI Machine Learning Repository <sup>2</sup>.

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<sup>2</sup><https://archive.ics.uci.edu/ml/datasets/Iris> 

Introduction

Have a Look at Data

Explore Individual Variables

Explore Multiple Variables

More Explorations

Save Charts to Files

Further Readings and Online Resources

# Size and Variables Names of Data

```
# number of rows
nrow(iris)
## [1] 150

# number of columns
ncol(iris)
## [1] 5

# dimensionality
dim(iris)
## [1] 150    5

# column names
names(iris)
## [1] "Sepal.Length" "Sepal.Width"  "Petal.Length" "Petal.Wid..."
## [5] "Species"
```

Below we have a look at the structure of the dataset with `str()`.

```
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...
##  $ Petal.Width : num  0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0...
##  $ Species      : Factor w/ 3 levels "setosa","versicolor",....
```

- ▶ 150 observations (records, or rows) and 5 variables (or columns)
- ▶ The first four variables are numeric.
- ▶ The last one, Species, is categoric (called “factor” in R) and has three levels of values.



```
attributes(iris)
## $names
## [1] "Sepal.Length" "Sepal.Width"  "Petal.Length" "Petal.Wid..."
## [5] "Species"
##
## $class
## [1] "data.frame"
##
## $row.names
##      [1]      1      2      3      4      5      6      7      8      9     10     11     12     13     ...
##     [16]     16     17     18     19     20     21     22     23     24     25     26     27     28     ...
##     [31]     31     32     33     34     35     36     37     38     39     40     41     42     43     ...
##     [46]     46     47     48     49     50     51     52     53     54     55     56     57     58     ...
##     [61]     61     62     63     64     65     66     67     68     69     70     71     72     73     ...
##     [76]     76     77     78     79     80     81     82     83     84     85     86     87     88     ...
##     [91]     91     92     93     94     95     96     97     98     99    100    101    102    103    1...
##    [106]    106    107    108    109    110    111    112    113    114    115    116    117    118    1...
##    [121]    121    122    123    124    125    126    127    128    129    130    131    132    133    1...
##    [136]    136    137    138    139    140    141    142    143    144    145    146    147    148    1...
```

```
iris[1:3,]
```

```
##      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1           5.1           3.5           1.4           0.2  setosa
## 2           4.9           3.0           1.4           0.2  setosa
## 3           4.7           3.2           1.3           0.2  setosa
```

```
head(iris, 3)
```

```
##      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1           5.1           3.5           1.4           0.2  setosa
## 2           4.9           3.0           1.4           0.2  setosa
## 3           4.7           3.2           1.3           0.2  setosa
```

```
tail(iris, 3)
```

```
##      Sepal.Length Sepal.Width Petal.Length Petal.Width  Spe...
## 148           6.5           3.0           5.2           2.0  virgi...
## 149           6.2           3.4           5.4           2.3  virgi...
## 150           5.9           3.0           5.1           1.8  virgi...
```

The first 10 values of Sepal.Length

```
iris[1:10, "Sepal.Length"]  
##    [1] 5.1 4.9 4.7 4.6 5.0 5.4 4.6 5.0 4.4 4.9  
  
iris$Sepal.Length[1:10]  
##    [1] 5.1 4.9 4.7 4.6 5.0 5.4 4.6 5.0 4.4 4.9
```

Introduction

Have a Look at Data

Explore Individual Variables

Explore Multiple Variables

More Explorations

Save Charts to Files

Further Readings and Online Resources

## Function summary()

- ▶ numeric variables: minimum, maximum, mean, median, and the first (25%) and third (75%) quartiles
- ▶ categorical variables (i.e., factors): frequency of every level

```
summary(iris)

##      Sepal.Length      Sepal.Width      Petal.Length      Petal.Wid...
## Min.       :4.300      Min.       :2.000      Min.       :1.000      Min.       :0....
## 1st Qu.:5.100      1st Qu.:2.800      1st Qu.:1.600      1st Qu.:0....
## Median :5.800      Median :3.000      Median :4.350      Median :1....
## Mean    :5.843      Mean    :3.057      Mean    :3.758      Mean    :1....
## 3rd Qu.:6.400      3rd Qu.:3.300      3rd Qu.:5.100      3rd Qu.:1....
## Max.     :7.900      Max.     :4.400      Max.     :6.900      Max.     :2....
##
##      Species
## setosa      :50
## versicolor:50
## virginica   :50
##
##
##
```

```
library(Hmisc)
# describe(iris) # check all columns
describe(iris[, c(1,5)]) # check columns 1 and 5
## iris[, c(1, 5)]
##
## 2 Variables      150 Observations
## -----...
## Sepal.Length
##      n missing distinct      Info      Mean      Gmd      ...
##    150      0      35    0.998    5.843    0.9462    4....
##    .10     .25     .50     .75     .90     .95
##    4.800    5.100    5.800    6.400    6.900    7.255
##
## lowest : 4.3 4.4 4.5 4.6 4.7, highest: 7.3 7.4 7.6 7.7 7.9
## -----...
## Species
##      n missing distinct
##    150      0      3
##
## Value      setosa versicolor virginica
## Frequency      50      50      50
## Proportion    0.333    0.333    0.333
## -----...
```

- ▶ Mean, median and range: `mean()`, `median()`, `range()`
- ▶ Quartiles and percentiles: `quantile()`

```
range(iris$Sepal.Length)
```

```
## [1] 4.3 7.9
```

```
quantile(iris$Sepal.Length)
```

```
##    0%   25%   50%   75%  100%
```

```
##  4.3   5.1   5.8   6.4   7.9
```

```
quantile(iris$Sepal.Length, c(.1, .3, .65))
```

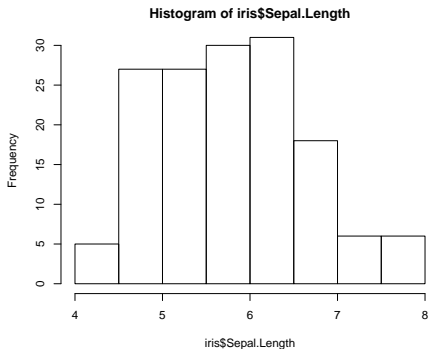
```
##   10%   30%   65%
```

```
## 4.80 5.27 6.20
```

# Variance and Histogram

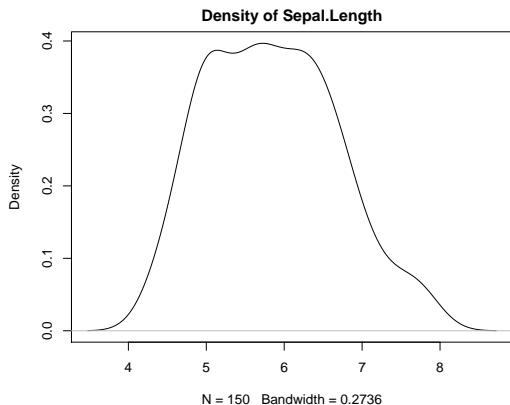
```
var(iris$Sepal.Length)
## [1] 0.6856935

hist(iris$Sepal.Length)
```





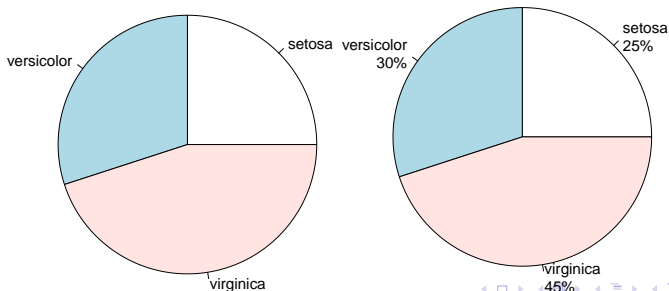
```
library(magrittr) ## for pipe operations  
iris$Sepal.Length %>% density() %>% plot(main='Density of Sepal.Length
```



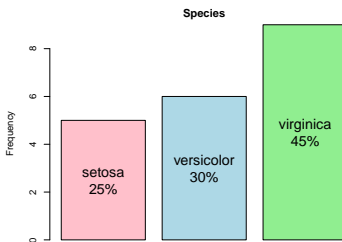
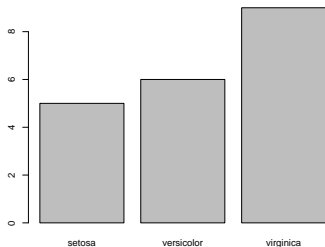
# Pie Chart

Frequency of factors: `table()`

```
library(dplyr)
set.seed(123)
iris2 <- iris %>% sample_n(20)
iris2$Species %>% table() %>% pie()
# add percentages
tab <- iris2$Species %>% table()
percentages <- tab %>% prop.table() %>% round(3) * 100
txt <- paste0(names(tab), '\n', percentages, '%')
pie(tab, labels=txt)
```



```
iris2$Species %>% table() %>% barplot()  
# add colors and percentages  
bb <- iris2$Species %>% table() %>%  
  barplot(axisnames=F, main='Species', ylab='Frequency',  
          col=c('pink', 'lightblue', 'lightgreen'))  
text(bb, tab/2, labels=txt, cex=1.5)
```



Introduction

Have a Look at Data

Explore Individual Variables

**Explore Multiple Variables**

More Explorations

Save Charts to Files

Further Readings and Online Resources

Covariance and correlation: `cov()` and `cor()`

```
cov(iris$Sepal.Length, iris$Petal.Length)
```

```
## [1] 1.274315
```

```
cor(iris$Sepal.Length, iris$Petal.Length)
```

```
## [1] 0.8717538
```

```
cov(iris[,1:4])
```

```
##           Sepal.Length Sepal.Width Petal.Length Petal.Width
## Sepal.Length    0.6856935  -0.0424340    1.2743154    0.5162707
## Sepal.Width     -0.0424340   0.1899794   -0.3296564   -0.1216394
## Petal.Length     1.2743154  -0.3296564    3.1162779    1.2956094
## Petal.Width      0.5162707  -0.1216394    1.2956094    0.5810063
```

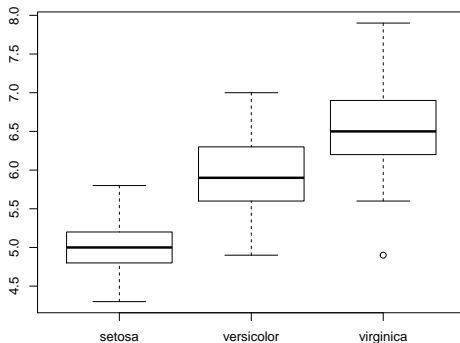
```
# cor(iris[,1:4])
```

Stats of Sepal.Length for every Species with aggregate()

```
aggregate(Sepal.Length ~ Species, summary, data=iris)
##      Species Sepal.Length.Min. Sepal.Length.1st Qu.
## 1      setosa           4.300           4.800
## 2 versicolor           4.900           5.600
## 3  virginica           4.900           6.225
##      Sepal.Length.Median Sepal.Length.Mean Sepal.Length.3rd Qu.
## 1              5.000           5.006           5.200
## 2              5.900           5.936           6.300
## 3              6.500           6.588           6.900
##      Sepal.Length.Max.
## 1              5.800
## 2              7.000
## 3              7.900
```

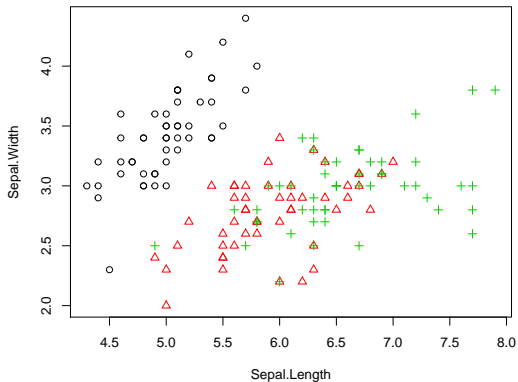
- ▶ The bar in the middle is median.
- ▶ The box shows the interquartile range (IQR), i.e., range between the 75% and 25% observation.

```
boxplot(Sepal.Length ~ Species, data=iris)
```



# Scatter Plot

```
with(iris, plot(Sepal.Length, Sepal.Width, col = Species,  
               pch = as.numeric(Species)))
```

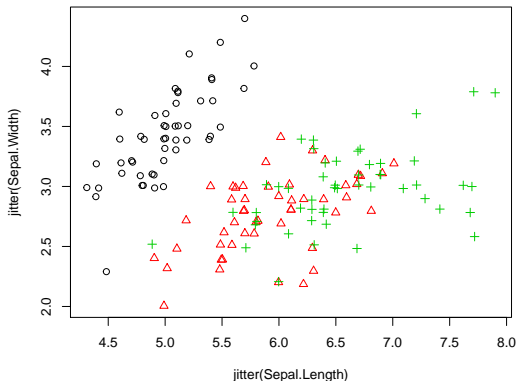




# Scatter Plot with Jitter

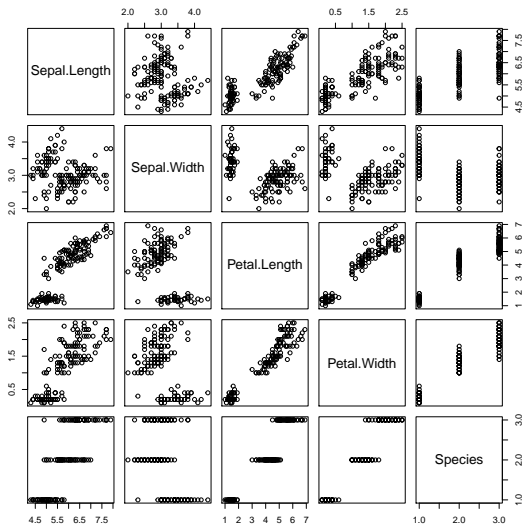
Function `jitter()`: add a small amount of noise to the data

```
with(iris, plot(jitter(Sepal.Length), jitter(Sepal.Width), col=Species,  
               pch=as.numeric(Species)))
```



# A Matrix of Scatter Plots

```
pairs(iris)
```



Introduction

Have a Look at Data

Explore Individual Variables

Explore Multiple Variables

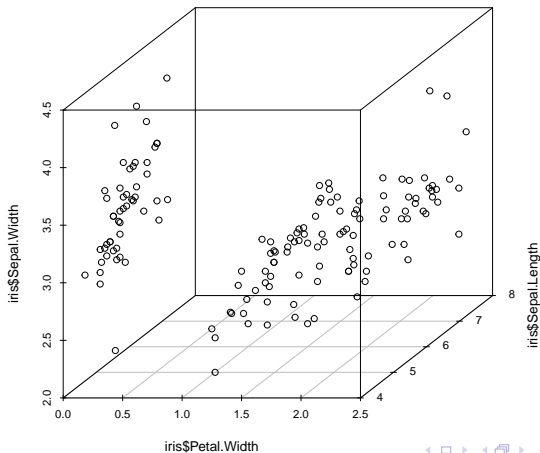
**More Explorations**

Save Charts to Files

Further Readings and Online Resources

# 3D Scatter plot

```
library(scatterplot3d)  
scatterplot3d(iris$Petal.Width, iris$Sepal.Length, iris$Sepal.Width)
```



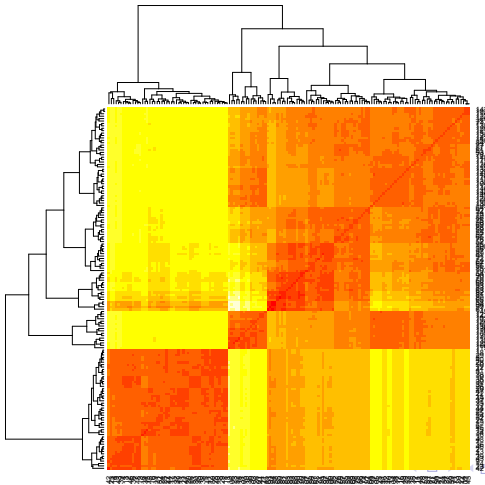
Package *rgl* supports interactive 3D scatter plot with `plot3d()`.

```
library(rgl)
plot3d(iris$Petal.Width, iris$Sepal.Length, iris$Sepal.Width)
```

# Heat Map

Calculate the similarity between different flowers in the iris data with `dist()` and then plot it with a heat map

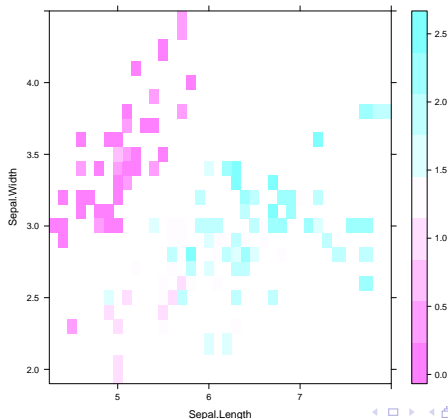
```
dist.matrix <- as.matrix(dist(iris[,1:4]))  
heatmap(dist.matrix)
```



# Level Plot

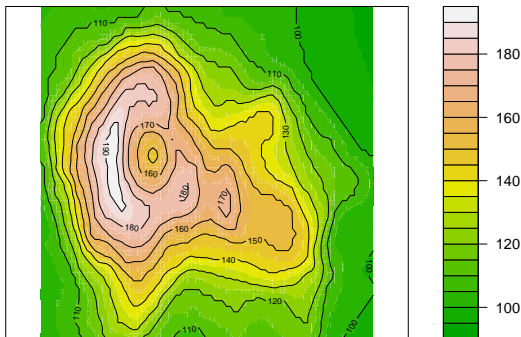
Function `rainbow()` creates a vector of contiguous colors.  
`rev()` reverses a vector.

```
library(lattice)
levelplot(Petal.Width ~ Sepal.Length * Sepal.Width,
          data=iris, cuts=8)
```



`contour()` and `filled.contour()` in package *graphics*  
`contourplot()` in package *lattice*

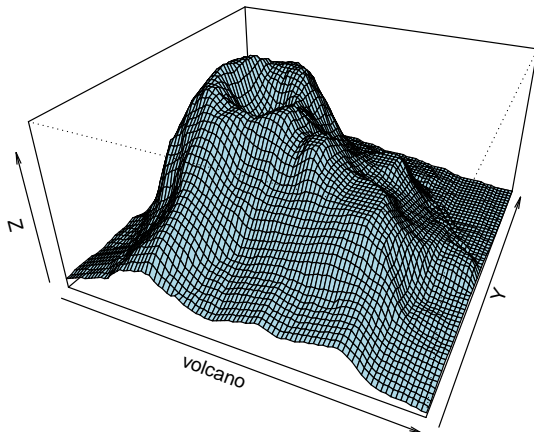
```
filled.contour(volcano, color=terrain.colors, asp=1,  
               plot.axes=contour(volcano, add=T))
```





# 3D Surface

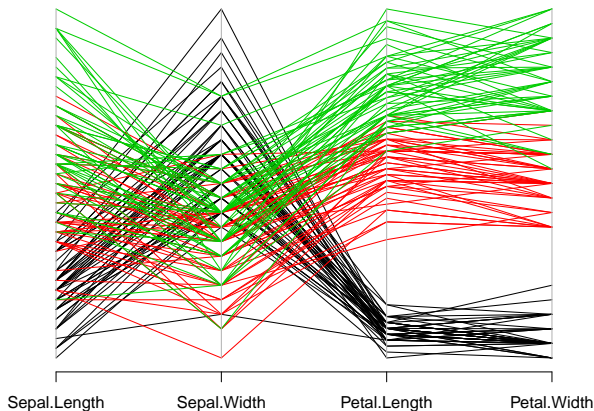
```
persp(volcano, theta=25, phi=30, expand=0.5, col="lightblue")
```



# Parallel Coordinates

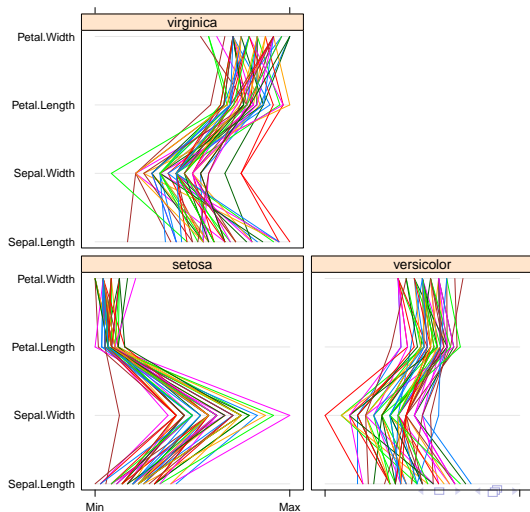
Visualising multiple dimensions

```
library(MASS)
parcoord(iris[1:4], col=iris$Species)
```



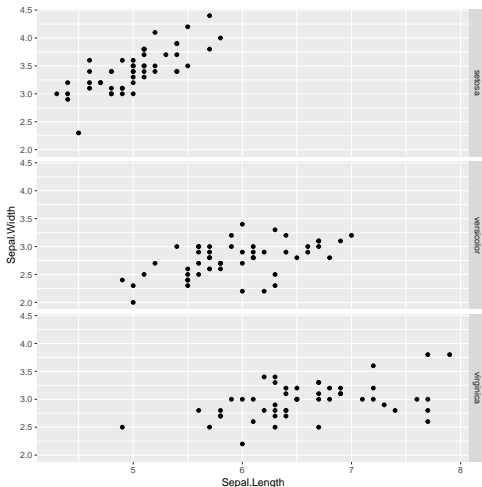
# Parallel Coordinates with Package *lattice*

```
library(lattice)
parallelplot(~iris[1:4] | Species, data=iris)
```



# Visualisation with Package *ggplot2*

```
library(ggplot2)
qplot(Sepal.Length, Sepal.Width, data=iris, facets=Species ~.)
```



Introduction

Have a Look at Data

Explore Individual Variables

Explore Multiple Variables

More Explorations

Save Charts to Files

Further Readings and Online Resources

- ▶ Save charts to PDF and PS files: `pdf()` and `postscript()`
- ▶ BMP, JPEG, PNG and TIFF files: `bmp()`, `jpeg()`, `png()` and `tiff()`
- ▶ Close files (or graphics devices) with `graphics.off()` or `dev.off()` after plotting

```
# save as a PDF file
pdf("myPlot.pdf")
x <- 1:50
plot(x, log(x))
graphics.off()
#
# Save as a postscript file
postscript("myPlot2.ps")
x <- -20:20
plot(x, x^2)
graphics.off()
```

`ggsave()`: by default, saving the last plot that you displayed. It also guesses the type of graphics device from the extension.

```
ggsave('myPlot3.png')  
ggsave('myPlot4.pdf')  
ggsave('myPlot5.jpg')  
ggsave('myPlot6.bmp')  
ggsave('myPlot7.ps')  
ggsave('myPlot8.eps')
```

Introduction

Have a Look at Data

Explore Individual Variables

Explore Multiple Variables

More Explorations

Save Charts to Files

Further Readings and Online Resources



- ▶ Examples of ggplot2 plotting:  
<https://ggplot2.tidyverse.org/>
- ▶ Package *iplots*: interactive scatter plot, histogram, bar plot, and parallel coordinates plot (iplots)  
<http://rosuda.org/software/iPlots/>
- ▶ Package *googleVis*: interactive charts with the Google Visualisation API  
[http://cran.r-project.org/web/packages/googleVis/vignettes/googleVis\\_examples.html](http://cran.r-project.org/web/packages/googleVis/vignettes/googleVis_examples.html)
- ▶ Package *ggvis*: interactive grammar of graphics  
<http://ggvis.rstudio.com/>
- ▶ Package *rCharts*: interactive javascript visualisations from R  
<https://ramnathv.github.io/rCharts/>

- ▶ Chapter 3: Data Exploration, in book *R and Data Mining: Examples and Case Studies*

<http://www.rdatamining.com/docs/RDataMining-book.pdf>

- ▶ R Reference Card for Data Mining

<http://www.rdatamining.com/docs/RDataMining-reference-card.pdf>

- ▶ Free online courses and documents

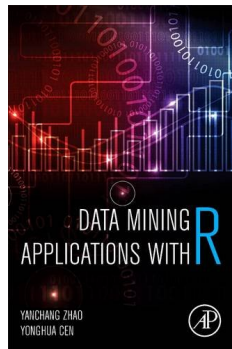
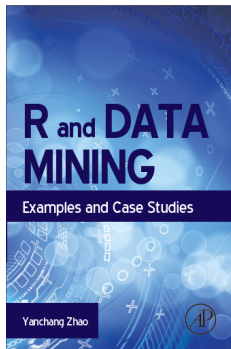
<http://www.rdatamining.com/resources/>

- ▶ RDataMining Group on LinkedIn (26,000+ members)

<http://group.rdatamining.com>

- ▶ Twitter (3,300+ followers)

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Thanks!

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Twitter: @RDataMining



Frank, A. and Asuncion, A. (2010).

UCI machine learning repository. university of california, irvine, school of information and computer sciences.  
<http://archive.ics.uci.edu/ml>.