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Getting Started

Mandy Vogel

May 31, 2015

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What's R?			

- R is a high-level language and an environment for data analysis and graphics
- influenced by S (Becker, Chamber, Wilks) and Scheme (Sussman)
- and created by Ross Ihaka and Robert Gentleman at the university of Auckland

- R is free.
- R is open source.
- R is a dialect of S system.

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What R can do			

 ${\sf R}$ provides a wide variety of statistical and graphical techniques including

- linear and nonlinear modelling
- classical statistical tests
- time-series analysis
- classification
- clustering and many more

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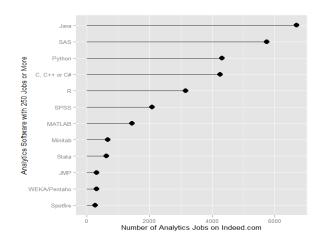
- linear and nonlinear modelling
- classical statistical tests
- time-series analysis
- classification
- clustering and many more

R is easily extensible, can produce publication-quality graphs including mathematical symbols; dynamic and interactive graphics are available through additional packages.

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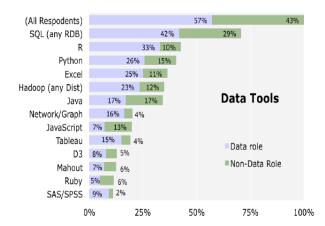
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Pros			

- R is free and R is open source
- there is a lot of material and books available
- there is a lot of help on the web, including developers who are active in mailing lists
- most of your problems are already solved and with a high probability the solution is available from one of the repositories (as package)
- there are a lot of intuitive GUIs
- the language is easy to learn and also intuitive
- the graphics capabilities are impressive



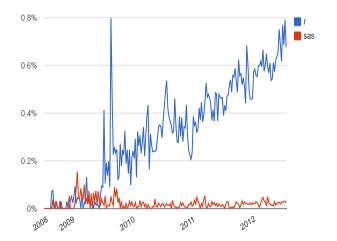
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Cons			

- there is a LOT of help on the web
- with a high probability there is more than one solution for your problem
- there are a lot of intuitive GUIs so you have to decide what you want (so first you have to *know* what you want)

- the real power of R (i.e. high flexibility) is not entirely available through GUIs
- and therefore the learning curve can be lengthy in the beginning (but soon accelerating ;)

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The best way to learn R is to use it!



Where can I get it?

For the basic installation CRAN is a good place to start

- CRAN stands for Comprehensive R Archive Network
- http://cran.r-project.org
 - Microsoft Windows: :: http://cran.r-project.org/bin/windows/base/
 - MacOS: :: http://cran.r-project.org/bin/macosx/
 - Linux: :: http://cran.r-project.org/bin/linux/
- for mac and pc users: just download and install the precompiled binaries
- for ubuntu users: add

```
deb http://ftp5.gwdg.de/pub/misc/cran/bin/linux/ubuntu
oneiric/
```

```
to
```

```
/etc/apt/sources.list;
```

detailed howto:

```
http://cran.r-project.org/bin/linux/ubuntu/
```

The R commander, developed by John Fox is a complete GUI for R. It is implemented in the package Rcmdr:

- Rcmdr has a comprehensive menu, which includes data reading, summaries, statistical analyses, etc.
- When the menu is activated, the Rcmdr will generate an R script. This script can be used as a log for documentation or for self learning.

• It has excellent graphical tools.

Packages

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The R-Commander

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•		
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> library(ggplot2) > data(Freatige) > library(relimp, pos-d)	nnati-1-20227, fast-getHands(1)agtas(1), manifelded), madatada	
> library(ggplot2) > data(Freatige) > library(relimp, pos-d)	nante" (Delle), fotogethade ('jogfot'), aantikkoli, madelyk	
> library(ggplot2) > data(Freatige) > library(relimp, pom-d)	nnan (1997), faisgettank (1997), faisgettakk, nakalikak, nakalikak	

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12.26	25879	4.02	69.1		prof
12.77					prof
11.42	8865	9.11	56.8		prof
	8403				prof
15.64					prof
15.09	8258		72.6		prof
15.44	14163		78.1	2141	
14.52	11377	1.03	73.1		prof
14.64	11023				prof
12.39	5902				prof
12.30	7059	7.83	60.0		prof
13.83	8425	15.33	53.8	2183	prof
14.44	8049	57.31	62.2		prof
14.36	7405	48.28	74.9		prof
14.21	6336	54.77	55.1	2331	prof
15,77	19263	5.13	82.3	2343	prof
14.15	6112	77.10	58.1	2351	prof
	9593			2391	prof
14.50		4.14		2511	prof
15.97					prof
13.62	5648	83.78	59.6	2731	prof
15.08	8034	46,80	66.1	2733	prof
15,96	25308	10,56	87.2	3111	prof
15,94	14558	4.32	66.7	3115	prof
14.71	17498	6.91	68.4		prof
12.46	4614	96.12	64.7	3131	prof
9,45	3485	76.14	34.9	3135	bc
13.62	5092	82.66	72.1	3137	prof
15.21	10432		69.3	3151	prof

Auswahl der aktiven Datenmatrix Aktualisiere aktive Datenmatrix Hilfe zur aktiven Datenmatrix (falls vorhanden) Variablen in aktiver Datenmatrix Fallbezeichnungen setzen ... Teilmenge der aktiven Datenmatrix ... Aggregate variables in active data set... Remove row(s) from active data set... Variablen übereinander plazieren ... Fälle mit fehlenden Werten entfernen ... Speichere aktive Datenmatrix ... Farbpalette ... Index-Plot Histogramm ... "Stamm und Blatt" Abbildung Boxplot ... Quantile-comparison plot... Streudiagramm ... Streudiagramm Matrix ... Liniengrafik ... XY conditioning plot... Plot für arithmetische Mittel Strip chart... Balkendiagramm ... Kreisdiagramm ... Speichere Abbildung in Datei イロン 人通 と 人 回 と 人 回 と

To install Rcmdr go to Packages \rightarrow Install package(s) (or simply type install.packages("Rcmdr")), then choose a CRAN mirror close to you, than OK. A window with a list of packages will pop-up, on this list choose Rcmdr and OK. A bundle of packages will be automatically installed.

To run the "R Commander" GUI type at the prompt line:

> library(Rcmdr)

This will start a GUI similar to other statistical software. Therefore, any typical process, like read data, produce plots, make statistical analyses, etc. will be made by clicking the appropriate menu.

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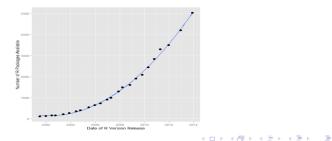
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Packages			

The capabilities of R are extended through user-created packages, which allow specialized statistical techniques, graphical devices, import/export capabilities, reporting tools, etc.

These packages are developed primarily in R, and sometimes in Java, C and Fortran. A core set of packages is included with the installation of R, with more than 6381 (as of Feb 2015) available at the Comprehensive R Archive Network (CRAN), 2095 (936 Software packages) on Bioconductor, and more on other repositories (e.g. R-Forge).



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R taskviews			

- of course: you can google your problem: but you should use http://www.rseek.org/ instead of www.google.com; rseek is a google custom search, can easily be added to the toolbar of popular browsers
- http://cran.r-project.org/web/views/
- before you install a new package: help.search() allows for searching the help system for documentation matching a given character string in the (file) name, alias, title, concept or keyword entries (or any combination thereof), using either fuzzy matching or regular expression matching.(installed help system)

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Packages			

- An R installation contains a library of packages. Some of these packages are part of the basic installation. These packages have the *recommended* status.
- Others (over 6000) can be downloaded from CRAN.
- A package is loaded into R using the library() or the require() command. For example to load the survival package you should enter
 - > library(survival)
- The loaded packages are not considered part of the workspace. You need to load a package when you start a new R session.

Getting Pack	arec		
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- you can download a package from CRAN and install by using the *package menu* (bottom right corner)
- another effective way to download and install a package is by command line. For example the following line install the R commander package with all its dependencies:
 - > install.packages("Rcmdr", dependencies=TRUE)

• install now the packages ggplot2 and faraway

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- start R (depending on you OS and UI) by double-clicking the R icon, typing R in a console or starting you favorite UI
- choose your working directory (via a menu or by typing setwd('/your/directory/')
- R works fundamentally by the question-and-answer model: you enter a line with a command and press Enter (↔). Then the program does something, prints the results, and asks for more input. When R is ready for input, it prints out its prompt, a ">". It is possible to use R as a text-only application, and also in batch mode.

One of the simplest possible tasks in R is to enter an arithmetic expression and receive a result.

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```
> 2 + 2
[1] 4
> exp(-2)
[1] 0.1353353
> round(exp(-2),3)
[1] 0.135
>
```

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• During a session you create a workspace. The workspace contains all variables created, for example typing

> y <- 2 * x + rnorm(100, mean=0, sd=0.5)

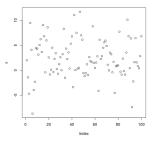
creates a vector variable with 100 random numbers from a normal distribution with mean 2 and standard deviation 4 and a second vector containing also 100 numbers dependend on \times

• to see the contents of this variables just type its names, e.g.

```
> x
[1] 2.663558 2.187709 -1.849147
5.566364 2.5016523.046095 ...
```

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- To plot these values type
 - > plot(x)



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Nothing is lost or	hidden		

- statistical packages provide *canned* procedures to address common statistical problems
- canned procedures are useful for routine analysis, but they are also limiting - you can only do what the programmer lets you do
- in R, the result of statistical calculation are always accessible, so
 - you can use them for further calculations
 - you can always see how calculations were done
- what you see in the first place is most of the time only a small part of the result

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• for example building a linear model gives you the following result

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```
> lm(y~x)
```

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Call: lm(formula = y ~ x) Coefficients: (Intercept) x 0.07101 1.98083

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No	othing is lost or	hidden		
		el in an object mm ary() or plot()	so you can run functio	ns on

- > mm <- lm(y~x)
- > summary(mm)

```
Call:
lm(formula = y ~ x)
```

Residuals:

```
Min 1Q Median 3Q Max
-1.67162 -0.36329 0.02206 0.29193 1.36333
```

```
Coefficients:
```

Estimate Std. Error t value Pr(>|t|) (Intercept) 0.07101 0.06353 1.118 0.266 x 1.98083 0.01404 141.043 <2e-16 *** ---Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1

Residual standard error: 0.5385 on 98 degrees of freedom Multiple R-squared: 0.9951, Adjusted R-squared: 0.995 F-statistic: 1.989e+04 on 1 and 98 DF, p-value: < 2.2e-16

Nothing is lost o	r hidden		
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• to explore the object you can use the object browser or str()

```
> str(mm)
List of 12
 $ coefficients : Named num [1:2] 0.071 1.981
  ..- attr(*, "names")= chr [1:2] "(Intercept)" "x"
 $ residuals : Named num [1:100] 0.6828 -0.1417 0.0992 0.1057 ...
  ..- attr(*, "names")= chr [1:100] "1" "2" "3" "4" ...
 $ effects : Named num [1:100] -48.23926 -75.95182 0.04414 ...
  ..- attr(*, "names")= chr [1:100] "(Intercept)" "x" "" "...
 $ rank : int 2
 $ fitted.values: Named num [1:100] 9.18 13.41 8.12 15.11 5.96 ...
  ..- attr(*, "names")= chr [1:100] "1" "2" "3" "4" ...
 $ assign : int [1:2] 0 1
 $ gr :List of 5
  ..$ qr : num [1:100, 1:2] -10 0.1 0.1 0.1 0.1 0.1 0.1 ...
  ....- attr(*, "dimnames")=List of 2
  ....$ : chr [1:100] "1" "2" "3" "4" ...
  .....$ : chr [1:2] "(Intercept)" "x"
  ....- attr(*, "assign")= int [1:2] 0 1
. . .
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• Entering the command

> help.start()

at the command line, will launch an extensive online help that can be read using a Web browser such as Firefox or Internet Explorer. Another way to access to these "help" pages is the help tab in the bottom right corner. Notice that the HTML version of the help system has a very useful "Search Engine and Keywords".

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	Statistica	l Data Analysis 📿			
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First Session			

• All variables, functions and diverse objects can be seen by the ls() and the newer version of it objects() function. Thus in our example we will have

> ls() [1] "x"

• quitting R is done with the q() function

> q()

at the command prompt. You will be asked to save your "workspace image". Give a name for your workspace for example "project1", if you want to save your workspace. Or use save() to save only the objects you want to keep.

• you can load this workspace in a new R session. On windows just click directly on the workspace file and R will be opened and automatically load the workspace

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Citation			
Input			

```
> citation()
```

To cite R in publications use:

```
R Development Core Team (2012). R: A language and environment for
statistical computing. R Foundation for Statistical Computing,
Vienna, Austria. ISBN 3-900051-07-0, URL http://www.R-project.org/.
```

```
A BibTeX entry for LaTeX users is
```

```
@Manual,
title = R: A Language and Environment for Statistical Computing,
author = R Development Core Team,
organization = R Foundation for Statistical Computing,
address = Vienna, Austria,
year = 2012,
note = ISBN 3-900051-07-0,
url = http://www.R-project.org/,
```

We have invested a lot of time and effort in creating R, please cite it when using it for data analysis. See also citation("pkgname") for citing R packages.

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Licence

Licence

R is mainly distributed under the terms of the GNU General Public License, either Version 2, June 1991 or Version 3, June 2007. Core Bioconductor packages are typically licensed under Artistic-2.0. You get detailed information with: license(), RShowDoc("COPYING"), packageDescription("packagename")\$License

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Exercises I			

start R in your favourite UI!

- Itry to load the MASS package (libary())! The MASS package belongs to the recommended packages and should be included in every standard installation of R. If contrary to expactations the MASS package is not installed, install it using the install.packages() command or the appropriate menue of your ui.
- after loading the package try to load some example data: the data() command loads data contained in packages. Here we want to load and inspect the Pima.tr data set. Type the following lines. What are they for?

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Exercises II			

- > library(MASS)
- > ## require(MASS)
- > data(Pima.te)
- > ?Pima.te
- > names(Pima.te)
- > head(Pima.te)
- > summary(Pima.te)

So Pima.te is here the name of the table. names() gives you the names of the columns. Remember: To access one particular column type

> table_name\$column_name

So to get the column skin out of the Pima.te data frame type

> Pima.te\$skin