

Statistical Analysis of Optimization Algorithms with R

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July 2012

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GECCO'12 Companion, July 7-11, 2012, Philadelphia, PA, USA.
ACM 978-1-4503-1178-6/12/07.

Your Instructors Today

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- ▶ Mike Preuss is research associate at the Computer Science Department, TU Dortmund. His main fields of activity are EAs for real-valued problems and their application in numerous engineering domains
- ▶ Martin Zaefferer is a research assistant at Cologne University of Applied Sciences. His research interests include computational intelligence, applications of knowledge discovery and sequential parameter optimization.

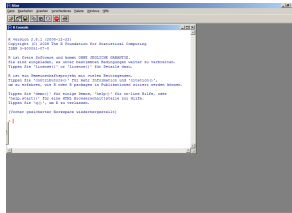
Agenda

- Introduction
- R Basics and Technical Details
- Exploratory Data Analysis
- Distributions and Random Number Generation
- Design of Experiments (DoE)
- R-based automated analysis and tuning, e.g., sequential parameter optimization
- Reporting results. Automated report generation using Sweave
- R-based optimization and benchmarking resources

Goals

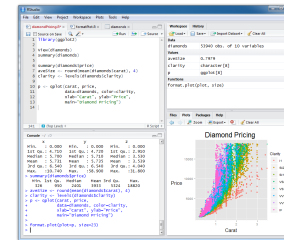
- ▶ Most effective approach for learning how to design, conduct, and analyze experiments that optimize performance in algorithms
- ▶ Show how to use statistically designed experiments to
 - ▶ Obtain information for characterization and optimization of algorithms
 - ▶ Improve their performance
 - ▶ Design and develop new operators and algorithms
- ▶ Learn how to evaluate algorithm alternatives, improve their field performance and reliability
- ▶ Conduct experiments effectively and efficiently
- ▶ Hands-on tutorial which
 - ▶ demonstrates how to analyze results from real experimental studies, e.g., experimental studies in EC
 - ▶ gives a comprehensive introduction in the R language
 - ▶ introduces the powerful GUI "rstudio" (<http://rstudio.org>)
 - ▶ will be held in an interactive manner, i.e., the analyses will be performed in real time.

Pure R



- Windows version comes with a simple build-in GUI

RStudio



- Powerful productivity tools
 - Syntax highlighting, code completion, and smart indentation
 - Execute R code directly from the source editor
 - Easily manage multiple working directories using projects
 - Quickly navigate code using typeahead search and go to definition

The Famous Iris Data Set

- Four features were measured from each sample
- Length and width of sepal and petal, respectively



Photo by[7]

The Famous Iris Data Set: Iris Setosa, Virginica, and Versicolor

- Based on the combination of the four features, Fisher [5] developed a linear discriminant model to determine which species from these four measurements
- Used as a typical test for many other classification techniques



- Iris setosa [3], iris virginica [10], and iris versicolor [8]

The Famous Iris Data Set: Hands-on Exercises

- How to generate a scatter plot of Fisher's Iris data with pure R code
- First, we load the data frame:

```
> data(iris)
```

- Next, we have a quick look at the data (here, only the first three rows are shown)

```
> 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
```

The Famous Iris Data Set: Hands-on Exercises

- The `summary()` command gives a quick overview

```
> options(width=70)
> summary(iris)

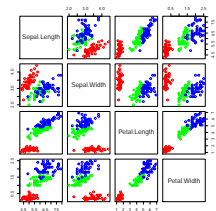
 Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
Min.      :4.300   Min.      :2.000   Min.      :1.000   Min.      :0.100
1st Qu.:5.100   1st Qu.:2.800   1st Qu.:1.600   1st Qu.:0.300
Median :5.800   Median :3.000   Median :4.350   Median :1.300
Mean    :5.843   Mean    :3.057   Mean    :3.758   Mean    :1.199
3rd Qu.:6.400   3rd Qu.:3.300   3rd Qu.:5.100   3rd Qu.:1.800
Max.    :7.900   Max.    :4.400   Max.    :6.900   Max.    :2.500

Species
setosa      :50
versicolor:50
virginica   :50
```

The Famous Iris Data Set: Hands-on Exercises

- Finally, a scatter plot is generated with the `pairs()` function.

```
> pairs(iris[,1:4], col=c("red", "green", "blue")[as.numeric(iris$Species)])
```



A Gentle Introduction to R

- Some of the following examples are based on [9]

- R can be used as a calculator

```
> 2+2
```

```
[1] 4
```

```
> 5+3*4
```

```
[1] 60
```

- Data entry

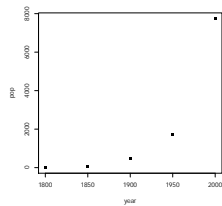
```
> year <- c(1800,1850,1900,1950,2000)
```

```
> pop <- c(18,54,500,1701,7731)
```

A Simple plot

Data entry:

```
> print( c(year,pop))  
[1] 1800 1850 1900 1950 2000 18 54 500 1701 7731  
> plot(pop~year, pch=15)
```



R sessions

The working directory.

```
> getwd()
```

```
[1] "C:/Users/bartz/Documents/workspace/SvnSpot.d/trunk/publications/Gecco2012Tutorial.d/Slides2012.d"
```

Use `ls` to list contents of R's workspace:

```
> ls()
```

```
[1] "iris" "mygd" "pop" "year"
```

Quitting: Note, `q()` is a function and can be used if R should be quit.

Expressions, Objects, and Methods

- ▶ Standard interaction mode in R is as follows:
 - ▶ Users enter an expression, which is evaluated by the R system. Result is printed on the screen
- ▶ Expressions work on *objects*
- ▶ Each object has a *class* attribute, which is a character vector

```
> x <- 10  
> class(x)  
[1] "numeric"
```

Logical Operators and Vectors in R

- ▶ R implements the following logical operators
 - ▶ `&`, the logical "and",
 - ▶ `|`, the logical "or", and
 - ▶ `!`, the logical "not" operator
- ▶ R commands to generate vectors:
 - ▶ `c()` ("concatenate"),
 - ▶ `seq()` ("sequence"), and
 - ▶ `rep()` ("replicate")
- ▶ Modes: logical, numeric, character, or list

```
> x <- c(1,2)  
> y <- c(3,4)  
> z <- c(x,y)  
> x ==y  
[1] FALSE FALSE
```

Vectors in R

- ▶ R's repeat command `rep()` can be used in two variants
 - ▶ To repeat the numerical value one ten times, we use


```
> rep(1,10)
[1] 1 1 1 1 1 1 1 1 1 1
```
 - ▶ The second argument to the `rep()` command can be a vector.


```
> v <- c(1,2,4)
> letters <- c("a","b","c")
> rep(letters, v)
[1] "a" "b" "b" "c" "c" "c" "c" "c"
```
- ▶ Here, the first element "a" is repeated once, the second element "b" twice, and the third element "c" four times

Vectors in R: Sequences

```
> seq(from=5, to=22, by=3)
[1] 5 8 11 14 17 20
```

Short form

```
> seq(5,22,3)
[1] 5 8 11 14 17 20
```

Default step size is one. Short form with colon

```
> seq(0,10) == 0:10
[1] TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE
```

Character vectors: vectors of text strings, entries are specified in quotes

```
> c("one","two","three")
[1] "one" "two" "three"
```

Calculations with Vectors in R

- ▶ Calculations with vectors of the same length (like ordinary numbers)


```
> x <- c(1,2,3)
> y <- c(1,2,4)
> x+y
[1] 2 4 7
```
- ▶ Relational expressions can be evaluated as follows


```
> x < y
[1] FALSE FALSE TRUE
```
- ▶ If vectors do not have the same length, the shorter vector is recycled


```
> y <- c(5,6)
> x+y
[1] 6 8 8
```
- ▶ Vector `v` modified to `(5,6,5)`, i.e., the first element is added at the end. Both vectors have the same size and can be added

Calculating the mean of vectors in R

Consider the vector

```
> x <- c(2,3,5,7)
```

To calculate its mean,

$$\bar{x} = \frac{1}{n} \sum_{i=1}^n x_i,$$

we can proceed as follows:

```
> sum(x)/length(x)
```

```
[1] 4.25
```

Calculating the standard deviation of vectors in R

To calculate its standard deviation,

$$sd(x) = \sqrt{\sum_{i=1}^n \frac{(x_i - \bar{x})^2}{n-1}},$$

we can proceed as follows:

```
> xbar <- sum(x)/length(x)
> sqrt( sum( (x - xbar)^2 / (length(x)-1) ) )
[1] 2.217356
```

Alternatively, we can use the build-in commands

```
> mean(x)
[1] 4.25
> sd(x)
[1] 2.217356
```

Subsets, and Indexing

Brackets are used to access certain elements of a vector. To select the i -th entry of the vector v , e.g., the third entry v_3 , we can use the command

```
> v <- c(10,20,30,40,50,60,70,80,90,100)
> v[3]
[1] 30
```

This procedure is referred to as *indexing* in the following. To select a subset, we can index with a vector.

```
> v[ c(3,4,5) ]
[1] 30 40 50
```

Use negative subscripts to omit elements in nominated subscript positions

```
> v[-c(2,3)]
[1] 10 40 50 60 70 80 90 100
```

Conditional Selection

► To modify elements of an vector, we can use the assignment operator

```
> v[1] <- -10
> v
[1] -10 20 30 40 50 60 70 80 90 100
```

► Conditional selection can be performed as follows

```
> v>50
[1] FALSE FALSE FALSE FALSE FALSE TRUE TRUE TRUE TRUE TRUE
> v[v>50]
[1] 60 70 80 90 100
```

► Logical operators can be used to combine several subset selection conditions.

► For example, to select entries, which are larger than 55 and smaller than 79, we can use the following command

```
> v[ v > 55 & v < 79 ]
[1] 60 70
```

Missing Values. The Symbol NA

```
> y <- c(1, NA, 3, 0, NA)
> y
[1] 1 NA 3 0 NA
```

Any operation that involves NA generates NA. The following does not work as expected, all values remain unchanged:

```
> y[y==NA] <- 0
> print(y)
[1] 1 NA 3 0 NA
```

To replace NA by 0, use `is.na()`:

```
> y[is.na(y)] <- 0
> print(y)
[1] 1 0 3 0 0
```

Some functions, e.g., `mean()` take the argument `"na.rm=T"`.

Factors

Categorical data such as gender can be "female" or "male", respectively. Categorical data should be specified as factors.

```
> gender <- c(rep("female", 3), rep("male", 5))
```

To generate a factor, use R's `factor()` command

```
> gender <- factor(gender)
```

Now: internally 3 1s are followed by 5 2s. We can use the function `as.numeric()` to extract the numerical coding as numbers "1" and "2".

```
> gender
```

```
[1] female female female male   male   male   male   male  
Levels: female male
```

```
> as.numeric(gender)
```

```
[1] 1 1 1 2 2 2 2 2
```

A factor has set of levels. "female" and "male" are the levels of the factor `gender`:

```
> levels(gender)
```

```
[1] "female" "male"
```

Matrices and Arrays

Matrices and arrays are vectors with dimensions.

```
> x <- 1:20  
> dim(x) <- c(5,4)  
> x
```

```
      [,1] [,2] [,3] [,4]  
[1,]    1    6   11   16  
[2,]    2    7   12   17  
[3,]    3    8   13   18  
[4,]    4    9   14   19  
[5,]    5   10   15   20
```

We can also use the `matrix()` command.

```
> matrix(1:20, nrow= 5)
```

```
      [,1] [,2] [,3] [,4]  
[1,]    1    6   11   16  
[2,]    2    7   12   17  
[3,]    3    8   13   18  
[4,]    4    9   14   19  
[5,]    5   10   15   20
```

Matrices and Arrays

To fill the matrix rowwise, we can use the argument `byrow=T`. To label the rows of a matrix, we can use the command `rownames()`.

```
> A <- matrix(1:20, nrow= 5, byrow=T)  
> rownames(A) <- LETTERS[1:nrow(A)]  
> colnames(A) <- 1:ncol(A)  
> A
```

```
      1  2  3  4  
A    1  2  3  4  
B    5  6  7  8  
C    9 10 11 12  
D   13 14 15 16  
E   17 18 19 20
```

Combining Matrices and Vectors

`cbind()` and `rbind()` combine objects such as matrices or vectors columnwise or rowwise, respectively

```
> A <- matrix(1:4, nrow= 2, byrow=T)  
> B <- matrix(10*(1:4), nrow= 2, byrow=T)  
> cbind(A,B)
```

```
      [,1] [,2] [,3] [,4]  
[1,]    1    2   10  20  
[2,]    3    4   30  40
```

```
> rbind(A,B)
```

```
      [,1] [,2]  
[1,]    1    2  
[2,]    3    4  
[3,]   10   20  
[4,]   30   40
```

Lists

Many R functions return results as a list. Flexible structures to store heterogeneous data, e.g., numerical or boolean values.

```
> l <- list( c("a", "b", "c"), 1:4, c(TRUE, TRUE, FALSE))  
> l
```

```
[[1]]  
[1] "a" "b" "c"
```

```
[[2]]  
[1] 1 2 3 4
```

```
[[3]]  
[1] TRUE TRUE FALSE
```

List l has three elements: 1) three strings, 2) numbers from one to four, and 3) three boolean values.

Addressing Lists

Single square brackets return a list.

```
> l[1]  
[[1]]  
[1] "a" "b" "c"
```

Here, l[1] returns a list of length one, whereas l[2:3] returns a list of length two.

```
> l[2:3]  
[[1]]  
[1] 1 2 3 4
```

```
[[2]]  
[1] TRUE TRUE FALSE
```

List elements can be addressed by double square brackets

```
> l[[1]]  
[1] "a" "b" "c"  
> l[[1]][2]  
[1] "b"
```

Adding and Deleting List Elements

List elements can be deleted by setting their values to NULL

```
> l  
[[1]]  
[1] "a" "b" "c"
```

```
[[2]]  
[1] 1 2 3 4
```

```
[[3]]  
[1] TRUE TRUE FALSE
```

Delete the first element

```
> l[[1]] <- NULL  
> l
```

```
[[1]]  
[1] 1 2 3 4
```

```
[[2]]  
[1] TRUE TRUE FALSE
```

To delete multiple list elements, we can use the minus sign

```
> l <- c(l,1)  
> l <- l[-c(3,4)]  
> l
```

```
[[1]]  
[1] 1 2 3 4
```

```
[[2]]  
[1] TRUE TRUE FALSE
```

Length of Lists

length() to determine the length of a list

```
> length(l)  
[1] 2
```

Adding a new element at the end using length()

```
> l[length(l)+1] <- c("x","y")  
> l
```

```
[[1]]  
[1] 1 2 3 4
```

```
[[2]]  
[1] TRUE TRUE FALSE
```

```
[[3]]  
[1] "x" "y"
```


Naming List Elements

We can use the `names()` function to add names to list elements.

```
> names(l) <- c("numbers", "booleans")
> l

$numbers
[1] 1 2 3 4

$booleans
[1] TRUE TRUE FALSE

$<NA>
[1] "x" "y"
```

Data Frames

Data frames can be used for grouping data. A data frame is a list of vectors of the same length.

```
> year <- c(1800, 1850, 1900, 1950, 2000)
> pop <- c(18, 54, 500, 1701, 7731)
> demography <- data.frame(y=year, p=pop)
> demography
```

	y	p
1	1800	18
2	1850	54
3	1900	500
4	1950	1701
5	2000	7731

Implicit Loops Using Apply

- ▶ `apply(x, margin, fun)` returns a vector or array or list of values obtained by applying a function to margins of an array or matrix
- ▶ Margin: vector giving the subscripts which the function will be applied over
- ▶ For example, for a matrix 1 indicates rows, 2 columns, `c(1, 2)` rows and columns, or in matrices named `dimnames`
- ▶ Implicit Loops Using `sapply()`, `lapply()`, and `tapply()`
 - ▶ `sapply()` returns a simplified result (vector or matrix),
 - ▶ `lapply()` returns a list, and
 - ▶ `tapply()` creates a table

Implicit Loops Using lapply()

List of car data:

```
> cars <- list(speed=c(180, 250, 300),
+ price = c(10.5, 55.6, 76.0),
+ consumption=c(5, 7.1, 12.5))
> cars

$speed
[1] 180 250 300

$price
[1] 10.5 55.6 76.0

$consumption
[1] 5.0 7.1 12.5
```

Consider the `sum()` function.

```
> lapply(cars, sum)

$speed
[1] 730

$price
[1] 142.1

$consumption
[1] 24.6
```

Implicit Loops Using Anonymous Functions

An anonymous function can be used as well

```
> lapply(cars, function(x) return(x[2]))
```

```
$speed  
[1] 250
```

```
$price  
[1] 55.6
```

```
$consumption  
[1] 7.1
```

Sorting

We can use the `sort()` function to sort a vector.

```
> sort(iris$Sepal.Length)[1:10]
```

```
[1] 4.3 4.4 4.4 4.4 4.5 4.6 4.6 4.6 4.6 4.7
```

The `order()` function generates a vector of the indices of the sorted values.

```
> a <- c(20,-1,4,3)  
> order(a)
```

```
[1] 2 4 3 1
```

Here, the smallest value "1" is at position 2, the next at position 4, whereas the largest value is at position 1.

sapply()

Given a list structure `x`, the function `unlist()` simplifies it to produce a vector. In order to obtain a vector of mean values instead of a list using the `lapply()` function, the following command can be used.

```
> unlist(lapply(cars, mean))
```

```
speed price consumption  
243.3333 47.36667 8.20000
```

Using `sapply()`, the same result can be obtained directly.

```
> sapply(cars, mean)
```

```
speed price consumption  
243.3333 47.36667 8.20000
```

For example, to apply `mean()` to each of the iris data set columns:

```
> data(iris)
```

```
> sapply(iris, mean)
```

```
Sepal.Length Sepal.Width Petal.Length Petal.Width Species  
5.843333 3.057333 3.758000 1.199333 NA
```

Sorting Using order()

```
> a
```

```
[1] 20 -1 4 3
```

```
> i <- order(a)
```

```
> i
```

```
[1] 2 4 3 1
```

```
> a[i]
```

```
[1] -1 3 4 20
```

Sorting a set of variables according to the values of some other variables:

```
> i <- order(iris$Sepal.Length)
```

```
> options(width=70)
```

```
> iris[i,][1:4,]
```

```
Sepal.Length Sepal.Width Petal.Length Petal.Width Species  
14 4.3 3.0 1.1 0.1 setosa  
9 4.4 2.9 1.4 0.2 setosa  
39 4.4 3.0 1.3 0.2 setosa  
43 4.4 3.2 1.3 0.2 setosa
```

Simple Formulas

```
> celsius<-(0:4)*10
> fahrenheit <- 9/5*celsius+32
> conversion <- data.frame(c=celsius, f=fahrenheit)
> print(conversion)
```

	c	f
1	0	32
2	10	50
3	20	68
4	30	86
5	40	104

Control Structures

Try using functions from the `apply(x)` family instead of loops.
`system.time()` returns CPU (and other) times used by process. First, an implementation without loops.

```
> require(stats)
> x <- 1:1000000
> system.time(y <- x^2)
```

user	system	elapsed
0.02	0.01	0.03

Next, the `for()` function to perform calculation with loops.

```
> x <- 1:1000000
> system.time(
+ for( i in 1:length(x)) y[i] <- x[i]^2
+ )
```

user	system	elapsed
2.49	0.02	2.53

Loops with `for()` and `while()`

Loops can be generated with the `for()` function as follows.

```
> for( i in 1:5) print(i)
```

```
[1] 1
[1] 2
[1] 3
[1] 4
[1] 5
```

We can also use the `while()` command.

```
> x<-1
> while(x <= 5){
+ print(x)
+ x <- x+1
+ }
```

```
[1] 1
[1] 2
[1] 3
[1] 4
[1] 5
```

Loops with `repeat()`

Alternatively, the `repeat()` command can be used in combination with the `break()` command.

```
> x <- 1
> repeat{
+ print(x)
+ x<-x+1
+ if (x > 5) break
+ }
```

```
[1] 1
[1] 2
[1] 3
[1] 4
[1] 5
```

R Functions and Arguments

R has many pre-defined functions, e.g., `mean`, `sum`, or `range`.

```
> year <- c(1800,1850,1900,1950,2000)
> pop <- c(18,54,500,1701,7731)
> range(year)
```

```
[1] 1800 2000
```

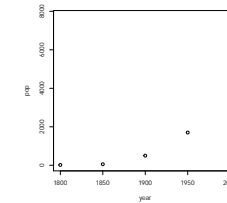
```
> summary(year)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1800	1850	1900	1900	1950	2000

Arguments and Positional Matching

R uses positional matching, i.e., the n -th argument corresponds to the n -th function variable. For example, `plot()` assumes: 1st argument (year) corresponds to x , whereas 2nd (population) corresponds to y

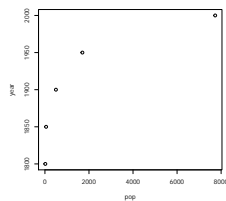
```
> plot(year, pop)
```



Named Actual Arguments

- Positional matching: very simple concept, becomes unhandily if many arguments occur
- R can handle *named actual arguments*, i.e., names are matched against their formal arguments
- Output from `plot()` with x and y values exchanged:

```
> plot(y=year, x=pop)
```



Writing R Functions

Using R's `function()` function, we can write our own functions. Note,

`%*%` denotes matrix multiplication

```
> norm <- function(x) sqrt(x%*%x)
> norm(1:4)
```

```
      [,1]
[1,] 5.477226
```

Curly braces can be used to define the body of the function.

```
> h <- function(x){
+   if (x<0) -1
+   else 1}
> h(1)
```

```
[1] 1
```

Note, we can use the command `ifelse()` as well.

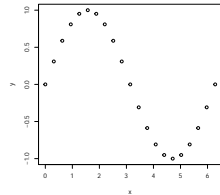
```
> heaviside <- function(x) ifelse(x<0,-1,1)
> heaviside(1)
```

```
[1] 1
```

Graphics: The Basic Plot Command `plot(x,y)`

- The basic plot command is `plot(x,y)`
- Alternatively, `plot(y ~ x)` can be used

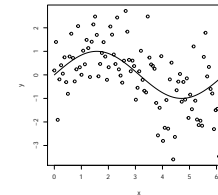
```
> x <- (0:20)*pi/10
> y <- sin(x)
> plot(y~x)
```



Combining Plots

- Add lines to this plot using the function `lines()`

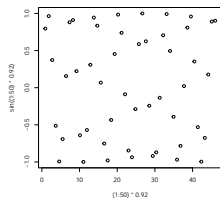
```
> n<-100
> x <- (0:n)*2*pi/100
> y <- sin(x)+rnorm(n+1)
> plot(y~x)
> lines(x,sin(x))
```



Modifying the Layout

- `par()` modifies layouts, e.g., margin sizes, line widths and types, colors, clipping, character sizes and fonts

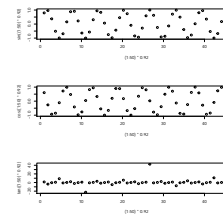
```
> plot( (1:50)*0.92, sin( (1:50)*0.92))
```



Modifying the Layout

- Now we plot the same figure again with a modified layout. In addition, two new figures are plotted

```
> par(mfrow=c(3,1))
> plot( (1:50)*0.92, sin( (1:50)*0.92))
> plot( (1:50)*0.92, cos( (1:50)*0.92))
> plot( (1:50)*0.92, tan( (1:50)*0.92))
```



Importing from Text Files

- ▶ `read.table()` is an easy to use method to importing data from a simple text file

- ▶ Simple test file, say "simple.txt":

```
x y
1 2
2 4
3 6
4 8

> df.simple <- read.table("simple.txt", header = TRUE)
> df.simple
```

```
x y
1 1 2
2 2 4
3 3 6
4 4 8
```

- ▶ The result of the `read.table()` is a data frame.

Exporting to Text Files

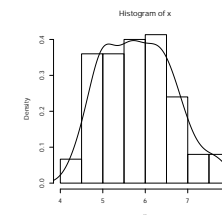
- ▶ `write.table()` prints its required argument `x` (after converting it to a data frame if it is not one nor a matrix) to a file or connection
- ▶ `write.csv()` and `write.csv2()` provide convenience wrappers for writing CSV files

Exploratory data analysis (EDA): Overview

- ▶ Idea: *let the data speak for themselves*
- ▶ Use of human brain's abilities as a pattern recognition device
- ▶ Reveal new information ("playing trumpet to the tulips")
- ▶ Ways how explore data prior to a formal analysis
- ▶ Standard tools:
 - ▶ Histograms and density plots
 - ▶ Stem-and-leaf plots
 - ▶ Scatter plots
 - ▶ Lattice: lowess smoother, trellis graphics
- ▶ Histograms: graphical representations of the frequency distribution of sets of data
- ▶ Areas of the plotted rectangles proportional to the number of observations with values within rectangle width
- ▶ Add density curves, they do not rely on breakpoints

Histograms and Density Plots

```
> data(iris)
> x <- iris$Sepal.Length
> dens <- density(x)
> hist(x,freq=F)
> lines(dens)
```



Stem-and-leaf plots

- ▶ The stem is on the left, leaves are on the right
- ▶ Smallest value reads 42. The value 44 appears four times

```
> stem(iris$Sepal.Length)
The decimal point is 1 digit(s) to the left of the |
```

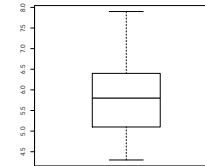
```
42 | 0
44 | 0000
46 | 000000
48 | 0000000000
50 | 000000000000000000
52 | 00000
54 | 00000000000000
56 | 0000000000000000
58 | 0000000000
60 | 00000000000000
62 | 00000000000000
64 | 00000000000000
66 | 0000000000
68 | 0000000
70 | 00
72 | 0000
74 | 0
76 | 00000
78 | 0
```

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Boxplots

- ▶ Boxplots summarize graphically the following information:
 - ▶ Outliers
 - ▶ Smallest and largest value (outliers excluded)
 - ▶ Lower and upper quantile
 - ▶ Median

```
> boxplot(iris$Sepal.Length)
```

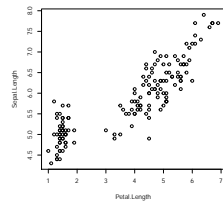


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Scatterplots

- ▶ Simple but effective tool for the analysis of pairwise relationships

```
> plot(Sepal.Length~Petal.Length,data=iris)
```



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What to Look for in Plots: Outliers

- ▶ Points that appear to be isolated from the main region of the data are called outliers
- ▶ Outliers can distort models to be fit to the data
- ▶ But there is no general definition for outliers
- ▶ This definition depends on our view of the data
- ▶ Boxplots are useful to detect outliers in one dimension, scatterplots are useful in two dimensions
- ▶ However, sometimes outliers will be apparent only in three or more dimensions.

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What to Look for in Plots

- ▶ Asymmetry
 - ▶ Most asymmetric distributions are positively or negatively skewed
 - ▶ Positively skewed distributions can be characterized as follows: There is a long tail to the right, values near the minimum are bunched up together, and the largest values are widely dispersed
- ▶ Different variabilities
 - ▶ Sometimes variability increases as data values increase
 - ▶ Then the logarithmic transformation can be helpful
- ▶ Clustering
 - ▶ Outliers can be considered as a special form of clustering
 - ▶ Clusters may suggest structures in the data which may or may not have been expected
 - ▶ Scatterplots can be useful to detect clusters.
- ▶ Non-linearity
 - ▶ Linear models should not be fitted to data where relationships are non-linear

Distributions

Example (Binomial distribution)

- ▶ Determine five random numbers following a binomial (100, 1/5) distribution

```
> set.seed(123)
> rbinom(5, size = 100, p=1/5)
[1] 18 23 19 25 26
```

- ▶ Hundred samplings with replacement from a box with 64 black and 16 red balls. Probability of drawing a red ball is $p = 16/(64 + 16) = 1/5$.

- ▶ Probability that ten red balls are drawn, i.e., $P(X = 10)$

```
> dbinom(10,100,1/5)
[1] 0.00336282
```

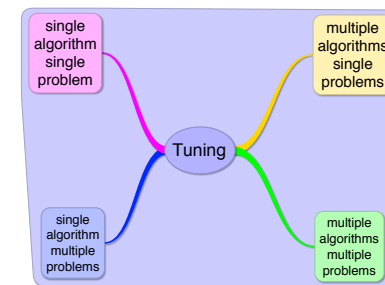
- ▶ CDF, i.e., compute $P(X \leq 10)$

```
> sum(dbinom(0:10, 100, 1/5))
[1] 0.005696381
> pbinom(10,100,1/5)
[1] 0.005696381
```

A Taxonomy of Algorithm and Problem Designs

- ▶ Classify parameters
- ▶ Parameters may be *qualitative*, like for the presence or not of an recombination operator or *numerical*, like for parameters that assume real values
- ▶ Our interest: understanding the contribution of these components
- ▶ Statistically speaking: parameters are called *factors*
- ▶ The interest is in the effects of the specific *levels* chosen for these factors

Problems and Algorithms



- ▶ How to perform comparisons?
- ▶ Adequate statistics and models?

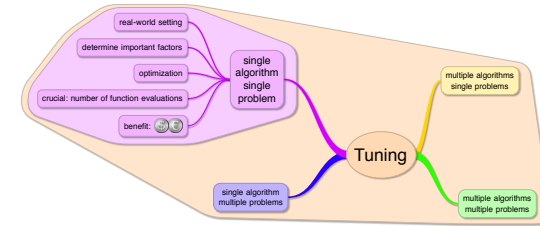
SASP: Algorithm and Problem Designs

- ▶ Basic design: assess the performance of an *optimization algorithm* on a single problem instance π
- ▶ Randomized optimization algorithms \Rightarrow performance Y on one instance is a random variable
- ▶ Experiment: On an instance π algorithm is run r times \Rightarrow collect sample data Y_1, \dots, Y_r (independent, identically distributed)
- ▶ One instance π , run the algorithm r times $\Rightarrow r$ replicates of the performance measure Y , denoted by Y_1, \dots, Y_r
- ▶ Samples are conditionally on the sampled instance and given the random nature of the algorithm, independent and identically distributed (i.i.d.), i.e.,

$$p(y_1, \dots, y_r | \pi) = \prod_{j=1}^r p(y_j | \pi). \quad (1)$$

- ▶ Y might be described by a probability density/mass function $p(y | \pi)$

SASP – Single Algorithm, Single Problem



SAMP: Algorithm and Problem Designs

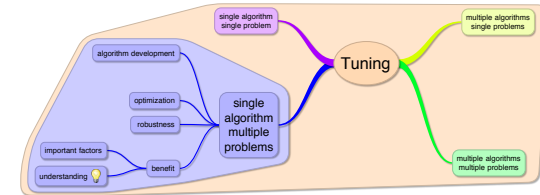
- ▶ Multiple problem instances occur if optimization problems have a set of input data which instantiate the problem
- ▶ Experiment: collect sample data Y_1, \dots, Y_R (independent, identically distributed)
- ▶ Goal: Drawing conclusions about a certain *class* or *population* of instances Π
- ▶ Single algorithm, multiple problems: performance Y of the algorithm on the class Π is described by the probability function

$$p(y) = \sum_{\pi \in \Pi} p(y | \pi) p(\pi), \quad (2)$$

with $p(\pi)$ being the probability of sampling instance π

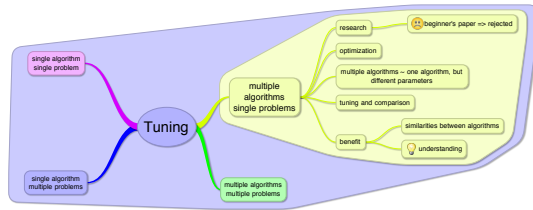
- ▶ In other terms, we are interested in the distribution of Y marginalized over the population of instances

SAMP – Single Algorithm, Multiple Problems



MASP: Algorithm and Problem Designs

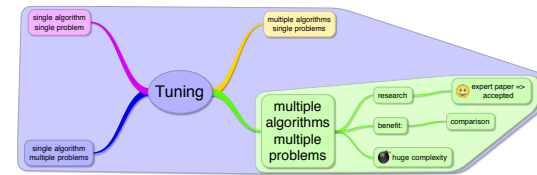
- ▶ Several optimization algorithms are compared on one fixed problem instance π
- ▶ Experiment: collect sample data Y_1, \dots, Y_R (independent, identically distributed)
- ▶ Goal: comparison of algorithms on one (real-world) problem instance π



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MAMP: Fixed Algorithm and Problem Designs

- ▶ Typically:
 - ▶ Take a few, *fixed* instances for the problem at hand
 - ▶ Collect the results of some runs of the algorithms on these instances
- ▶ Statistically, instances are also *levels of a factor*
- ▶ Instances treated as *blocks*
- ▶ All algorithms are run on each single instance
- ▶ Results are therefore *grouped* per instance



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MAMP: Randomized Problem Designs

- ▶ Sometimes, several hundred (or even more) problem instances to be tested \Rightarrow interest not just on the performance of the algorithms on a few specific instances, but rather on the generalization of the results to the entire population of instances
- ▶ Procedure: instances are chosen at random from a large set of possible instances of the problem
- ▶ Statistically, instances are also *levels of a factor*
- ▶ However, factor is of a different nature from the fixed algorithmic factors described above
- ▶ Levels are chosen at random and the interest is not in these specific levels but in the population from which they are sampled
- ▶ \Rightarrow levels and the factor are *random*
- ▶ This leads naturally to a mixed model [4]

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MAMP: Randomized Problem Designs

- ▶ Organize our presentation in different cases according to the number and type of factors involved
- ▶ Identify the cases with the following notation:

$$\left\langle \begin{matrix} \text{algorithm} \\ \text{factors} \end{matrix}, \begin{matrix} \text{number of} \\ \text{instances} \end{matrix} \left(\begin{matrix} \text{instance} \\ \text{factors} \end{matrix} \right), \begin{matrix} \text{number of} \\ \text{runs} \end{matrix} \right\rangle.$$

- ▶ Lower-case letters when referring to the number of factors, upper-case letters when referring to the number of levels
- ▶ Dash (-) indicates absence of fixed factors, round parenthesis indicates nesting
- ▶ Example: $\langle N, q(M), r \rangle$ means N algorithmic factors, q instances sampled from each combination of M instance factors, and r runs of the algorithm per instance

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MAMP: Nested Linear Mixed Models

- ▶ In statistics, the effects described are modeled as linear combinations, and mathematical theory has been developed to make inferences about the populations on the basis of the results observed in the samples.
- ▶ The mixed nature of the factors leads to so-called *nested linear mixed models*
- ▶ Nontrivial designs, go beyond the classical multifactorial ANOVA, where all factors are instead treated as fixed
- ▶ Mathematical formula involved and the inference derived are different in the case of mixed-effects models and this may lead to a different inference
- ▶ [4] give an example where this difference clearly arises

Algorithm and Problem Designs

- ▶ Classify parameters
 - ▶ Continuous, categorical, etc.
- ▶ Designs
 - ▶ Factorial, fractional factorial, space filling, etc.
- ▶ Models
 - ▶ ANOVA, regression, kriging, tree-based models, etc.
- ▶ R packages for experimental designs: Groemping's CRAN Task View: Design of Experiments (DoE) & Analysis of Experimental Data
<http://cran.r-project.org/web/views/ExperimentalDesign.html>

Summary: A Taxonomy of Algorithm and Problem Designs

- ▶ Taxonomy combining ideas from [1] and [4]
- ▶ Experimental design notation:

$$\left\langle \begin{matrix} \text{algorithm} \\ \text{factors} \end{matrix}, \begin{matrix} \text{number of} \\ \text{instances} \end{matrix} \left(\begin{matrix} \text{instance} \\ \text{factors} \end{matrix} \right), \begin{matrix} \text{number of} \\ \text{runs} \end{matrix} \right\rangle.$$

- ▶ Case $\langle -, q(-), r \rangle$: Random-Effects Design: one algorithm is evaluated on q instances randomly sampled from a class Π
- ▶ Case $\langle N, q(-), r \rangle$: Mixed-Effects Design: h algorithms are evaluated on q instances randomly sampled from a class Π
- ▶ Case $\langle 1, 1(1), r \rangle$: Fixed-Effects Design: one algorithm is evaluated r times on one fixed instance π
- ▶ ...

Comparison of Two Simulated Annealing Parameter Settings

- ▶ Case $\langle 2, 1(1), r \rangle$: Fixed-Effects Design: one algorithm is evaluated on one instance π (fixed), i.e., SASP

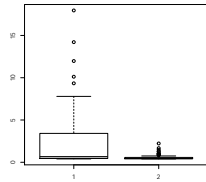
```
> set.seed(123)
> library(SPOT)
> fn <- spotBraninFunction #test function to be optimized by SANN
> x0 <- c(-2,3) #starting point that SANN uses when optimizing Branin
> maxit <- 100 #number of evaluations of Branin allowed for SANN
> temp <- 10
> tmax <- 10
> n <- 100
> y <- rep(1,n)
> y0<-sapply(y, function(x) x<-optim(par=x0, fn=fn, method="SANN"
+                               , control=list(maxit=maxit,
+                               temp=temp, tmax=tmax))$value)
> temp <- 4
> tmax <- 62
> y <- rep(1,n)
> y1<-sapply(y, function(x) x<-optim(par=x0, fn=fn, method="SANN"
+                               , control=list(maxit=maxit,
+                               temp=temp, tmax=tmax))$value)
```

Comparison: Simple EDA Using Boxplots

```
> summary(y0)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
0.3984  0.4444   0.6587   2.2770   3.4020  17.9600

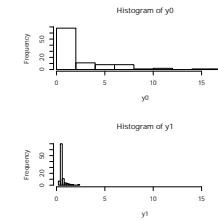
> summary(y1)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
0.3985  0.4150   0.4439   0.5609   0.5736   2.2250

> boxplot(y0,y1)
```



Comparison: Simple EDA Using Histograms

```
> par(mfrow=c(2,1))
> hist(y0,xlim = c( min(y0,y1), max(y0,y1)))
> hist(y1,xlim = c( min(y0,y1), max(y0,y1)))
> par(mfrow=c(1,1))
```



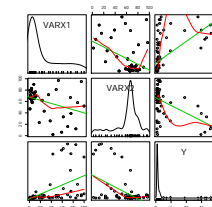
Simple EDA: Let the Data Speak

```
> df1 <- read.table("Data.d/NULL.res",header=T)
> options(width=80)
> df1[1:10,]
```

	Function	XDIM	YDIM	STEP	SEED	CONFIG	VARX1	VARX2	Y
1	UserSuppliedFunction	2	1	0	1234	1	23	62	0.5183426
2	UserSuppliedFunction	2	1	0	1235	1	23	62	0.4020790
3	UserSuppliedFunction	2	1	0	1234	2	62	33	3.5002149
4	UserSuppliedFunction	2	1	0	1235	2	62	33	16.6525805
5	UserSuppliedFunction	2	1	0	1234	3	38	26	4.7735424
6	UserSuppliedFunction	2	1	0	1235	3	38	26	0.3987177
7	UserSuppliedFunction	2	1	0	1234	4	70	16	1.4725001
8	UserSuppliedFunction	2	1	0	1235	4	70	16	18.1272253
9	UserSuppliedFunction	2	1	0	1234	5	8	90	0.5871467
10	UserSuppliedFunction	2	1	0	1235	5	8	90	0.6200017

Analysis: Simple EDA Using Scatterplots

```
> library(car)
> scatterplotMatrix(~VARX1+VARX2+Y, reg.line=lm, smooth=TRUE,
+ spread=FALSE, span=0.5, diagonal = 'density', data=df1)
```



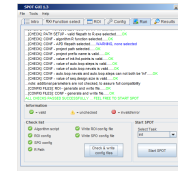
Sequential Parameter Optimization SPO

Use statistical techniques and methods from design of experiment to solve optimization problems.

1. Take initial samples from design space and evaluate on target function/algorithm
2. Build surrogate model (Linear, Tree-based, Kriging, ...) based on known evaluations
3. Determine promising new solutions with model
4. Evaluate new solutions
5. If termination criterion not reached: go to 2.
6. Summarize Results / Create Report

SPO Toolbox (SPOT)

- ▶ Currently maintained and developed as an R-Package
- ▶ Interfaces to several other R-packages
- ▶ Provides Demos and Documentation
- ▶ Graphical User Interface
- ▶ Alternative version is available for matlab



SPOT: Installation, Help, Demos

- ▶ Install from CRAN:
`> install.packages("SPOT")`
- ▶ Load package to Workspace:
`> require("SPOT")`
- ▶ Get help on some spot functions
`> ?spot`
`> ?spotOptim`
- ▶ Get a list of SPOT demos
`> demo(package="SPOT")`
- ▶ Run a SPOT demo
`> demo("spotDemo18ForresterOptim", ask=F)`
- ▶ Start the GUI
`> spotGui()`

Applications: algorithms tuned by SPOT

- ▶ Several types of evolution strategies
- ▶ Time series prediction and anomaly detection
- ▶ Classification
- ▶ Symbolic Regression
- ▶ Simulated Annealing
- ▶ For more applications see [2]

Simulated Annealing SANN

- ▶ Randomized optimization algorithm
- ▶ Two parameters: starting temperature TEMP and number of function evaluations at each temperature TMAX
- ▶ implementation used: optim, part of R-base

```
> #Find minimum of 2D-sphere function with SANN
> fn<-function(x){return(sum(x^2))}
> result<-optim(par=c(2,-4),fn,method="SANN")
> result$value
```

```
[1] 0.0002277956
```

```
> result$par
```

```
[1] 0.006771081 0.013488814
```

Tuning SANN: Define Problem to solve

- ▶ Target function: Branin-Function (2-D function with three global minima)

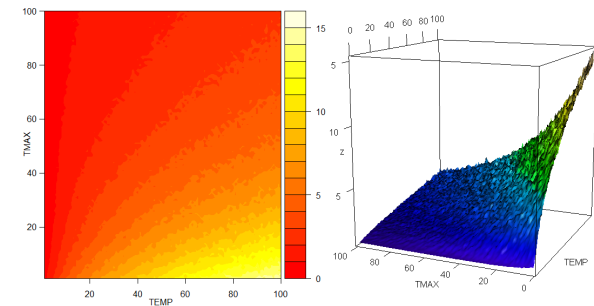
```
> require(SPOT)
> fn <- spotBraninFunction #test function to be optimized by SANN
> x0 <- c(-2,3) #starting point that SANN uses when optimizing Branin
> maxit <- 100 #number of evaluations of Branin allowed for SANN
> testalgorithm <- function(pars,x0,fn,maxit){
+   temp<-pars[1]
+   tmax<-pars[2]
+   y <- optim(x0, fn, method="SANN",
+     control=list(maxit=maxit,
+       temp=temp, tmax=tmax))
+   return(y$value)
+ }
```

SANN sweep

- ▶ Since this is a simple test problem: Complete sweep
- ▶ Understand underlying fitness shape
- ▶ 1000 repeats for each setting (takes rather long)

```
> target <- function(x,y,x0,fn,maxit){
+   zz<-matrix(0,length(x))
+   repeats=1000
+   for(i in 1:repeats){
+     set.seed(i)
+     zz =zz + apply(cbind(x,y),1,testalgorithm,x0=x0,fn=fn,maxit=maxit)
+   }
+   return(zz/repeats)
+ }
> x <- seq(1, 100, length.out = 100)
> y <- x
> z <- outer(x, y, target,x0=x0,fn=fn,maxit=maxit)
> filled.contour(x, y, z, color.palette=heat.colors,xlab="temp",ylab="tmax")
> pal <- topo.colors(100)
> require(rgl)
> persp3d(x,y,z,col=pal[cut(z,100)],xlab="TEMP",ylab="TMAX")
```

Plots from sweep



Tuning SANN: Configure SPOT

- ROI: Region of interest, in which parameters are tuned
- Surrogate: Kriging based on Forrester et. al. [6]
- Settings are minimalistic (uses a lot of default values)

```
> roi<-spotROI(c(1,1),c(100,100),type=c("INT","INT"))
> config<-list(alg.func=testalgorithm,
+   alg.ROI=roi,
+   init.design.size=20,
+   seq.predictionModel.func="spotPredictForrester",
+   seq.predictionOpt.func="spotPredictOptMulti",
+   seq.predictionOpt.method="cmaes",
+   seq.predictionOpt.budget=1000,
+   report.func="spotReportSens",
+   spot.fileMode=T,
+   io.verbosity=3,
+   auto.loop.nevals=100)
```

Tuning SANN: Run SPOT

- Pass configuration to SPOT
- Pass additional parameters to SPOT, needed by target function

```
> res<-spot(spotConfig=config,x0=x0,fn=fn,maxit=maxit)
```

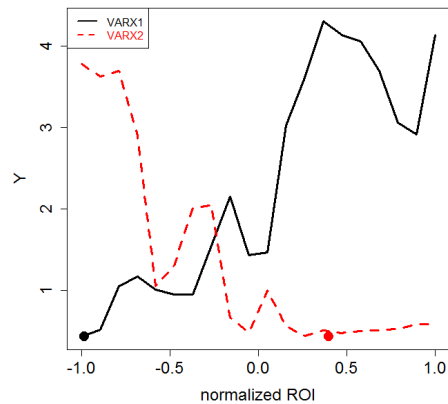
Sensitivity plot for this ROI:

	lower	upper	type	BEST
VARX1	1	100	INT	1.84744
VARX2	1	100	INT	70.36899

Best solution found with 103 evaluations:

	Y	VARX1	VARX2	COUNT	CONFIG
245	0.409769	1.84744	70.36899	5	24

Standard deviation of best solution:
 0.409769033349987 +- 0.0112398099327513



Tuning SANN: Raw results

- Result file, logged information separated by space

```
Function XDIM YDIM STEP SEED CONFIG VARX1 VARX2 Y
UserSuppliedFunction 2 1 0 1234 1 23 62 0.518342556082896
UserSuppliedFunction 2 1 0 1235 1 23 62 0.402079045134601
UserSuppliedFunction 2 1 0 1234 2 62 33 3.50021485407806
```

- Results in R command line

```
str(res$alg.currentResult)

'data.frame': 103 obs. of 9 variables:
 $ Function: Factor w/ 1 level "UserSuppliedFunction": 1 1 1 1 1 1 1 1 1 ...
 $ XDIM : num 2 2 2 2 2 2 2 2 2 ...
 $ YDIM : int 1 1 1 1 1 1 1 1 1 ...
 $ STEP : int 0 0 0 0 0 0 0 0 0 ...
 $ SEED : num 1234 1235 1234 1235 1234 ...
 $ CONFIG : int 1 1 2 2 3 3 4 4 5 ...
 $ VARX1 : num 23 23 62 62 38 38 70 70 8 8 ...
 $ VARX2 : num 62 62 33 33 26 26 16 16 90 90 ...
 $ Y : num 0.518 0.402 3.5 16.653 4.774 ...
```

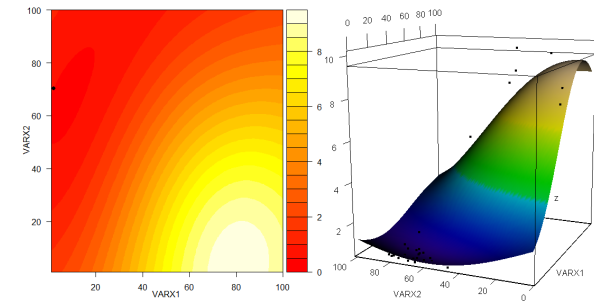
Tuning SANN: Other report functions

- ▶ Other reports/graphics can be created
- ▶ `spotReportContour` for a contour plot


```
> spot(spotConfig=append(list(
+   report.func="spotReportContour",
+   report.interactive=F),
+   res),
+   spotTask="rep")
```
- ▶ `spotReport3d` for 3d plot


```
> spot(spotConfig=append(list(
+   report.func="spotReport3d",
+   report.interactive=F),
+   res),
+   spotTask="rep")
```

Plots from SPOT



Existing features

- ▶ Single and multi criteria optimization
- ▶ Automated tuning, or manual steps
- ▶ modular concept: Use different combinations of models / methods
- ▶ Available surrogate models: Linear, Tree, Kriging, Support Vector Machine, Random Forest, ...
- ▶ Tuning real valued parameters as well as factors (i.e. with tree-based models)
- ▶ User can use custom models
- ▶ Different means of budget allocation
- ▶ Logging and Report generation

Development

- ▶ Extend report functions
- ▶ Implementation of ensembles of surrogate models
- ▶ Improve multi criteria optimization
- ▶ Adaptive ROI
- ▶ New test problems or applications

Overview

We will focus on:

- ▶ Available optimization algorithms
- ▶ Benchmarking resources

But first a very short glimpse on our targets...

Optimization Algorithms

CRAN Task View: Optimization and Mathematical Programming
<http://cran.r-project.org/web/views/Optimization.html>

- ▶ Huge list of available algorithms
- ▶ Also: Mathematical programming solvers
- ▶ We focus on (some) general purpose continuous solvers
- ▶ You can also deliver your implementations there (to Stefan Theussl)

The Adaptability Perspective

When adapting algorithms to a problem (or multiple), two things are of basic interest [11]:

- ▶ How good do we get?
- ▶ How long does it take to get there?

What to do with that?

- ▶ We can expect that different algorithms have different properties
- ▶ It depends on the optimization context which one is more important (algorithm selection problem)
- ▶ We encourage to further look at these aspects (together)

Evolutionary Methods Packages

- [cmaes](#) Covariance matrix adaptation evolution strategy
- [genalg](#) Genetic algorithm
- [rgenoud](#) GA plus quasi-Newtonian approach hybridization
- [pso](#) Particle swarm optimization
- [DEoptim](#) Differential evolution

Other Interesting Methods

- `optim` (built-in function of the stats package)
 Broyden-Fletcher-Goldfarb-Shanno (BFGS) method,
 bounded BFGS, conjugate gradient, Nelder-Mead, and
 simulated annealing (SANN)
- `optimx` new common frame for `optim()` methods and many more,
 e.g. `bobyqa`, `uobyqa`, and `newuoa`
- `nloptr` supports several global optimization routines (e.g.
 DIRECT), local derivative-free and gradient-based (e.g.
 BFGS) methods used as subroutines

And many more, even interfaces to solvers (COIN-OR, CPLEX)

BBOB Function Overview

Function groups:

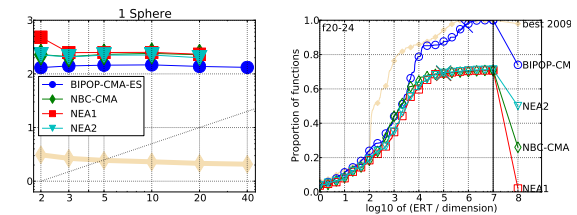
- ▶ Separable (sphere, ellipsoidal, Rastrigin, Büche-Rastrigin, linear slope)
- ▶ Low or moderate conditioning (attractive sector, step ellipsoidal, Rosenbrock original, Rosenbrock rotated)
- ▶ High conditioning, unimodal (ellipsoidal, discus, bent cigar, sharp ridge, different powers)
- ▶ Multi-modal with global structure (Rastrigin, Weierstrass, Schaffers F7, Schaffers F7, moderately ill-conditioned, Composite Griewank-Rosenbrock F8F2)
- ▶ Multi-modal with weak global structure (Schwefel, Gallagher's Gaussian 101-me Peaks, Gallagher's Gaussian 21-hi Peaks, Katsuura Function, Lunacek bi-Rastrigin)

Benchmarking: BBOB

Black-Box Optimization Benchmarking (BBOB) 2012 library
<http://coco.gforge.inria.fr/doku.php?id=bbob-2012>
 (see the GECCO workshop)

- ▶ 24 selected problems
- ▶ Interfaces from Matlab, C, Java, R, Python
- ▶ Lots of already existing results to compare with (BBOB 2009, BBOB 2010)
- ▶ Very powerful visualization for free (Python-based post-processing)
- ▶ You can also just use the problems

BBOB Sample Graphics



Real-World Problems

Noisy real-world test cases (as e.g. used in [12])

`http://ls11-www.cs.tu-dortmund.de/rudolph/kriging/applications`

Currently available:

- ▶ Gaming related: Car setup optimization (related to the former competition)
- ▶ Hydrogeologic Testcase: well placement
- ▶ More to come (hopefully)
- ▶ If you have other interesting problems, let us know

Acknowledgments

- ▶ This work has been supported by the Federal Ministry of Education and Research (BMBF) under the grants FIWA (AIF FKZ 17N1009) and CIMO (FKZ 17002X11)



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