



# An Introduction to Scientific Image Data Analysis using R

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- Introduction
- First steps with the R environment and Language
- Statistics and Plots in R
- Data Transfer and Analysis with ImageJ and R
- Useful Packages and Resources



- R is a language and environment for statistical computing and graphics
- OpenSource (GPL, GPL2)
- R is an implementation of the [S programming language](#) combined with [lexical scoping](#) semantics inspired by [Scheme](#)
- First binary 1993 (source avail. 1995) by Ross Ihaka and Robert Gentleman
- First production release 2000 (1.0)
- Current Release 2.15.2

# Why Use R?

- Open Source and available for all popular OS environments
- Huge community
- Well documented (Many specialized books)
- Statistical routines not yet available in other packages
- Can be extended with over 4000 Packages (at the moment)!
- Packages can be written in different languages (C, C++, Java, Fortran)
- Can be called from Java (libraries available)
- Many different free GUI-Applications available
- Many Plot types available

# Installation

**Windows:** “base” installation

**MacOSX:** pkg installation

**Linux:** sudo yum install R-core.i386 or R-core.x86\_64 (Fedora)

sudo apt-get install r-base (Ubuntu)

## Start Linux:

e.g. /usr/sbin/R (Shell)

or:

/usr/sbin/R -g TK & (with Window)

Packages Linux:

e.g.: /usr/lib/R/site-library

**Data Editor Matrices and Dataframes:**

edit(object)

# R Workspace

The so-called workspace is the current **R** working environment. It contains all defined objects.

This workspace can be saved or reloaded with the following commands:

## **Save Workspace:**

```
save.image(„yourFile.RData“)
```

## **Save objects to a file:**

```
save(object list,file="myfile.RData")
```

## **Load Workspace:**

```
load(„yourFile.RData“)
```

# Basic Data Structures

## Vectors

Values have the same data type (Character or numeric.. etc.)

## Factors

A Factor is an ordered collection of items (categorical values). The different values are called levels. They are stored internally as integers

## Matrices

A two-dimensional structure which can contain one datatype. Values are in rows and columns. Matrices are 2-d arrays. For more dimensions arrays are needed!

## Data frames

For tabular data. Contains vectors with same length. Each column can have a different datatype.

## Lists

A list is a collection of objects. A list can contain e.g. a list (2d) of arbitrary length. Different objects in one list are possible.

## Arrays

The **array** is similar to a matrix but can have more dimensions

## Conversion

- |         |           |           |
|---------|-----------|-----------|
| •matrix | as.matrix | is.matrix |
| •vector | as.vector | is.vector |
| •array  | as.array  | is.array  |

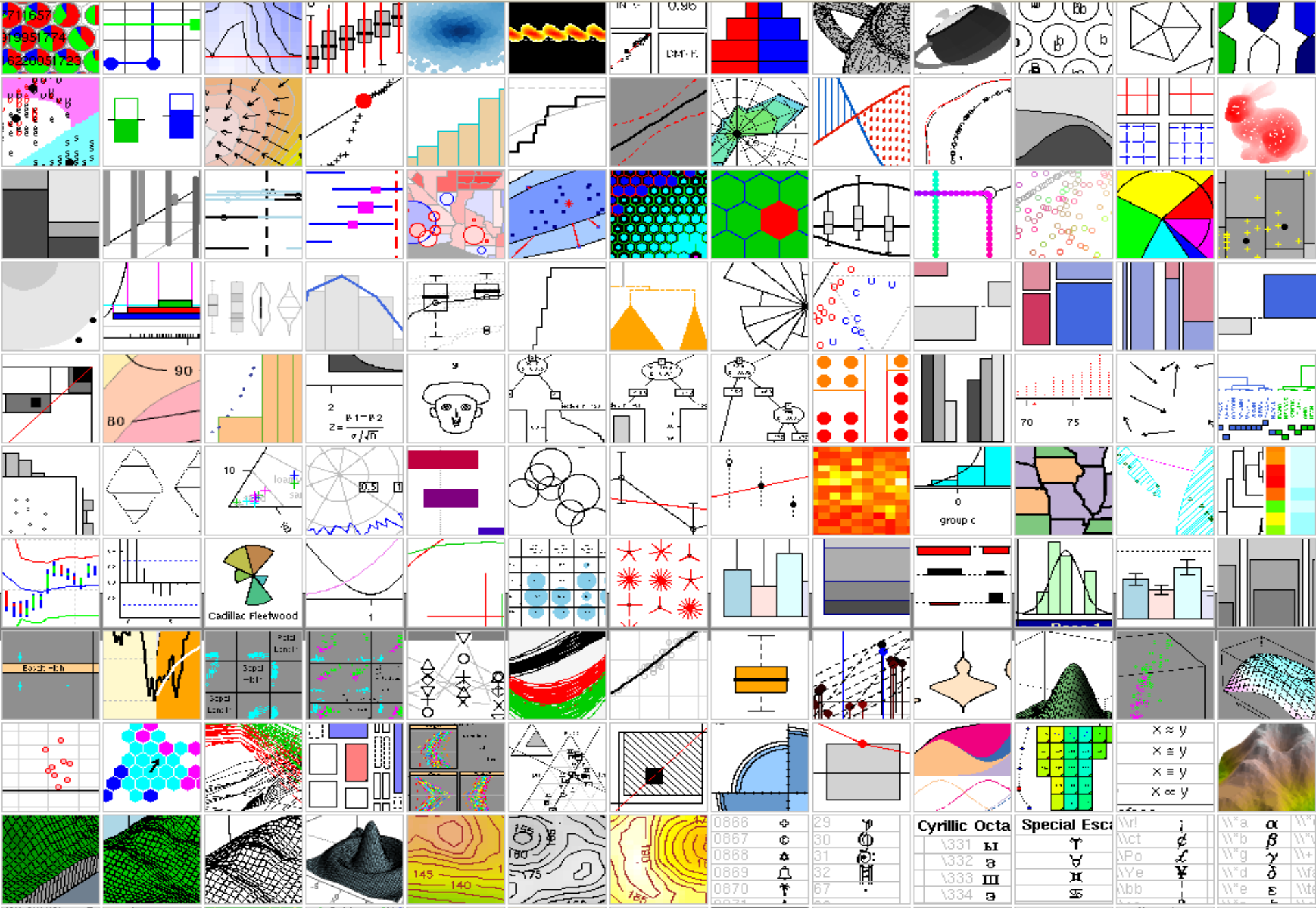
## Information

- class(x)
- dim(x)
- length(x)



# Basic Data Types

|            |              |            |
|------------|--------------|------------|
| •integer   | as.integer   | is.integer |
| •double    | as.double    | ...        |
| •numeric   | as.numeric   | is.numeric |
| •complex   | as.complex   | ...        |
| •character | as.character | ...        |
| •logical   | as.logical   | ...        |
| •raw       | as.raw       | ...        |



<http://gallery.r-enthusiasts.com/>

erri plain  
erif italic  
erif bold  
f bold italic  
rif cyrillic

160

0866  
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0868  
0869  
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Cyrillic Octa  
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Special Escapes  
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\e

# **First Steps with the R Environment and Language**

# Cran Task Views

|   |   |
|---|---|
| <a href="#">Bayesian</a>                  | Bayesian Inference  |
| <a href="#">ChemPhys</a>                  | Chemometrics and Computational Physics                                |
| <a href="#">ClinicalTrials</a>            | Clinical Trial Design, Monitoring, and Analysis                       |
| <a href="#">Cluster</a>                   | Cluster Analysis & Finite Mixture Models                              |
| <a href="#">DifferentialEquations</a>     | Differential Equations  |
| <a href="#">Distributions</a>             | Probability Distributions   |
| <a href="#">Econometrics</a>              | Computational Econometrics  |
| <a href="#">Environmetrics</a>            | Analysis of Ecological and Environmental Data                         |
| <a href="#">ExperimentalDesign</a>        | Design of Experiments (DoE) & Analysis of Experimental Data           |
| <a href="#">Finance</a>                   | Empirical Finance   |
| <a href="#">Genetics</a>                  | Statistical Genetics  |
| <a href="#">Graphics</a>                  | Graphic Displays & Dynamic Graphics & Graphic Devices & Visualization |
| <a href="#">HighPerformanceComputing</a>  | High-Performance and Parallel Computing with R                        |
| <a href="#">MachineLearning</a>           | Machine Learning & Statistical Learning                               |
| <a href="#">MedicalImaging</a>            | Medical Image Analysis  |
| <a href="#">Multivariate</a>              | Multivariate Statistics   |
| <a href="#">NaturalLanguageProcessing</a> | Natural Language Processing   |
| <a href="#">OfficialStatistics</a>        | Official Statistics & Survey Methodology                              |
| <a href="#">Optimization</a>              | Optimization and Mathematical Programming                             |
| <a href="#">Pharmacokinetics</a>          | Analysis of Pharmacokinetic Data                                      |
| <a href="#">Phylogenetics</a>             | Phylogenetics, Especially Comparative Methods                         |
| <a href="#">Psychometrics</a>             | Psychometric Models and Methods                                       |
| <a href="#">ReproducibleResearch</a>      | Reproducible Research   |
| <a href="#">Robust</a>                    | Robust Statistical Methods  |
| <a href="#">SocialSciences</a>            | Statistics for the Social Sciences                                    |
| <a href="#">Spatial</a>                   | Analysis of Spatial Data  |
| <a href="#">Survival</a>                  | Survival Analysis   |
| <a href="#">TimeSeries</a>                | Time Series Analysis  |
| <a href="#">gR</a>                        | gRaphical Models in R   |

# Image Packages

## CRAN packages Medical Images:

[adimpro](#) (core)

[AnalyzeFMRI](#) (core)

[arf3DS4](#) (core)

[cudaBayesreg](#)

[DATforDCEMRI](#) (core)

[dcmriS4](#) (core)

[dpmixsim](#) (core)

[dti](#) (core)

[fmri](#) (core)

[mmand](#) (core)

[mritc](#) (core)

[neuRosim](#) (core)

[occ](#) (core)

[oro.dicom](#) (core)

[oro.nifti](#) (core)

[PET](#) (core)

[PTAk](#)

[Rniftilib](#) (core)

[RNiftyReg](#) (core)

[tractor.base](#) (core)

## General Image Packages

### EBImage

EBImage is an R package which provides general purpose functionality for the reading, writing, processing and analysis of images.

### biOps

Image processing and analysis

### rimage?

Image Processing Module for R

### RImageJ?

R bindings for ImageJ

## Spatial Packages (Raster data)

### rgdal

Bindings to Frank Warmerdam's Geospatial Data Abstraction Library (GDAL)

### raster

Reading, writing, manipulating, analyzing and modeling of gridded spatial data

### spatstat

A package for analysing spatial data, mainly Spatial Point Patterns

# Spatstat

**im** create a pixel image  
**as.im** convert other data to a pixel image  
**pixellate** convert other data to a pixel image  
**as.matrix.im** convert pixel image to matrix  
**as.data.frame.im** convert pixel image to data frame  
**plot.im** plot a pixel image on screen as a digital image  
**contour.im** draw contours of a pixel image  
**persp.im** draw perspective plot of a pixel image  
**rgbim** create colour-valued pixel image  
**hsvim** create colour-valued pixel image  
[.im extract a subset of a pixel image  
[<-.im replace a subset of a pixel image  
**shift.im** apply vector shift to pixel image  
**X** print very basic information about image X  
**summary(X)** summary of image X  
**hist.im** histogram of image  
**mean.im** mean pixel value of image  
**integral.im** integral of pixel values  
**quantile.im** quantiles of image  
**cut.im** convert numeric image to factor image  
**is.im** test whether an object is a pixel image  
**interp.im** interpolate a pixel image  
**blur** apply Gaussian blur to image  
**connected** find connected components  
**compatible.im** test whether two images have comp. dim.  
**harmonise.im** make images compatible  
**commonGrid** find a common pixel grid for images  
**eval.im** evaluate any expression involving images  
**scaletointerval** rescale pixel values  
**zapsmall.im** set very small pixel values to zero  
**levelset** level set of an image  
solutionset region where an expression is true  
**imcov** spatial covariance function of image  
**convolve.im** spatial convolution of images  
**transect.im** line transect of image

The package supports

- creation, manipulation and plotting of point patterns
- exploratory data analysis
- simulation of point process models
- parametric model-fitting
- hypothesis tests and model diagnostics

e.g.:

Summary statistics for a point pattern

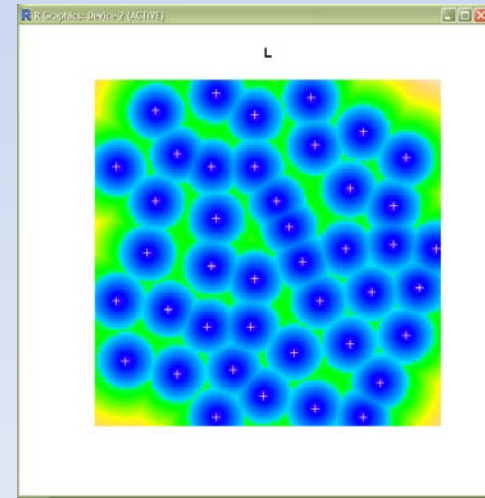
Summary statistics for a multitype point pattern

Line segment patterns

Tessellations

Three-dimensional point patterns

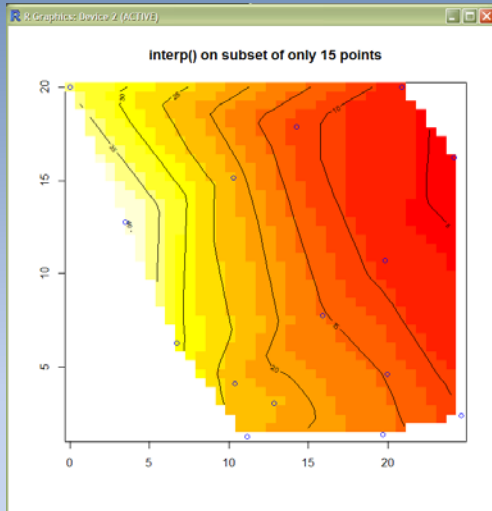
.....



# Interpolation, Spatial Modelling

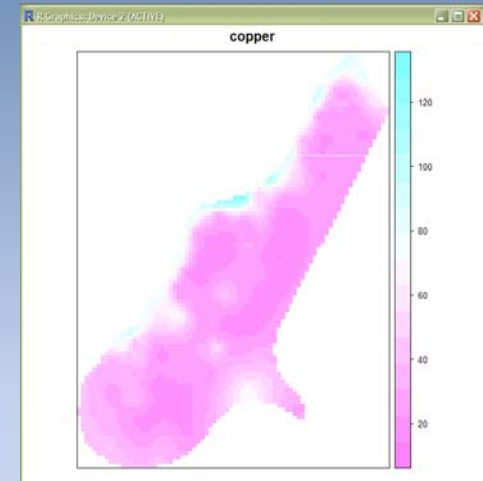
## akima

Linear or cubic spline interpolation for irregular gridded data



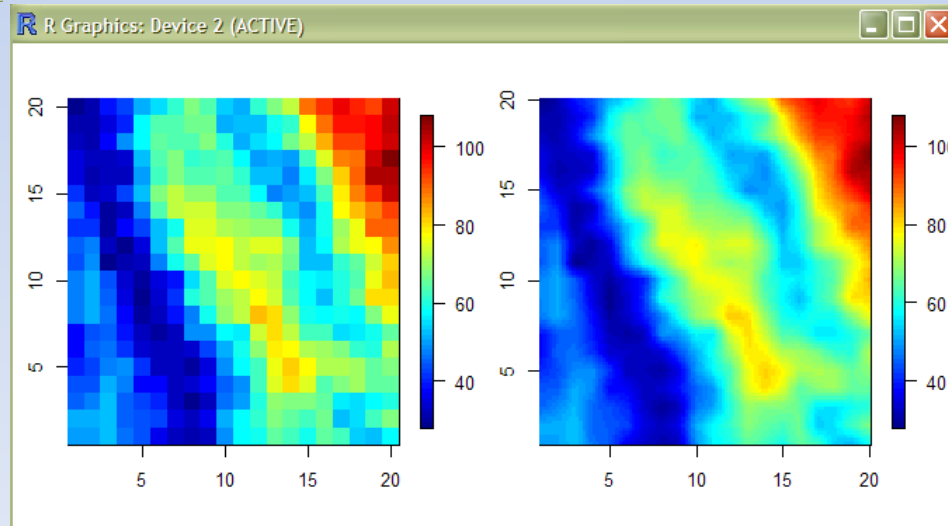
## gstat

spatial and spatio-temporal geostatistical modelling, prediction and simulation



## fields

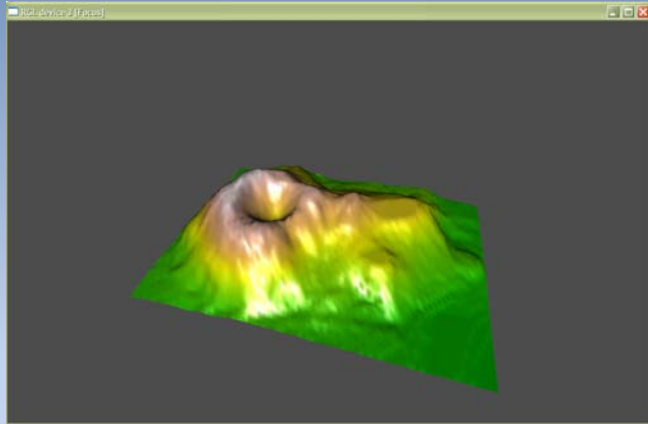
Fields is for curve, surface and function fitting with an emphasis on splines, spatial data and spatial statistics



# Plot Packages

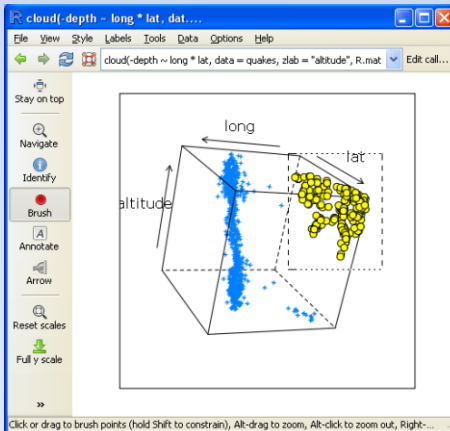
**rgl**

<http://rgl.neoscientists.org/about.shtml>



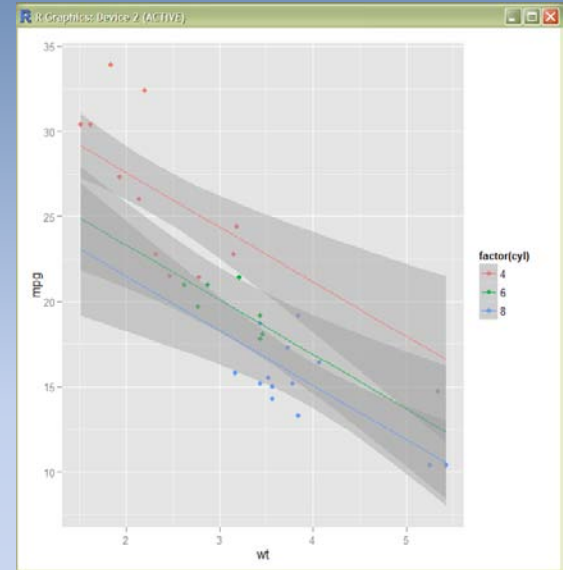
**playwith**

<http://code.google.com/p/playwith/>



**ggplot2**

<http://ggplot2.org/>



And:

**animation**  
**scatterplot3d**  
**plotrix**

.....



# Reporting in R

## Sweave

“Sweave is a tool that allows to embed the R code for complete data analyses in latex documents. The purpose is to create dynamic reports, which can be updated automatically if data or analysis change.”

Source: <http://www.statistik.lmu.de/~leisch/Sweave/>

## Knitr

“The **knitr** package was designed to be a transparent engine for dynamic report generation with R, solve some long-standing problems in Sweave, and combine features in other add-on packages into one package.”

Source: <http://yihui.name/knitr/>

# Knitr example

```
library(ggplot2)
pie <- ggplot(diamonds, aes(x = factor(1), fill = cut)) +
  xlab("cut") + geom_bar(width = 1)
pie + coord_polar(theta = "y") # a pie chart
pie + coord_polar() # the bullseye chart
```

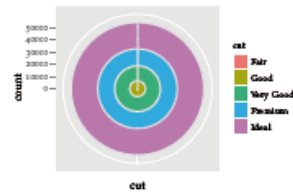


Figure 6: Two plots were produced in this chunk, but only the last one is kept. This can be useful when we experiment with many plots, but only want the last result (Adapted from the ggplot2 website)

When multiple plots are produced by a code chunk, we may want to show them as an animation with the option `fig.show='animate'`. Figure 7 shows a simple clock animation; you may compare the code to Figure 5 to understand that high-level plots are always recorded, regardless of where they appeared.

```
par(mar = rep(3, 4))
for (i in seq(pi/2, -4/3 * pi, length = 12)) {
  plot(0, 0, pch = 20, ann = FALSE, axes = FALSE)
  arrows(0, 0, cos(i), sin(i))
  axis(1, 0, "VI"); axis(2, 0, "IX")
  axis(3, 0, "XII"); axis(4, 0, "III"); box()
}
```

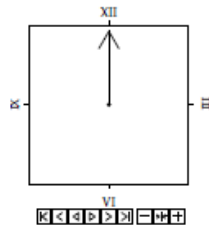
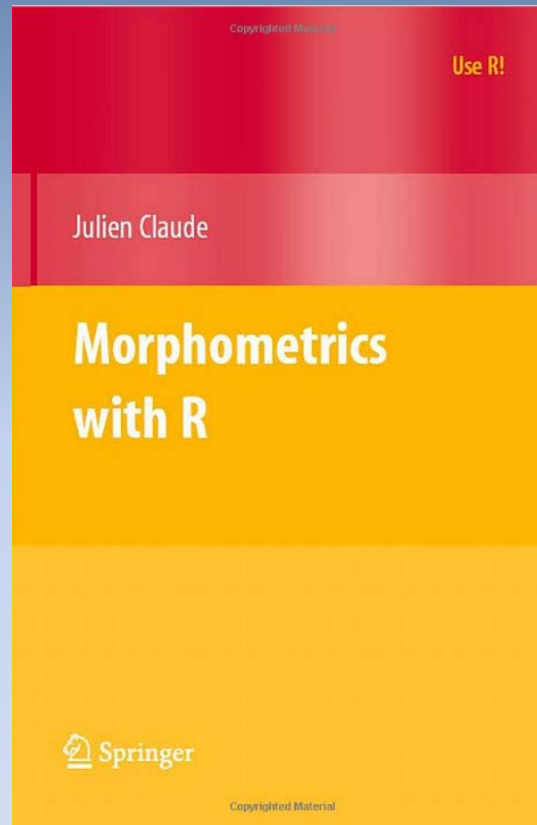


Figure 7: A clock animation. You have to view it in Adobe Reader: click to play/pause; there are also buttons to speed up or slow down the animation.

We can also set the alignment of plots easily with the `fig.align` option; this document uses `fig.align='center'` as a global option, and we can also set plots to be left/right-aligned. Figure 8 is an example

# Morphometrics



Packages:

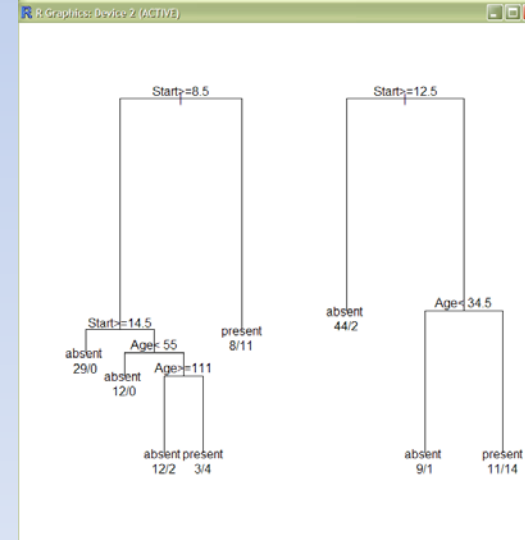
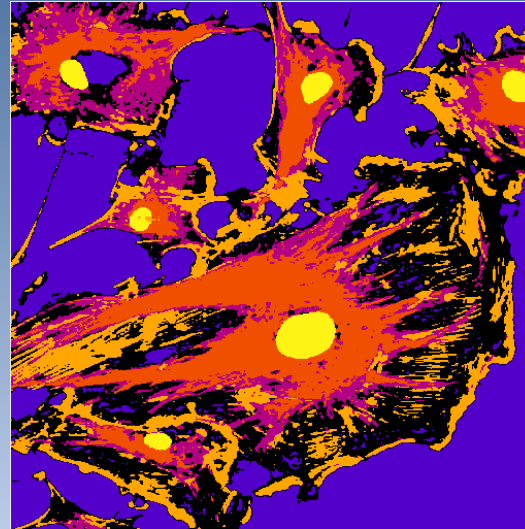
Momocs  
shapes

Source: <http://www.isem.cnrs.fr/spip.php?article835>

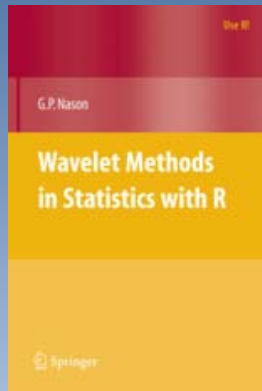
# Maschine Learning / Supervised or Unsupervised Learning

## CRAN packages :

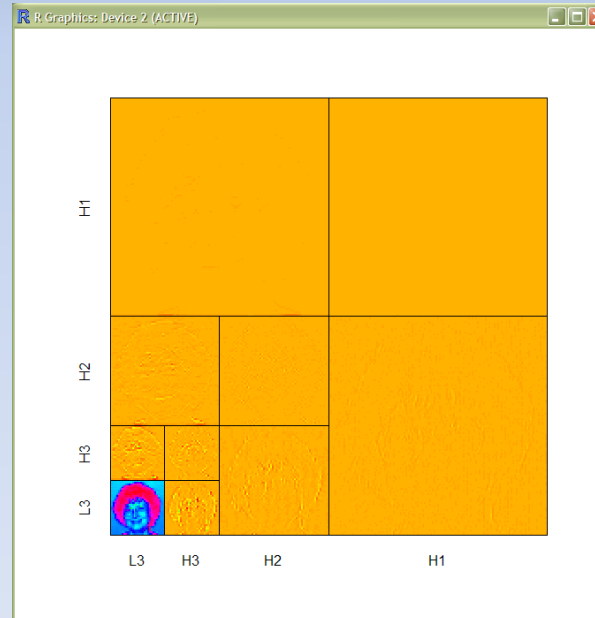
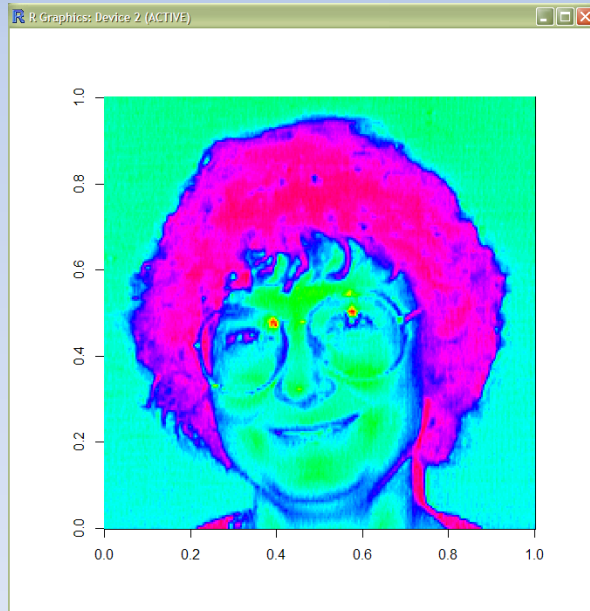
- [ahaz](#)
- [arules](#)
- [BayesTree](#)
- [Boruta](#)
- [BPHO](#)
- [bst](#)
- [C50](#)
- [caret](#)
- [CORElearn](#)
- [CoxBoost](#)
- [Cubist](#)
- [e1071](#) (core)
- [earth](#)
- [elasticnet](#)
- [ElemStatLearn](#)
- [evtree](#)
- [gafit](#)
- [GAMBoost](#)
- [gamboostLSS](#)
- [gbm](#) (core)
- [glmnet](#)
- [glmpath](#)
- [GMMBoost](#)
- [grplasso](#)
- [grpreg](#)
- [hda](#)
- [ipred](#)
- [kernlab](#) (core)
- [klaR](#)
- [lars](#)
- [lasso2](#)
- [Liblinear](#)
- [LogicForest](#)
- [LogicReg](#)
- [longRPart](#)
- [mboost](#) (core)
- [mvpart](#)
- [ncvreg](#)
- [nnet](#) (core)
- [oblique.tree](#)
- [pamr](#)
- [party](#)
- [partykit](#)
- [penalized](#)
- [penalizedSVM](#)
- [predbayescor](#)
- [quantregForest](#)
- [randomForest](#) (core)
- [randomSurvivalForest](#)
- [rattle](#)
- [rda](#)
- [rdetools](#)
- [REEMtree](#)
- [relaxo](#)
- [rgenoud](#)
- [rgp](#)
- [rminer](#)
- [ROCR](#)
- [rpart](#) (core)
- [rpartOrdinal](#)
- [RPMM](#)
- [RSNNS](#)
- [RWeka](#)
- [sda](#)
- [SDDA](#)
- [svmpath](#)
- [tgp](#)
- [tree](#)
- [TWIX](#)
- [varSelRF](#)



# Wavelett Packages



- wavethresh
- wavelets
- waveslim



Source: Package "waveslim", Author: Brandon Whitcher

# Using R in Java

## Libraries and Projects

### **rJava**

<http://www.rforge.net/rJava/>

### **Rserve**

<http://rforge.net/Rserve/>

### **RCaller**

<http://www.mhsatman.com/rcaller.php>

### **JVM-based Interpreter for the R Language**

renjin

<http://code.google.com/p/renjin/>

# R GUI's

## **R Commander**

<http://socserv.mcmaster.ca/jfox/Misc/Rcmdr/>

## **JGR**

<http://rforge.net/JGR/>

## **Deducer**

<http://www.deducer.org/>

## **RStudio**

<http://rstudio.org/>

## **RKward**

<http://rkward.sourceforge.net/>

## **Tinn-R**

<http://www.sciviews.org/Tinn-R/>

## **StatET**

<http://www.walware.de/goto/statet>

## **Elastic-R**

<http://www.elastic-r.net/>

## **Bio7**

<http://bio7.org>

**Thank You!**



# Numeric Precision

[http://cran.r-project.org/doc/FAQ/R-FAQ.html#Why-doesn 0027t-R-think-these-numbers-are-equal 003f](http://cran.r-project.org/doc/FAQ/R-FAQ.html#Why-doesn%20t-R-think-these-numbers-are-equal)

[http://rwiki.sciviews.org/doku.php?id=misc:r\\_accuracy:high\\_precision\\_arithmetic](http://rwiki.sciviews.org/doku.php?id=misc:r_accuracy:high_precision_arithmetic)

# Batch Execution of R

R CMD BATCH [options] infile [outfile]

Run **R** non-interactively with input from infile and send output (stdout/stderr) to another file.