R notes

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1 Basic

```
### Time
  proc.time()
  system.time()

### debug # c, n, where, Q;
  # options(error = recover)
  debug()
  browser()
```

```
trace()
 recover()
  traceback()
### Search in environment:
 ?ls; ?objects # List Objects
 ?search # Give Search Path for R Objects
 ?apropos; ?find # Find Objects by (Partial) Name
 ?get # Return the Value of a Named Object
### Object analysis:
 x <- matrix(1:4-0.5,2,2)
 class(x) # matrix
mode(x) # numeric
 storage.mode(x) # double
 typeof(x) # double
 attributes(x) # dim [1] 2 2 names(x) # NULL
 terms(lm(Sepal.Length ~ Sepal.Width, data=iris)) # For models
 # S3
                                         # the list of all S3 methods
 methods(class="matrix")
 methods("summary")
                                  # functions of `summary` by class
 getS3method("summary", "matrix") # Get the body of the function
 getClass("matrix")
 # S4
 library("Matrix")
 mat <- Matrix(1:6, 3, 2) # S4 object
 getClass(class(mat)) # or `showClass(class(mat))`
 getSlots(class(mat)) # or `slotNames(mat)`
 mat@Dim
 methods(class = "Matrix")
 showMethods(class = "Matrix")
### Set operations:
 union(x, y)
 intersect(x, y)
 setdiff(x, y)
 setequal(x, y)
 is.element(el, set)
 subset(old,logical)
### Index, match
 a %in% b
 match()
 which()
### Loopless
           # Apply a Function Over a Ragged array. General, good and fast
 tapply()
 by() # `tapply` for `data.frame` (or matrix)
 aggregate() # Compute Summary Statistics of Data Subsets
 apply() # Apply Functions Over Array Margins
 mapply() # Apply a Function to Multiple List or Vector Arguments
 outer() # Outer Product of Arrays
 Vectorize() # Vectorize a Scalar Function
 # Apply a Function over a List or Vector:
 lapply()
 sapply()
 replicate()
```

```
### Sequences
 rep(1:4, each = 2) # 1 1 2 2 3 3 4 4
 rep(1:4, c(2,1,2,1)) # 1 1 2 3 3 4
 seq(1, 9, by = pi) # 1.000000 4.141593 7.283185
sequence(c(3,2)) # 1 2 3 1 2
 sequence(c(3,2)) # 1 2 3 1
seq_along(5:10) # 1 2 3 4 5 6
 unique()
 duplicated()
### Combinatorics, possible combinations, brudforce:
 combn(x, m, FUN = NULL) # All Combinations of n Elements, Taken m at a Time (with sunction)
 expand.grid() # All combinations (with simetry)
 choose(n, k) #pasirinkimu skaicius (C is n po K)
 factorial(n)
 combinations()
                     #from gtools
 permutations() #from gtools
### Vector, matrix
 # manipulation
 subset()
 with();within()
 # Ordering
 order()
 sort.list()
 sort()
 arrange() # drom plyr package
 rank()
 # appending
 cbind()
 rbind()
 append()
 # calculation
 kronecker(X, Y, FUN = "*") # kronecer product {%x%}
 # Misc
 colSums(); rowSums()
 split(); cut() # grouping
 rev() # reverse
 embed(x,2) # laged matrix
### Functions, solutions
 optim()  # General optimisation
optimize()  # one dimension optimisation
 constrOptim() # constrain optimisation
 uniroot() # the root of monotone functions
              # Non-linear minimisation
 nlm()
                 # More robust (non-)constrained non-linear minimisation
 nlminb()
### Strings:
 paste()
 format()
 formatC()
 # Substrings (vectorised, can raplece)
 substr(x, start, stop)
 substring(text, first, last = 1000000L)
 # replacments
```

```
sub(pattern, replacement, x)
 gsub(pattern, replacement, x)
 # Find patern
 grep(pattern, x);
 grepl(pattern, x)
 # find expresion
 regexpr(pattern, text)
 gregexpr(pattern, text)
 # interpretation
 eval(parse(text = "(d \leftarrow 4 + 7)"))
 substitute(expr, env)
 # misc
 strsplit("a,b;c", ",|;") # a, b, c
 nchar() # the length of string
 chartr() # traslete: L -> L
 tolower()
 toupper()
 as.numeric(as.character(f)) # factor - > number
### Files
 ?files
               # Help on low-level interface to file system
 list.files() # List files in a give directory
 file.info() # Get information about files
### Misc:
 interaction()
                 # Joining factors
 x %% y
                 #mod
 x %/% y
               #div
 identical()
 suppressMessages
### Built-in constants:
 pi;letters;LETTERS  # Pi, lower & uppercase letters, e.g. letters[7] = "g"
 month.abb;month.name # Abbreviated & full names for months
### Graphics
 help(package=graphics) # List all graphics functions
                       # Generic function for plotting of R objects
 plot()
                       # Set or query graphical parameters
 par()
 curve(5*x^3,add=T) # Plot an equation as a curve
 points(x,y)
                       # Add another set of points to an existing graph
                      # Draw arrows [see errorbar script]
 arrows()
 abline()
                      # Adds a straight line to an existing graph
                      # Join specified points with line segments
 lines()
                      # Draw line segments between pairs of points
 segments()
 hist(x)
                      # Plot a histogram of x
 pairs()
                      # Plot matrix of scatter plots
                      # Plot columns of matrices
 matplot()
 boxplot()
 ?identify
  ?device
                       # Help page on available graphical devices
  ?dev.control
 postscript()
                       # Plot to postscript file
 pdf()
                       # Plot to pdf file
 png()
                       # Plot to PNG file
```

```
jpeg()
                       # Plot to JPEG file
                       # Plot to X window
 X11()
                       # Draws perspective plot
 persp()
 contour()
                      # Contour plot
                       # Plot an image
 image()
 dev.off()
 x= recordPlot()
                                           #save the current plot device output in the object x
 replayPlot(x)
                                          #replot object x
                                     #specify where multiple graphs go on the page
 layout (mat)
                                     #experiment with the magic code from Paul Murrell to do fancy graphi
 layout(rbind(c(1, 1, 2, 2, 3, 3),
              c(0, 4, 4, 5, 5, 0)))
 for (i in 1:5) {
   plot(i, type="n")
   text(1, i, paste("Plot", i), cex=4)
 }
### Statistical
 help(package=stats)
                       # List all stats functions
 ?Chisquare
                       # Help on chi-squared distribution functions
 ?Poisson
                       # Help on Poisson distribution functions
 help(package=survival) # Survival analysis
                       # Perform correlation test
 cumsum(); cumprod(); cummin(); cummax() # Cumuluative functions for vectors
 density(x)
                       # Compute kernel density estimates
 ks.test()
                       # Performs one or two sample Kolmogorov-Smirnov tests
 loess(); lowess()
                       # Scatter plot smoothing
                       # Calculate median absolute deviation
 mean(x); weighted.mean(x); median(x); min(x); max(x); quantile(x)
 rnorm(); runif()  # Generate random data with Gaussian/uniform distribution
                       # Perform spline interpolation
 splinefun()
 smooth.spline() # Fits a cubic smoothing spline
 sd()
                      # Calculate standard deviation
 summary(x)
                     # Returns a summary of x: mean, min, max etc.
                      # Student's t-test
 t.test()
                      # Calculate variance
 var()
 sample()
                      # Random samples & permutations
                      # Empirical Cumulative Distribution Function
 ecdf()
 qqplot()
                      # quantile-quantile plot
### Help
 ?Control
              # Help on control flow statements (e.g. if, for, while)
              # Help on operators acting to extract or replace subsets of vectors
 ?Extract
 ?Logic
              # Help on logical operators
              # Help on functions which support complex arithmetic in R
 ?Mod
 ?Paren
              # Help on parentheses
              # Help on regular expressions used in R
 ?regex
 ?Syntax
              # Help on R syntax and giving the precedence of operators
             # Help on special functions related to beta and gamma functions
 ?Special
```

2 Data

2.1 Input-Output

Main base functions:

Table 1: Usefull packages of 'data, input-output'.

package	rating	heading
foreign	9.0	Read Data Stored by Minitab, S, SAS, SPSS, Stata, Systat, Weka, dBase,
XML2R	8.0	EasieR XML data collection
xlsx	7.0	Read, write, format Excel 2007 and Excel 97/2000/XP/2003 files
XLConnect	5.0	Excel Connector for R

See also: Database

Special examples

```
text <- "
1 2 3
4 5 6
7 8 9
"
read.table(textConnection(text))

## V1 V2 V3
## 1 1 2 3
## 2 4 5 6
## 3 7 8 9
```

2.2 Manipulation

Table 2: Usefull packages of 'data, manipulation'.

package	rating	heading
plyr	9.1	Tools for splitting, applying and combining data
reshape2	9.0	Flexibly reshape data: a reboot of the reshape package
sqldf	7.0	Perform SQL Selects on R Data Frames
gdata	5.0	Various R programming tools for data manipulation
abind	5.0	Combine multi-dimensional arrays

Special examples

```
# list -> matrix
list2mt <- function(lst){</pre>
  return(do.call(rbind, lst))
# matrix -> list
mt2list <- function(mt){</pre>
        return(split(mt, row(mt)))
# sort matrix by colums.
msort <- function(mt, sortnr=1, desc=FALSE){</pre>
  if(desc) mt <- -mt</pre>
  if(length(sortnr)==1){
    return(mt[order(mt[,sortnr]),])
  }else{
    return(mt[do.call(order,mt2list(t(mt[,sortnr]))),])
# sort DF. See also: arrange {plyr}
esort <- function(df, sortvar, ...) {</pre>
        attach(df,warn.conflicts = FALSE )
        df <- df[with(df,order(sortvar,...)),]</pre>
         detach(df)
         return(df)
```

2.3 Database

Table 3: Usefull packages of 'data, database'.

package	rating	heading
RMySQL	7.0	R interface to the MySQL database
RODBC	7.0	ODBC Database Access
RSQLite	7.0	SQLite interface for R

```
## [1] "data"
dbListFields(con, "data")
                               "v"
                                            "v"
## [1] "row_names" "x"
res <- dbSendQuery(con, "select x, sum(v) as sum_v from data group by x")
head(fetch(res, -1)) # or use dbGetQuery to do everything
##
     x sum_v
## 1 A 111.2
## 2 B 200.9
## 3 C 293.6
## 4 D 304.7
## 5 E 116.7
## 6 F 187.7
dbClearResult(res)
## [1] TRUE
```

2.4 Big-Data

Table 4: Usefull packages of 'data, big-data'.

package	rating	heading
bigmemory	7.0	Manage massive matrices with shared memory and memory-mapped files
ff	7.0	memory-efficient storage of large data on disk and fast access functions
biglm	7.0	bounded memory linear and generalized linear models
data.table	7.0	Extension of data.frame

- Package 'bigmemory' is from the family of 'big data'. The whole list: bigmemory, biganalytics, bigtabulate, bigalgebra.
- Nowadays it is in very active development MapReduce jobs in Hadoop, that is implimented in 'rmr' package.

```
library("data.table") # it is very fast for big data (in proper ussing)
# ?data.table

# create
DT = data.table(x=rep(c("a","b","c"),each=3), y=c(1,3,6), v=1:9)
setkey(DT,x,y)
DT

# colums
head(DT[,v])
head(DT[,2,with=FALSE]) # 2nd column

# rows
DT[2]
DT[2,]
DT[2:3, sum(v)] # sum(v) over rows 2 and 3
```

```
DT[c(FALSE,TRUE)]
# subseting
DT[x=="b" \& y==3,] # works but is using data.table badly
DT[J("b", 3)]  # goodway
DT[!J("b", 3)]
                 # revers
DT[x=="b" & y<5,] # works but is using data.table badly
DT[J("b", 1:4)]
                  # wiht NA
                          # fill NA
DT[J("b", 1:4),roll=TRUE]
DT[J("b", 1:4),nomatch=0]
                            # remuve NA
# BY
DT[,sum(v),by=x]
DT[,sum(v),by=list(y\\\2)] # by expression
DT[,.SD[2],by=x]
                         # 2nd row of each group
DT[,list(MySum=sum(v),
        MyMin=min(v),
        MyMax=max(v)),
   by=list(x,y%%2)]
# compound query
DT[,sum(v),x][V1<20]
# adding computed colum
DT[,m:=mean(v),by=x]
```

3 Programing

3.1 Debuging

```
##### Debuging. RStudio has quite good debuging tools for deep debuging.
    # The strategy there to start
    # 1) Then the error occurs use
    traceback()
    # 2)Use
    options(error=recover);
    # or
    options(error=browser);
    # (after debuging set `options(error=NULL);` )
    # and rerun the code. In error you can start exploring.
    # 3) Finily use
    with_debug(install()) # in package developing (with devtools)
    debug()
   browser()
    # or break points in RStudio for deap investigation.
    # 4) for source code and scripts use
    findLineNum()
    setBreakpoint()
```

3.2 Compiling

```
##### Compiling. Good for looping or smth.
    # If caling external function - have no speed benefit.
    # Be aware of recursion - must compile with same name (prefered) or resursion sould use `Rcall'
    # Compiling with same name is OK
library("compiler")
# ?cmpfun
### Matrix multiplication with a lot of loops
MM <- function(A, B){
  if(dim(A)[2]!=dim(B)[1]) stop("incompatable")
  v \leftarrow dim(A)[2]
  ans <- matrix(NA, dim(A)[1], dim(B)[2])
  for(i in 1:dim(ans)[1]){
    for(j in 1:dim(ans)[2]){
      dum <- 0
      for(k in 1:v){
        dum \leftarrow dum + A[i, k]*B[k, j]
      ans[i,j] \leftarrow dum
  }
  ans
MMC <- cmpfun(MM)
### test
A <- matrix(1:6, 2, 3)
B \leftarrow matrix(1:12, 3, 4)
all.equal(A %*% B, MM(A, B))
## [1] TRUE
all.equal(A %*% B, MMC(A, B))
## [1] TRUE
### speed
require("rbenchmark")
## Loading required package: rbenchmark
A <- matrix(rnorm(20*10), 20, 10)
B <- matrix(rnorm(10*30), 10, 30)
benchmark(MM(A, B), MMC(A, B), A%*%B)
##
          test replications elapsed relative user.self sys.self user.child sys.child
## 3
                               0.001
                                                   0.004
                                                             0.000
       A %*% B
                         100
                                            1
                                                                            0
                                                                                       0
## 1 MM(A, B)
                         100
                               0.709
                                           709
                                                   0.696
                                                             0.012
                                                                            0
                                                                                       0
## 2 MMC(A, B)
                         100
                               0.282
                                           282
                                                   0.276
                                                             0.000
                                                                            0
                                                                                       0
### Function compiling it self - it is OK
MM <- cmpfun(MM)
benchmark(MM(A, B), MMC(A, B), A%*%B)
##
          test replications elapsed relative user.self sys.self user.child sys.child
## 3
       A %*% B
                         100
                               0.001
                                            1
                                                   0.004
                                                                 0
                                                                            0
                                                                                       0
## 1 MM(A, B)
                         100
                               0.259
                                           259
                                                   0.260
                                                                 0
                                                                            0
                                                                                       0
                         100
                               0.235
                                                   0.236
                                                                 0
                                                                             0
                                                                                       0
## 2 MMC(A, B)
                                           235
```

3.3 Recursion

```
##### Compiling. Good for looping or smth.
    # 1. Avoid it, if you know alternatives
    # 2. Do not use Recall (unless you really need renaming)
    # 3. If possible use some kind memoise
    # 4. Compile (with same name)
# Very simple Fibonachi functin
fibonacci1 <- function(seq) {</pre>
  if (seq == 1) return(1);
  if (seq == 2) return(1);
  return (fibonacci1(seq - 1) + fibonacci1(seq - 2));
# same function using `Recall`
fibonacci2 <- function(seq) {</pre>
  if (seq == 1) return(1);
  if (seq == 2) return(1);
  return (Recall(seq - 1) + Recall(seq - 2)); # Do not use! it jus slow down
# Fibonachi with memose
library("memoise")
fibonacci1M <- memoise(fibonacci1) # must be with new name</pre>
# best alternative is use your own memose
fibonacci_M <- local({</pre>
  memo \leftarrow c(1, 1, rep(NA, 100000))
  f <- function(x) {</pre>
    if(x == 0) return(0)
    if(x < 0) return(NA)</pre>
    if(x > length(memo))
      stop("x too big for implementation")
    if(!is.na(memo[x])) return(memo[x])
    ans <- f(x-2) + f(x-1)
    memo[x] <-- ans
    ans
})
require("rbenchmark")
n <- 20
benchmark(fibonacci1(n), fibonacci2(n), fibonacci1M(n), fibonacci_M(n), order=NULL)
##
                test replications elapsed relative user.self sys.self user.child sys.child
## 1 fibonacci1(n)
                              100
                                    1.445
                                               1445
                                                         1.440
                                                                       0
                                                                                  0
## 2 fibonacci2(n)
                              100
                                               1870
                                                         1.864
                                                                       0
                                                                                  0
                                                                                             0
                                     1.870
## 3 fibonacci1M(n)
                              100
                                     0.007
                                                  7
                                                         0.008
                                                                       0
                                                                                  0
                                                                                             0
                              100
## 4 fibonacci_M(n)
                                     0.001
                                                         0.000
```

3.4 Rcpp(C++)

The package Rcpp allows to impliment C++ code easily and with minimal knowledge. Main links:

- http://adv-r.had.co.nz/Rcpp.html
- $\bullet \ \, http://dirk.eddelbuettel.com/code/rcpp/Rcpp-quickref.pdf$

- http://cran.r-project.org/web/packages/Rcpp/index.html
- http://www.rcpp.org/
- http://dirk.eddelbuettel.com/code/rcpp/html/index.html

All R types are supported (vectors, functions, environment, etc ...): IntegerVector, NumericVector, LogicalVector, CharacterVector, IntegerMatrix, NumericMatrix, LogicalMatrix, CharacterMatrix.

The types that do not have C types gos only with capital letter: List, Function, Environment,..

Good reference could be fould in Rcpp-quickref.pdf. Here is very basic

```
// Geting values
  x[i]
  x(i,j)
//geting info
  .size()
  .nrow()
  .ncol()
  .length()
//Very importas is iterator, see in examples
  ::iterator
  .begin()
  .end()
//basic procedures
  .create
  .fill
  .import
  .insert
//misc
  .erase
  .eval
  .get_na
  .is_na
  .offset
  .sort
```

Very first example of making C++ in R. It is jus like 'Hellow world':

```
library("Rcpp")
cppFunction('
  int add(int x, int y, int z) {
    int sum = x + y + z;
    return sum;
  }'
)
add(1, 2, 3)
## [1] 6
```

An example of misc basics. Writing mean function in several ways. See comments:

```
sourceCpp(code=
x <- rnorm(10<sup>2</sup>)
# having error form C++:
meanC(x, type = 0)
## Error: There are no such type. (stop).
# Comparing
all.equal(mean(x), meanC(x, type=1), meanC(x, type=2), meanC(x, type=3))
## [1] TRUE
library("microbenchmark")
microbenchmark(mean(x), meanC(x, type = 1), meanC(x, type = 2), meanC(x, type = 3))
```

```
## Unit: microseconds
##
                 expr min
                              lq median
                                           uq
##
              mean(x) 5.898 6.685 6.989 7.497 39.02
## meanC(x, type = 1) 1.969 2.177 2.510 2.971 6.50
                                                      100
## meanC(x, type = 2) 1.946 2.240 2.516 3.312 13.85
                                                      100
   meanC(x, type = 3) 1.958 2.268 2.654 3.131 15.55
                                                      100
head(x)
## [1] 1000.00000
                    0.18463 -0.01969
                                       -1.08211
                                                              -1.41652
                                                    0.55824
```

A quickref for matrix

```
SEXP x;
NumericMatrix xx(x);
// Matrix of 4 rows & 5 columns (filled with 0)
NumericMatrix xx(4, 5);
// Fill with value
int xsize = xx.nrow() * xx.ncol();
for (int i = 0; i < xsize; i++) {</pre>
  xx[i] = 7;
// Same as above, using STL fill
std::fill(xx.begin(), xx.end(), 8);
// Assign this value to single element
// (1st row, 2nd col)
xx(0,1) = 4;
// Reference the second column
// Changes propagate to xx (same applies for Row)
NumericMatrix::Column zzcol = xx( _, 1);
zzcol = zzcol * 2;
// Copy the second column into new object
NumericVector zz1 = xx( _, 1);
// Copy the submatrix (top left 3x3) into new object
NumericMatrix zz2 = xx( Range(0,2), Range(0,2));
```

The example of using C++ librarys and iterator. The equivalent of the function findInterval.

```
sourceCpp(code=
```

```
)
x <- rnorm(10^1)
breaks <- c(-Inf, -3, -2, -1, 0, 1, 2, 3, Inf)
all.equal(findInterval(x, breaks), findInterval2(x, breaks))
## [1] TRUE
microbenchmark(findInterval(x, breaks), findInterval2(x, breaks))
## Unit: microseconds
## expr min lq median uq max neval
## findInterval(x, breaks) 2.344 2.562 2.764 2.907 15.93 100
## findInterval2(x, breaks) 1.526 1.704 1.946 2.269 23.58 100
```

An example of best abstraction: List and Function, without knowing anything C code wors just fine (but slow).

```
# Function and List: works in CPP, so you can actualy use R function in CPP,
# but it acually quite slow. Therefore, if posible use R function ir R and C funciton in C.
# But list is quite usefull - in fact it can contain any object with out knowing.
cppFunction(code='
List lapply1(List input, Function f) {
  int n = input.size();
  List out(n);
  for(int i = 0; i < n; i++) {
    out[i] = f(input[i]);
  return out;
')
foo \leftarrow function(n) 2^n + 1
microbenchmark(lapply1(1:3, foo), lapply(1:3, foo))
## Unit: microseconds
##
                 expr
                         min
                                 lq median
                                              uq
                                                    max neval
## lapply1(1:3, foo) 58.390 60.315 61.624 67.79 143.06
                                                           100
    lapply(1:3, foo) 4.672 5.423 6.621 7.21 16.42
```

An example of regular arrays(C) and compatability with R.

```
# Technicly, array in R is jus a vector with attributes. So C eqvivalnet is Vector.
# There is no eqvivalnet to get C array (but we have matrix, or 3D cube - they are special calsses)
sourceCpp(code='
#include <Rcpp.h>
using namespace Rcpp;
```

```
// REturning a list with all our toys
// [[Rcpp::export]]
List arr(NumericVector input, IntegerVector dim, IntegerVector dummy) {
  Dimension d(dim);
                                     // get the dim object
  // making R array with dimensions
                                     // create vec. with correct dims
  NumericVector array(d);
  std::copy(input.begin(), input.end(), array.begin()); // copy - it is not optimal, it just an example
  // now `array` is NumericVector, but in R it will be an array with dimesnions
  // much better way is to assign dim attribute
  input.attr("dim") = d;
  return List::create(
    _["input"] = input,
    array
          // no name
  );
1)
x = 1:8
y = arr(x, c(2,2,2), 1:5)
```

From C to R, From R to C

```
sourceCpp(code='
#include <iterator>
#include <vector>
#include <Rcpp.h>
using namespace Rcpp;
// [[Rcpp::export]]
List RC(NumericVector input) {
  // From C to R #1
  double mynum[] = \{0.5,1776,7,4\}; // Creating regular C array
                                    // Creating regualr C vector
  std::vector<double> foo;
  foo.assign(mynum, mynum + sizeof mynum / sizeof mynum[0]); //assigning values (could be done in creation
  SEXP bar = wrap (foo);
  // From C to R #2 - directly in R eqvivanlent
  int myint[] = \{1776,7,4\};
  NumericVector bar2(myint, myint + sizeof myint / sizeof myint[0]);
  // From R to C #1
  double* a = &input[0];
  std::cout << a[0] << ", " << a[1]<< ", ...\\n";
  // From R to C #2
  double a2[100];
  std::copy(input.begin(), input.end(), a2);
  std::cout << a2[0] << ", " << a2[1] << ", ...\\n";
  return List::create(
   bar,
    bar2
  );
```

```
}
')

x = 1:8
RC(x)
```

Fast linear algebra, using extra package (and it tipes, that are compatable with R)

```
sourceCpp(code='
#include <RcppArmadillo.h>
//[[Rcpp::depends(RcppArmadillo)]]
#include <Rcpp.h>
using namespace Rcpp;
// [[Rcpp::export]]
List fastLm(NumericVector yr, NumericMatrix Xr) {
    int n = Xr.nrow(), k = Xr.ncol();
    arma::mat X(Xr.begin(), n, k, false);
                                           // reuses memory and avoids extra copy
    arma::colvec y(yr.begin(), yr.size(), false);
    arma::colvec coef = arma::solve(X, y);
                                              // fit model y ~ X
    arma::colvec resid = y - X*coef;
                                               // residuals
    double sig2 = arma::as_scalar( arma::trans(resid)*resid/(n-k) );
                                               // std.error of estimate
   arma::colvec stderrest = arma::sqrt( sig2 * arma::diagvec( arma::inv(arma::trans(X)*X)) );
   return Rcpp::List::create(
       Rcpp::Named("coefficients") = coef,
       Rcpp::Named("stderr") = stderrest
    );
//[[Rcpp::export]]
arma::mat mult(arma::mat A, arma::mat B) {
  return A*B;
')
A \leftarrow matrix(1:9, 3, 3);
B <- matrix(9:1, 3, 3);
microbenchmark(A%*%B, mult(A, B))
## Unit: microseconds
##
         expr min
                       lq median uq max neval
##
       A %*% B 1.113 1.370 1.825 1.958 64.32
## mult(A, B) 2.782 3.625 4.040 4.427 30.11
```

3.5 Developing

Creating and developing packages.

Table 5: Usefull packages of 'programing, developing'.

package	rating	heading
devtools	7.0	Tools to make developing R code easier
formatR	7.0	Format R Code Automatically
roxygen2	7.0	In-source documentation for R
testthat	7.0	Testthat code. Tools to make testing fun:)
profr	7.0	An alternative display for profiling information
rbenchmark	7.0	Benchmarking routine for R

```
library("devtools")
help(package="devtools")
##### 1. Creating package folder with necesary infrastructure (use one folder up)
   # And put the content that you allready have.
    # Sugestions:
   # a) basic in-buld sunction
   package.skeleton()
   # b) devtools function
   create()
    # c) RStudio meniu
    # d) From other packages. Very usefull for optional failes.
##### 2. Codding. Start coding in R folder.
    # The loan can be done by
   load_all() # devtools
    # If you allready have some version of the package that is curently in use you
    # can isolate developing code in
    dev_mode() # devtools
##### 3. Then having starting code you should write tests to make sure everything works as
    # intended to be.
    library("testthat")
    ?test
    ?test_that
##### 4. Profiling
   Rprof
    summaryRprof
    library("profr")
   help(package="profr")
##### 5. Preparing code. Making well format, commenting (+documentation in comment with roxygen2)
    library("formatR")
   help(package="formatR")
##### 6. Documentation
  ### For functions. use one of the strategies (you can mix them, but it is not rezomended):
    # a) manual documentation
    ?prompt #Produce Prototype of an R Documentation File
    check_doc()
    # b) in-code documentation
```

```
library("roxygen2")
    ?document # Use roxygen to make documentation
    ?roxygenize
 ### Do not foget to
   # Writing ReadMe
   # writing vinigete.
   # demo
   # package description
   dev_help("Vmisc")
   build_vignettes()
##### 7. local install
   install(quick=TRUE)
   library("Vmisc")
   help(package="Vmisc")
##### 8. final scheck, and bild
   check()
   build()
    # ?release
```

3.5.1 Vignettes

- A vignettes should be write in *vignettes* package. All necessary files should be in this directory (or sub-directory).
- The vignettes could be Rn(pd) or Rd(html). Let assume it is Rnw.
- Meta data should be included in the comments. Main list:

```
%\VignetteEngine{knitr::knitr}
%\VignetteIndexEntry{Just a pdf example}
%\VignetteDepends{}
%\VignetteKeywords{string, misc}
%\VignettePackage{Vmisc}
```

- If Vignette source file is Rwn, then it could be clasical Sweave or other (e.g. knitr). If using knitr, then
 - it sould be decleard in meta-date comment (note the first line).
 - it sould be declear in DESCRIPTION file, line 'VignetteBuilder: knitr'
 - knitr package should be included in DESCRIPTION file 'Suggests' filed, e.g : Suggests: knitr, ...

4 Special topics

4.1 Search in environment

Special examples

4.2 Strings

```
### Strings:
 paste()
 format()
 formatC()
  # Substrings (vectorised, can raplece)
  substr(x, start, stop)
  substring(text, first, last = 1000000L)
  # replacments
  sub(pattern, replacement, x)
  gsub(pattern, replacement, x)
  # Find patern
  grep(pattern, x);
  grepl(pattern, x)
  # find expresion
  regexpr(pattern, text)
  gregexpr(pattern, text)
  # interpretation
  eval(parse(text = "(d <- 4 + 7)"))
  substitute(expr, env)
  # misc
  strsplit("a,b;c", ",|;") # a, b, c
  nchar() # the length of string
  chartr() # traslete: L -> L
  tolower()
  toupper()
  as.numeric(as.character(f)) # factor - > number
```

Table 6: Usefull packages of 'strings'.

package	rating	heading
evaluate	7.0	Parsing and evaluation tools that provide more details than the default
stringr	7.0	Make it easier to work with strings

Special examples

```
### Basic operations
text <- "Hellow, # comment

The line after empty line with lots of sapces.</pre>
```

```
doc <- readLines(textConnection(text))</pre>
# remove comments
doc <- gsub("#.+$", "", doc)</pre>
# remove dublicate space
doc <- gsub("\\s+", " ", doc)</pre>
# Trim the rest
doc \leftarrow gsub("^\s+|\s+$", "", doc)
# remove empty lines
doc <- doc[nchar(doc)>0]
doc
### reverse
(a = "this is a string"); paste(rev(substring(a,1:nchar(a),1:nchar(a))),collapse="")
### vecorise string substitution
(x \leftarrow c("aaaa","bbbb","cccc")); substring(x, 2) \leftarrow c("...", "+++"); x;
### Binary string concat
"%.%" <- function(x,y) paste(x,y,sep="")
"I love " %.% "R."
### String matrix spliting
SplitLines <- function(x, chMode, sep=" "){</pre>
  ats = do.call(rbind, strsplit(x, sep))
  if (!missing(chMode)){
    storage.mode(ats)<-chMode
  return(ats)
text <-c(
  "1 2 3"
  ,"4 5 6"
  ,"7 8 9"
X <- SplitLines(text, "double")</pre>
# Alternative from data input:
X <- read.table(textConnection(text))</pre>
```

5 Graphics

Table 7: Usefull packages of 'graph'.

package	rating	heading
igraph	7.0	Network analysis and visualization
ggplot2	7.0	An implementation of the Grammar of Graphics
colorspace	7.0	Color Space Manipulation
RColorBrewer	7.0	ColorBrewer palettes
scales	7.0	Scale functions for graphics
labeling	7.0	Axis Labeling
rgl	7.0	3D visualization device system (OpenGL)
lattice	7.0	Lattice Graphics
gplots	7.0	Various R programming tools for plotting data
vcd	7.0	Visualizing Categorical Data
scatterplot3d	7.0	3D Scatter Plot
plotrix	7.0	Various plotting functions
aplpack	7.0	Another Plot PACKage: stem.leaf, bagplot, faces, spin3R, plotsummary, plothulls
latticeExtra	7.0	Extra Graphical Utilities Based on Lattice
munsell	7.0	Munsell colour system
iplots	7.0	iPlots - interactive graphics for R

6 Data science

6.1 Machine and Statistical Learning

Table 8: Usefull packages of 'learning'.

package	rating	heading
e1071	7.0	Misc Functions of the Department of Statistics (e1071), TU Wien
Metrics	7.0	Evaluation metrics for machine learning
dtw	7.0	Dynamic time warping algorithms
randomForest	7.0	Breiman and Cutler's random forests for classification and regression
FNN	7.0	Fast Nearest Neighbor Search Algorithms and Applications
nnet	7.0	Feed-forward Neural Networks and Multinomial Log-Linear Models
neuralnet	7.0	Training of neural networks
RSNNS	7.0	Neural Networks in R using the Stuttgart Neural Network Simulator (SNNS)
kernlab	7.0	Kernel-based Machine Learning Lab
tree	7.0	Classification and regression trees
rpart	7.0	Recursive Partitioning and Regression Trees
cluster	7.0	Cluster Analysis Extended Rousseeuw et al
FactoMineR	7.0	Multivariate Exploratory Data Analysis and Data Mining with R
rattle	5.0	Graphical user interface for data mining in R

```
# randomForest
library("randomForest")
rf <- randomForest(train, labels)</pre>
```

```
### SVM
library("kernlab") # pasirupina gerais defoltais ir yra daugiau metodu
sv <- ksvm(train, labels)</pre>
library("e1071") # siame pakete yra visko, cia svm yra esmine realizacija
sv <- svm(train, labels, kernel="sigmoid") # linear sigmoid polynomial radial</pre>
# k-nearest nabahood
library(FNN)
PredTest_knn = knn(train, test, labels, k = 5)
# recursive partition
library("rpart")
rp <- rpart(labels ~ ., data = TrainDF)</pre>
# neuron networks
library("nnet")
nn <- nnet(labels ~ ., data = TrainDF[,1:20, 980:1001], size = 40)</pre>
library(neuralnet)
?neuralnet
```

A Drafts