

R Programming Language

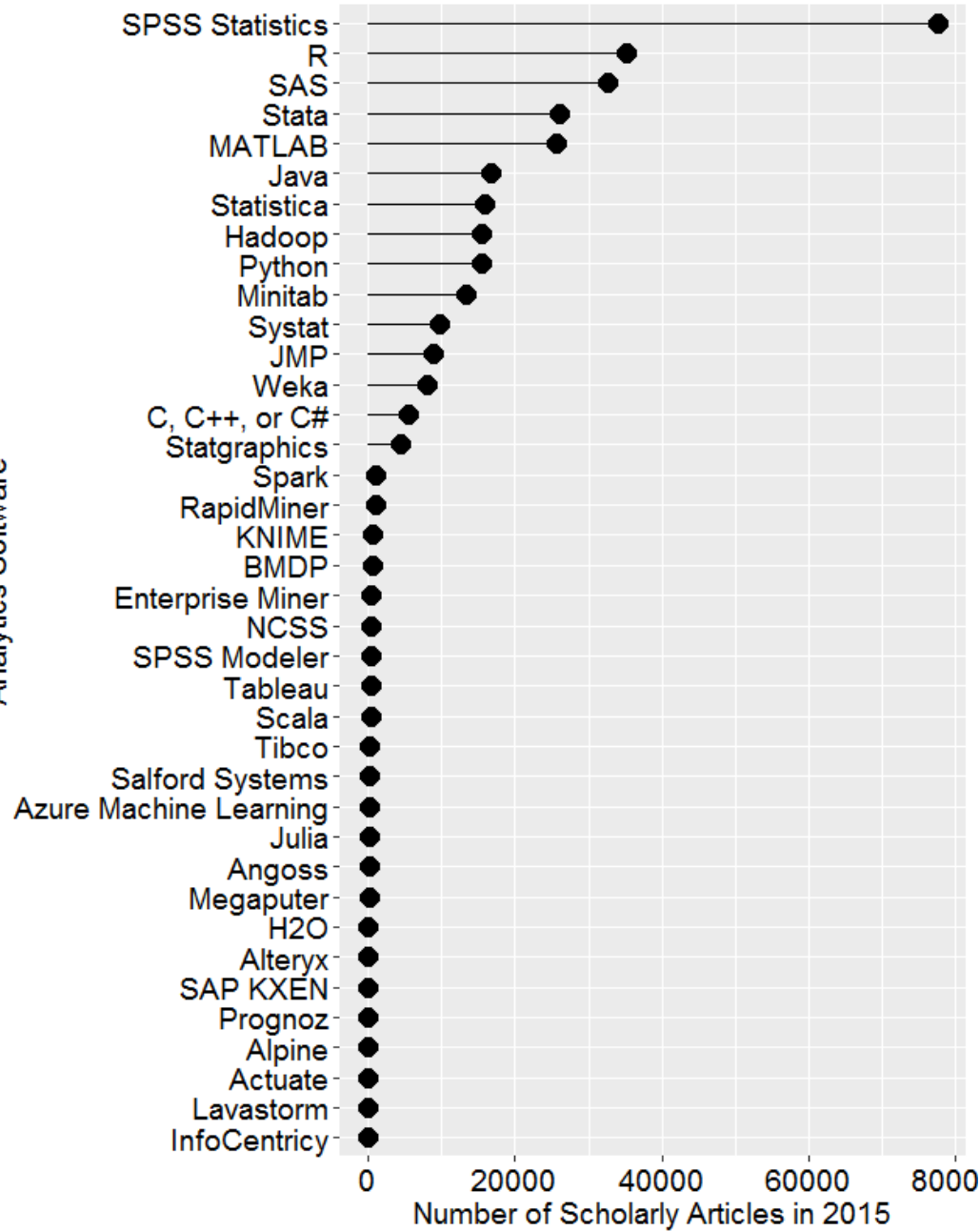
Objectives of the workshop

- Introduction to R programming language
- Import data to Rstudio
- Summary measures
 - Mean, median, standard deviation, Centiles, N, min, max
- Visualize data
 - Bar plot, Histogram, Box plot, Scatter plot
- Hypothesis testing
 - T test
 - Pearson's Chisqaure test

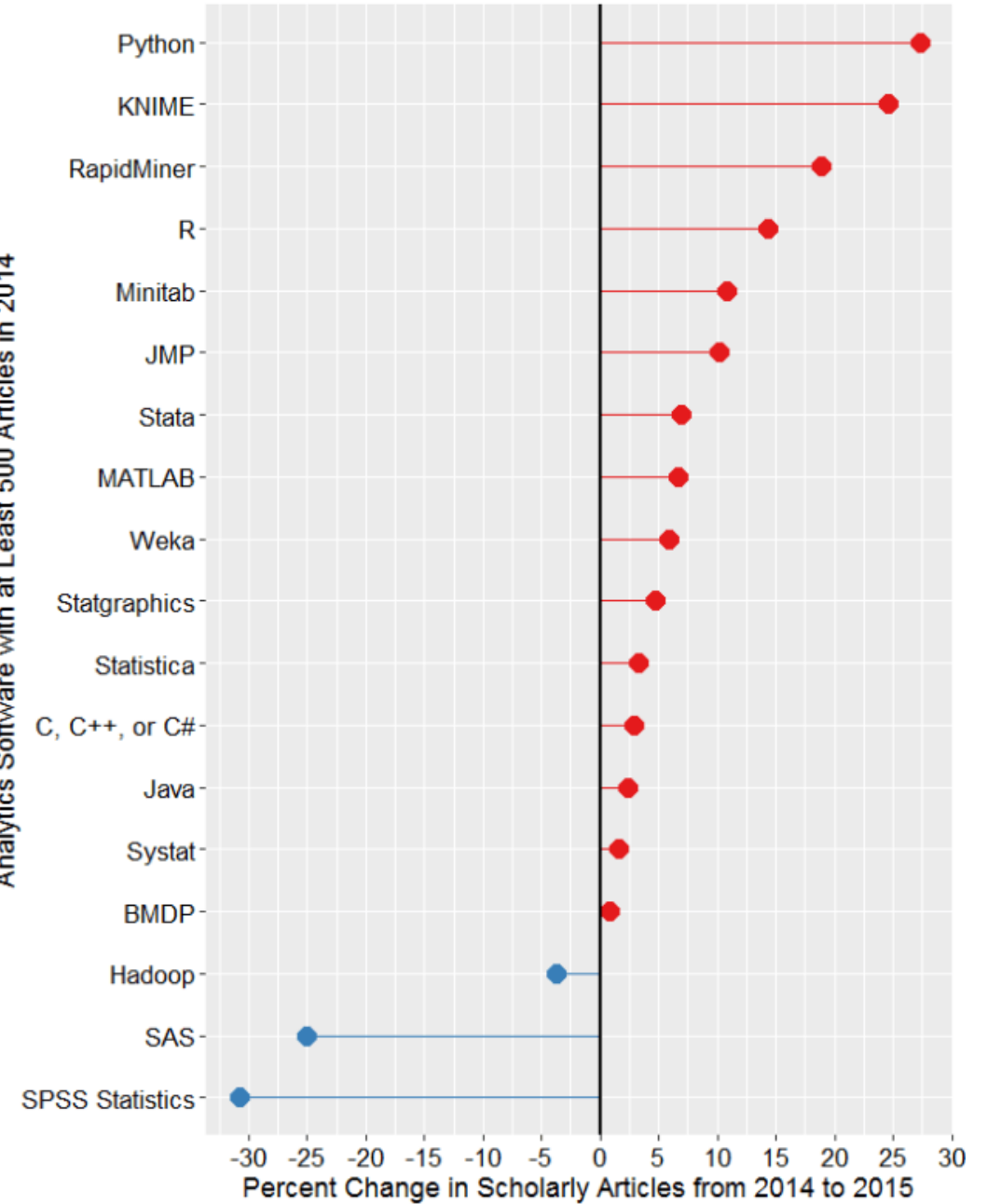
R Programming Language

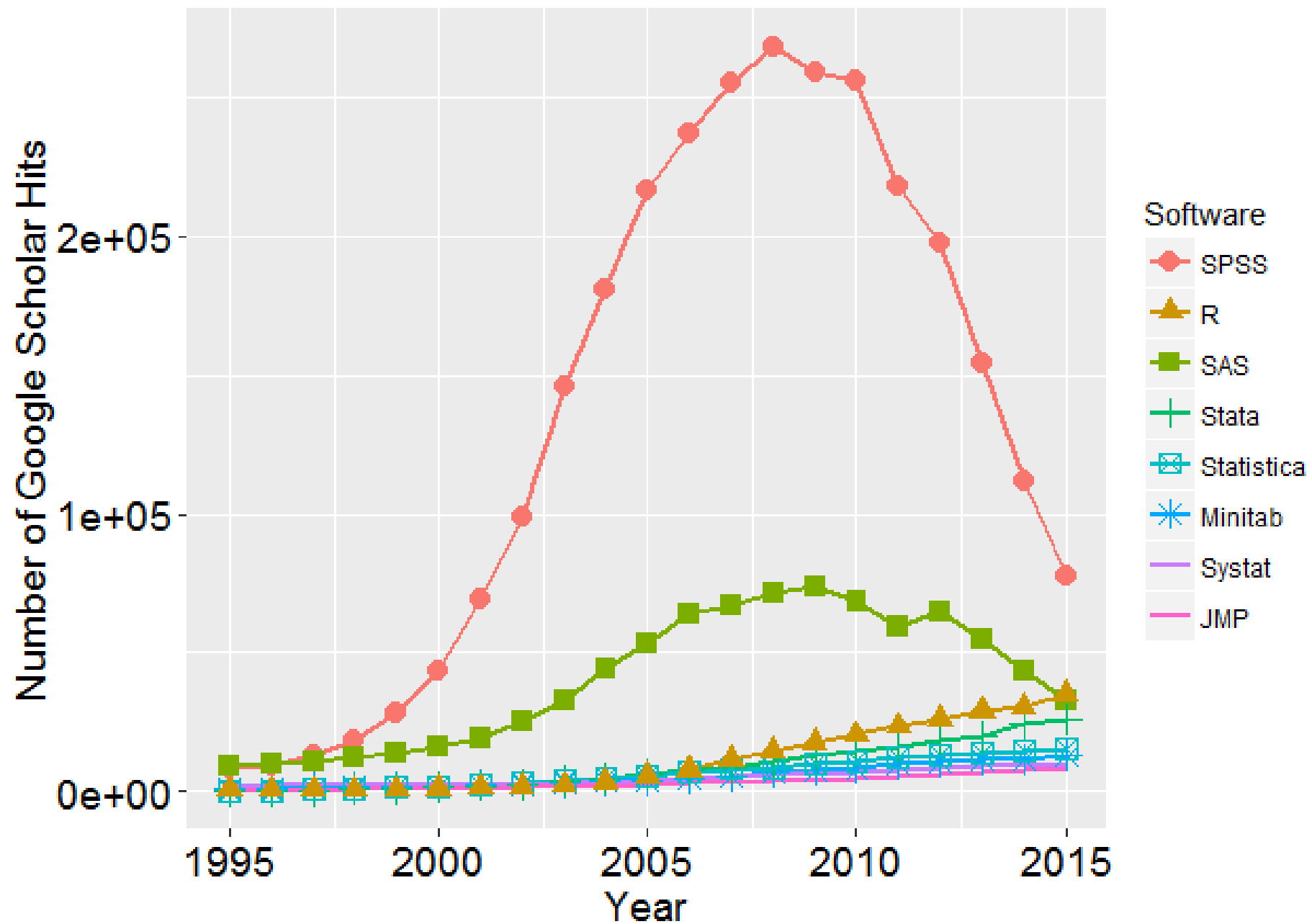
- R is a **free software** environment for statistical computing and graphics (Ref: <https://www.r-project.org/>)
- It compiles and runs on a wide variety of UNIX platforms, Windows and MacOS.
- R was created by **Ross Ihaka** and **Robert Gentleman** at the University of Auckland, New Zealand
- R is an implementation of the **S programming language**
 - by John Chambers while at Bell Labs
 - much of the code written for S runs unaltered

Analytics Software



Analytics Software with at Least 500 Articles in 2014



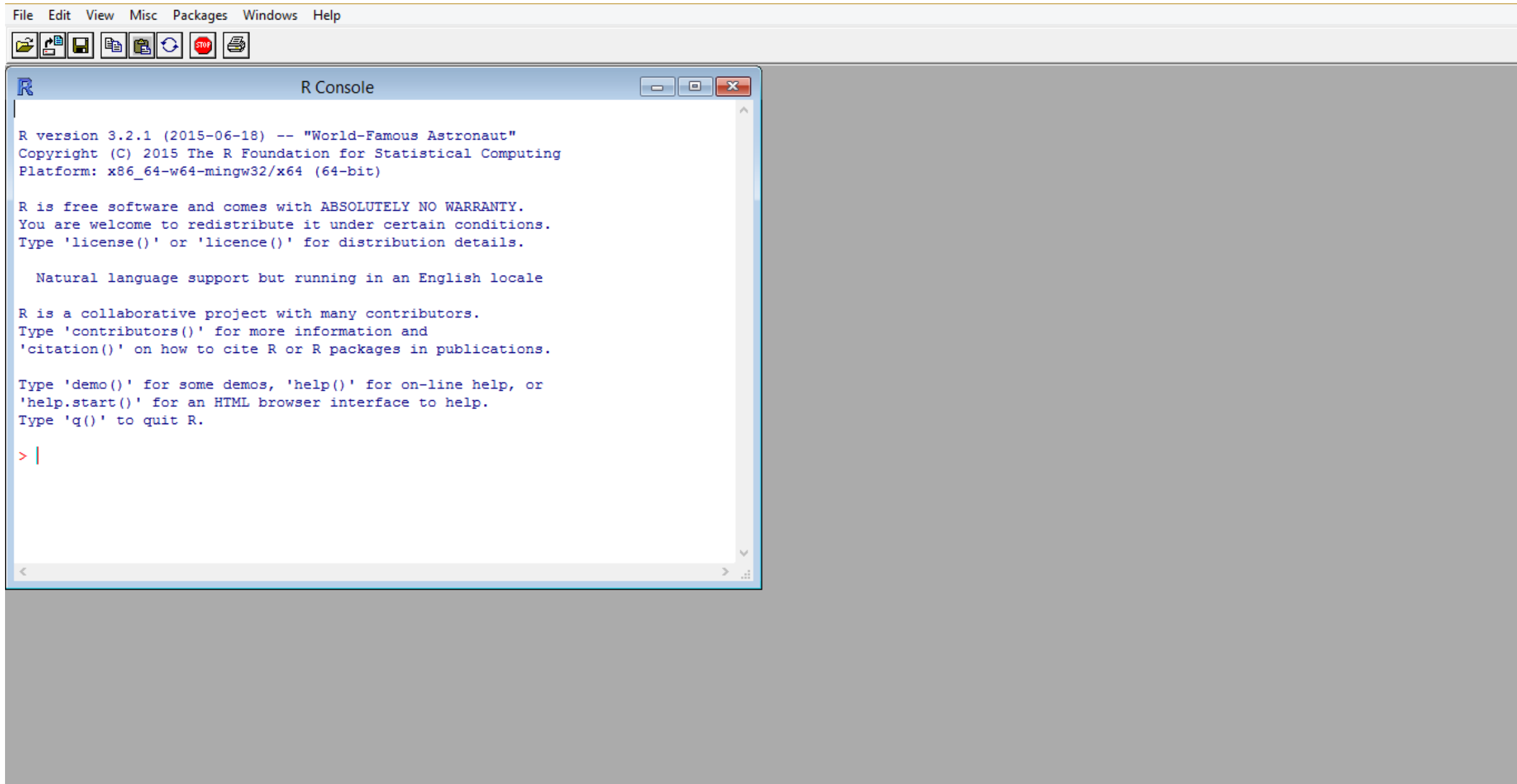


Tutorials

- P. Kuhnert & B. Venables, *An Introduction to R: Software for Statistical Modeling & Computing*
- J.H. Maindonald, *Using R for Data Analysis and Graphics*
- B. Muenchen, *R for SAS and SPSS Users*
- W.J. Owen, *The R Guide*
- D. Rossiter, *Introduction to the R Project for Statistical Computing for Use at the ITC*
- W.N. Venables & D. M. Smith, *An Introduction to R*

Data analysis with R...





RStudio

- RStudio is a **free** and open source integrated development environment (IDE) for R
- RStudio is available in two editions:
 - RStudio Desktop, where the program is run locally as a regular desktop application
 - RStudio Server, which allows accessing RStudio using a web browser while it is running on a remote Linux server.

C:/D-Datos/Göttingen/Papers/LiDAR variables selection Edu/Box-Cox - RStudio

File Edit Code View Plots Session Build Debug Tools Help

Go to file/function

01-Introduction.Rnw 02-SRS.Rnw Data analysis Kalimantan.R

Source on Save Run Source

```
200
201 # Biomass calculation per tree
202 kalimantan$w.brown<-brown.moist.d(kalimantan$dbh)
203 kalimantan$w.yamakura<-yamakura.stem(kalimantan$dbh, kalimantan$h)+yamakura.branch(yamakura.stem(k
204 kalimantan$w.basuki<-basuki.mixed.d(kalimantan$dbh)
205 kalimantan$w.samalca<-samalca.d(kalimantan$dbh)
206 kalimantan$w.hashimoto<-hashimoto.d(kalimantan$dbh)
207 kalimantan$w.kenzo<-kenzo.d(kalimantan$dbh)
208 kalimantan$w.forda<-forda.d(kalimantan$dbh)
209 kalimantan$w.jaya<-jaya.d(kalimantan$dbh)
210 kalimantan$w.novita<-novita.d(kalimantan$dbh)
211 kalimantan$w.nugroho.d<-nugroho.d(kalimantan$dbh)
212 kalimantan$w.nugroho.d.h<
213
214 plot(kalimantan$dbh, kalimantan$w.brown, col="brown", xlab="DBH", ylab="Biomass", main="Biomass estimation per plot with different models", xlab="DBH
215 points(kalimantan$dbh, kalimantan$w.yamakura, col="darkred", xlab="DBH", ylab="Biomass", main="Biomass estimation per plot with different models", xlab="DBH
216 points(kalimantan$dbh, kalimantan$w.basuki, col="darkgreen", xlab="DBH", ylab="Biomass", main="Biomass estimation per plot with different models", xlab="DBH
217 points(kalimantan$dbh, kalimantan$w.samalca, col="darkblue", xlab="DBH", ylab="Biomass", main="Biomass estimation per plot with different models", xlab="DBH
218 points(kalimantan$dbh, kalimantan$w.hashimoto, col=5)
219 points(kalimantan$dbh, kalimantan$w.kenzo, col=6)
220 points(kalimantan$dbh, kalimantan$w.forda, col=7)
221 points(kalimantan$dbh, kalimantan$w.jaya, col=8)
222 points(kalimantan$dbh, kalimantan$w.novita, col=9)
223 points(kalimantan$dbh, kalimantan$w.nugroho.d, col=10)
224 points(kalimantan$dbh, kalimantan$w.nugroho.d.h, col=11)
225
226 legend(10,8000, c("Brown", "Yamakura", "Basuki", "Samalca", "Hashimoto", "Kenzo", "Forda", "Jaya",
227
228
229 # summing all values per plot and nested plot
230 bio.plot.brown<-as.data.frame(tapply(kalimantan$w.brown, list(kalimantan$plot_id, kalimantan$subplot_id), FUN=sum))
231
```

310:1 (Untitled) R Script

Console Compile PDF

```
C:/D-Datos/Göttingen/Indonesia Projects/Kalimantan Project/Final Data/
> kal.plot<-merge(kal.plot, Dmed.Hmed.plot, by="Plot")
>
> # calculating the
> kal.plot$dg<-sqrt((4*kal.plot$
>
> write.csv(kal.plot, "kalimanta
>
```

Environment History

Global Environment

- hil.trees 716 obs. of 23 variables
- kal.plot 94 obs. of 18 variables
- kalimantan 1993 obs. of 44 variables
- lsi.plots 59 obs. of 19 variables
- lsi
- pub
- wet

valu

EFT	12.4339739290343
EFm	49.7359197162173
EFS	198.943678864869
N.tot	2696.5863280181

Files Plots Packages Help Viewer

Zoom Export Clear All

Biomass estimation per plot with different models

R script

R console

Graphical output

R basics

- R is a case sensitive language
 - FOO, Foo, and foo are three different objects
- Results of calculations can be stored in objects using the assignment operators:
 - An arrow (<-) formed by a smaller than character and a hyphen without a space!
 - The equal character (=)
- Object names
 - cannot contain 'strange' symbols like !, +, -, #
 - dot (.) and an underscore () are allowed
 - can contain a number but cannot start with a number

Types of brackets used in R

- Round brackets / Parenthesis ()
 - For functions
 - `mean()`, `sd()`, `max()`
- Square brackets
 - For manipulating data
 - `Data[1,1]`
 - `Data[1,]`
 - `Data[,1]`

Data frames

- Way of storing data with different types
- Data can be of different types
- There are different ways to create and manipulate data frames
 - Create
 - Import data

Comma-separated values (CSV) file

- Common data exchange format
 - Widely supported by software applications.
 - Move tabular data between programs (SPSS, SAS, etc)
- Use commas as field separators
- Interpreted as a sequence of characters
 - human-readable by text editor
 - Each line - data record

Selecting data

- Data frame
 - Elements
 - Rows
 - Columns

17 observations of 9 variables

	YEAR	CBE	PPO	CPO	PFO	DINC	CFO	RDINC	RFP
1	1925	58.6	60.5	65.8	65.8	51.4	90.9	68.5	877
2	1926	59.4	63.3	63.3	68.0	52.6	92.1	69.6	899
3	1927	53.7	59.9	66.8	65.5	52.1	90.9	70.2	883
4	1928	48.1	56.3	69.9	64.8	52.7	90.9	71.9	884
5	1929	49.0	55.0	68.7	65.6	55.1	91.1	75.2	895
6	1930	48.2	59.6	66.1	62.4	48.8	90.7	68.3	874
7	1931	47.9	57.0	67.4	51.4	41.5	90.0	64.0	791
8	1932	46.0	49.5	69.7	42.8	31.4	87.8	53.9	733
9	1933	50.8	47.3	68.7	41.6	29.4	88.0	53.2	752
10	1934	55.2	56.6	62.2	46.4	33.2	89.1	58.0	811
11	1935	52.2	73.9	47.7	49.7	37.0	87.3	63.2	847
12	1936	57.3	64.4	54.4	50.1	41.8	90.5	70.5	845
13	1937	54.4	62.2	55.0	52.1	44.5	90.4	72.5	849
14	1938	53.6	59.9	57.4	48.4	40.8	90.6	67.8	803
15	1939	53.9	51.0	63.9	47.1	43.5	93.8	73.2	793
16	1940	54.2	41.5	72.4	47.8	46.5	95.5	77.6	798
17	1941	60.0	43.9	67.4	52.2	56.3	97.5	89.5	830

country	year	cases	population
Afghanistan	1999	745	19987071
Afghanistan	2000	2666	20595360
Brazil	1999	37737	172006362
Brazil	2000	80488	174504898
China	1999	212258	1272915272
China	2000	213766	1280425583

variables

country	year	cases	population
Afghanistan	1999	745	19987071
Afghanistan	2000	2666	20595360
Brazil	1999	37737	172006362
Brazil	2000	80488	174504898
China	1999	212258	1272915272
China	2000	213766	1280425583

observations

country	year	cases	population
Afghanistan	1999	745	19987071
Afghanistan	2000	2666	20595360
Brazil	1999	37737	172006362
Brazil	2000	80488	174504898
China	1999	212258	1272915272
China	2000	213766	1280425583

values

Summary measures

- Mean - `mean(variable)`
- Standard deviation - `sd(variable)`
- Quantiles - `quantile(variable, c(.32, .57, .98))`
- Number of elements - `length()`
- Minimum - `min()`
- Maximum - `max()`
- Tabulate – `table()`
- Missing values – `is.na()`

Graphs

- Histogram - `hist()`
- Box plot - `boxplot()`
- Scatter plot - `plot(x,y)`
- Stem & leaf - `stem()`
- Pie chart - `pie()`
- Bar plot - `barplot()`

R Help

- `help.start()`
- `help(mean)` or `? Mean`
- `RSiteSearch("mean")`
- `apropos("mean")`
- `example(mean)`
- `data()`

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function

Addins

Prediction_2011_2013.R × sample_size.R × v1_data.R × Mixed_effect_models.R × Untitled1* ×

Source on Save Run Source

```
1 # Load required library
2 library(lme4)
3
4 # Sample data (replace with your actual data)
5 y <- rnorm(100) # Level 1 outcome variable
6 group1 <- gl(10, 10) # Grouping factor (10 groups with 10 observations each)
7 group2 <- gl(4, 25) # Grouping factor (4 groups with 25 observations each)
8 x1 <- rnorm(100) # Level 1 predictor
9 x2 <- rnorm(100) # Level 1 predictor
10
11 # Define the model formula (specifies diagonal structure)
12 model <- lmer(y ~ x1 + (1 | group1) , REML = FALSE)
13 summary(model)
14
15 model <- lmer(y ~ x1 + x2 + (1 | group1) , REML = FALSE)
16 summary(model)
17
18 model <- lmer(y ~ x1 + x2 + (1 | group1) + (1 | group2) , REML = FALSE)
19 summary(model)
20
21 ?var.test
22
23 x <- rnorm(50, mean = 0, sd = 2)
24 y <- rnorm(30, mean = 1, sd = 1)
25 var.test(x, y) # Do x and y have the same variance?
26 var.test(lm(x ~ 1), lm(y ~ 1)) # The same.
27
```

Saved to this PC

Console Terminal Background Jobs

R 4.3.3 · ~/

> help.start()
If nothing happens, you should open
'http://127.0.0.1:31820/doc/html/index.html' yourself
> |

Files Plots Packages Help Viewer Presentation

The R Language Find in Topic

Statistical Data Analysis

Manuals

[An Introduction to R](#)
[Writing R Extensions](#)
[R Data Import/Export](#)

[The R Language Definition](#)
[R Installation and Administration](#)
[R Internals](#)

Reference

[Packages](#)

[Search Engine & Keywords](#)

Miscellaneous Material

[About R](#)
[License](#)
[NEWS](#)

[Authors](#)
[Frequently Asked Questions](#)
[User Manuals](#)

[Resources](#)
[Thanks](#)
[Technical papers](#)

Material specific to the Windows port

[CHANGES up to R 2.15.0](#)

[Windows FAQ](#)

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function

Addins

Prediction_2011_2013.R x sample_size.R x v1_data.R x Mixed_effect_models.R x Untitled1* x

Source on Save Run Source

```
1 # Load required library
2 library(lme4)
3
4 # Sample data (replace with your actual data)
5 y <- rnorm(100) # Level 1 outcome variable
6 group1 <- gl(10, 10) # Grouping factor (10 groups with 10 observations each)
7 group2 <- gl(4, 25) # Grouping factor (4 groups with 25 observations each)
8 x1 <- rnorm(100) # Level 1 predictor
9 x2 <- rnorm(100) # Level 1 predictor
10
11 # Define the model formula (specifies diagonal structure)
12 model <- lmer(y ~ x1 + (1 | group1) , REML = FALSE)
13 summary(model)
14
15 model <- lmer(y ~ x1 + x2 + (1 | group1) , REML = FALSE)
16 summary(model)
17
18 model <- lmer(y ~ x1 + x2 + (1 | group1) + (1 | group2) , REML = FALSE)
19 summary(model)
20
21 ?var.test
22
23 x <- rnorm(50, mean = 0, sd = 2)
24 y <- rnorm(30, mean = 1, sd = 1)
25 var.test(x, y) # Do x and y have the same variance?
26 var.test(lm(x ~ 1), lm(y ~ 1)) # The same.
27
```

Console

Files Plots Packages Help Viewer Presentation

An Introduction to R Find in Topic

An Introduction to R

This is an introduction to R (“GNU S”), a language and environment for statistical computing and graphics. R is similar to the award-winning¹ S system, which was developed at Bell Laboratories by John Chambers et al. It provides a wide variety of statistical and graphical techniques (linear and nonlinear modelling, statistical tests, time series analysis, classification, clustering, ...).

This manual provides information on data types, programming elements, statistical modelling and graphics.

This manual is for R, version 4.3.3 (2024-02-29).

Copyright © 1990 W. N. Venables
Copyright © 1992 W. N. Venables & D. M. Smith
Copyright © 1997 R. Gentleman & R. Ihaka
Copyright © 1997, 1998 M. Maechler
Copyright © 1999–2023 R Core Team

Permission is granted to make and distribute verbatim copies of this manual provided the copyright notice and this permission notice are preserved on all copies.

Permission is granted to copy and distribute modified versions of this manual under the conditions for verbatim copying, provided that the entire resulting derived work is distributed under the terms of a permission notice identical to this one.


Permission is granted to copy and distribute translations of this manual into another language, under the above conditions for modified versions, except that this permission notice may be stated in a translation approved by the R Core Team.

Table of Contents

[Preface](#)
[1 Introduction and preliminaries](#)
 [1.1 The R environment](#)
 [1.2 Related software and documentation](#)
 [1.3 R and statistics](#)
 [1.4 R and the window system](#)
 [1.5 Using R interactively](#)

26:30 (Top Level) R Script

Environment History Connections Tutorial

 Kelaniya
Medicine

Student's T test

- `help(t.test)`
- independent 2-group t-test
 - `t.test(y~x)` # where y is numeric and x is a binary factor
- independent 2-group t-test
 - `t.test(y1,y2)` # where y1 and y2 are numeric
- paired t-test
 - `t.test(y1,y2,paired=TRUE)` # where y1 & y2 are numeric
- one sample t-test
 - `t.test(y,mu=3)` # $H_0: \mu=3$

Compare Two Variances

- `var.test(y1,y2)`

Chisquared test

- `help(chisq.test)`
 - `chisq.test(table)`
 - `fisher.test (table)`

Simple Linear Regression

- $y = mx + c$
- `plot(x,y)`
- Formula
 - `model<- lm(y ~ x)`
 - `summary(model)`

Feedback

1. what did you like about this session?
2. what didn't you like about this session?
3. what did you learn from this session?

Thank you