Working with R

transition: rotate



Lecture 6

What is R?

R is a free software environment for statistical computing and graphics.

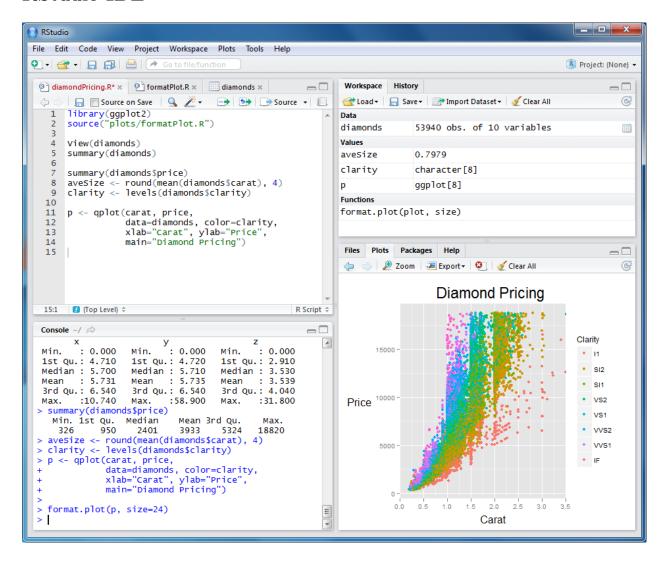
History

- 1976: In Bell Labs S language was created
- 1988: Commercial version S-PLUS appeared
- 1993: Fork R appeared
- ~2000: R getting more and more popular

Pure R

```
R version 3.0.2 (2013-09-25) -- "Frisbee Sailing"
Copyright (C) 2013 The R Foundation for Statistical Computing
Platform: x86_64-apple-darwin10.8.0 (64-bit)
R -- это свободное ПО, и оно поставляется безо всяких гарантий.
Вы вольны распространять его при соблюдении некоторых условий.
Введите 'license()' для получения более подробной информации.
R -- это проект, в котором сотрудничает множество разработчиков.
Введите 'contributors()' для получения дополнительной информации и
'citation()' для ознакомления с правилами упоминания R и его пакетов
в публикациях.
Введите 'demo()' для запуска демонстрационных программ, 'help()' -- для
получения справки, 'help.start()' -- для доступа к справке через браузер.
Введите 'q()', чтобы выйти из R.
> head(iris)
  Sepal.Length Sepal.Width Petal.Length Petal.Width Species
                       3.5
                                    1.4
           5.1
                                                 0.2 setosa
2
3
4
5
6
           4.9
                                    1.4
                       3.0
                                                 0.2 setosa
                                                 0.2 setosa
           4.7
                       3.2
                       3.1
                                                 0.2 setosa
           4.6
                                    1.5
                       3.6
                                                 0.2 setosa
           5.4
                       3.9
                                    1.7
                                                 0.4 setosa
```

RStudio IDE



CRAN & packages

Free software "tradition":

- CTAN: Comprehensive TeX archive network
- CRAN: Comprehensive R archive network
- ...

Another sources:

- Bioconductor
- GitHub

Main IDE frames

Editor

Create and edit scripts

In regular R: separate editors could be used.

Ctrl + Enter to execute selected part

Console

Main window:

```
> a <- 5 * 5
```

Environment

All variables and types

+ Viewer

Help & plots

In regular R:

- help inside console
- plots in separate windows

Primitives

```
# help on function
?lm

# math functions
2 + 3
10 / 3
sqrt(9)

# assigning
a <- 'Hello, world!'</pre>
```

Adding packages

```
install.packages("ggplot2")
library(ggplot2)
require(ggplot2)
```

Working directory

```
getwd()

[1] "/Users/quatsch/Documents/RIA_lectures"

# Set required WD
setwd('C:/Documents/my_R_project')

# Show files in current WD
dir()
```

Loading tables

View raw file
file.show()

```
read.table(fname)
read.csv(fname)
read.csv2(fname)
read.delim(fname)
read.delim2(fname)
require(xlsx)
read.xlsx(fname, sheetName)
```

Writing files

```
write.table
write.csv
write.csv2
```

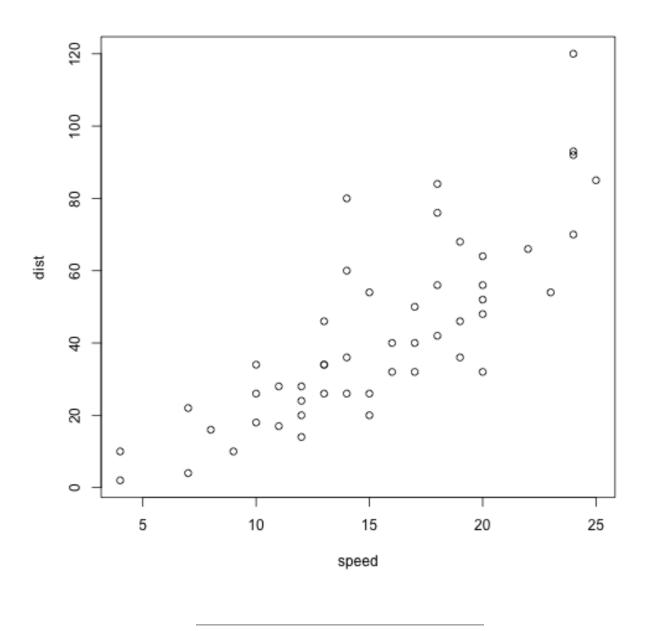
Demo data

```
head(cars, 5)
```

```
iris
mtcars
Titanic
# etc...
```

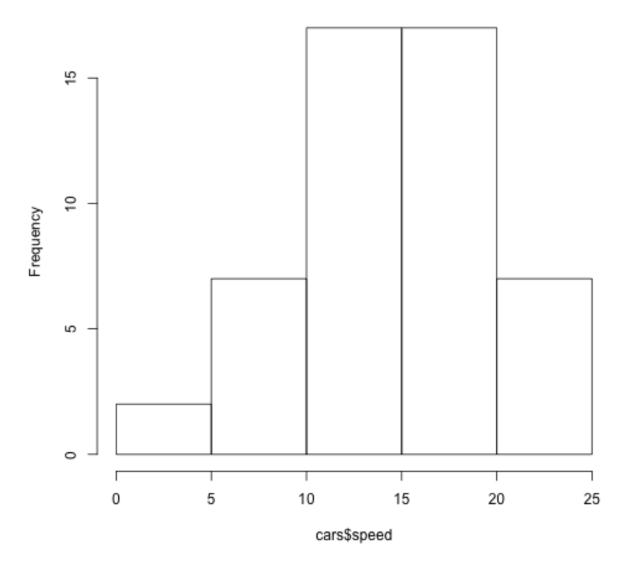
First graphic

```
plot(cars)
```



hist(cars\$speed)

Histogram of cars\$speed



Basic data types

- Numeric 10.5 / Integer 10 / Complex $1\ +\ 2i$
- Factor

```
factor(c('m', 'v', 'm', 'v', 'v'), levels = c('m', 'v'))
```

```
[1] m v m v v Levels: m v
```

- logical True (T) / False (F)
- character

Missing & special data

```
# NA
vec <- c(3, 10, 8, NA, 5, 6)
mean(vec)

[1] NA
mean(vec, na.rm=T)

[1] 6.4
```

Missing & special data

```
# Additional: NaN, Inf, and -Inf
pi/0

[1] Inf

0/0

[1] NaN

as.logical(0/0)
[1] NA
```

Converting

```
vec <- factor(c('1982', '1983', '1982', '1984', '1985'))

# Wrong
as.numeric(vec)

[1] 1 2 1 3 4

# Right
as.numeric(as.character(vec))

[1] 1982 1983 1982 1984 1985</pre>
```

Sequences - Vector

```
# concatenate
vec <- c(TRUE, 1, 0.5, 'item')

# vector elements should be of the same class
class(vec)

[1] "character"

# slicing
vec[3:4]

[1] "0.5" "item"</pre>
```

Sequences - Matrix

```
v <- 1:9
v

[1] 1 2 3 4 5 6 7 8 9

dim(v) <- c(3,3)
v

[,1] [,2] [,3]
[1,] 1 4 7
[2,] 2 5 8
[3,] 3 6 9
```

Sequences - List

```
test.list <- list(i1 = TRUE, i2 = 1, i3 = 0.5, i4 = 'item')
# every element is independent
class(test.list)

[1] "list"
# slicing
test.list[4]</pre>
$i4
[1] "item"
```

```
test.list$i4
```

[1] "item"

Sequences - Data Frame

Imagine list of multiple vectors (columns):

```
var1 var2 var3

f 1982 TRUE

m 1982 FALSE

m 1983 FALSE

f 1985 FALSE
```

Data Frames Slicing

```
nrow(iris)

[1] 150

iris.filtered <- iris[iris$Species == 'setosa', ]
nrow(iris.filtered)

[1] 50

head(iris[, 'Species'])

[1] setosa setosa setosa setosa setosa setosa</pre>
```

Vectorized computations

Levels: setosa versicolor virginica

```
    var1
    var2
    var3
    var4

    1
    1
    10
    11
    10

    2
    2
    20
    22
    40

    3
    3
    33
    90

    4
    4
    40
    44
    160

    5
    5
    55
    250

    6
    6
    60
    66
    360
```

apply VS for

```
#head(trees, 3)
apply(trees, 2, mean)

Girth Height Volume
13.25 76.00 30.17

for(row in names(trees)){
   print(paste(row, mean(trees[, row])))
}

[1] "Girth 13.2483870967742"
[1] "Height 76"
[1] "Volume 30.1709677419355"
```

Basic functions for stat analysis

```
mean()
median()
sd()
var() # sd() ^ 2
```

Your own functions

```
square <- function(vec)
{
  vec <- vec ^ 2
  return(vec)
}

my.vec <- c(1,2,3,4,5)
square(my.vec)</pre>
```

```
[1] 1 4 9 16 25
```

The End

type: sub-section

Questions & answers

http://www.r-tutor.com/elementary-statistics/qualitative-data/frequency-distribution-qualitative-data/frequency-data/fr