R Package miscset User Manual

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1 Introduction

The package **miscset** provides miscellaneous functions to simplify R workspace and data life. For example, one can scale numbers by a range, or use enhanced functions with regular expression and string operations. Latex tables, just empty plots or arrangements of ggplots can be exported. Reshaping of matrices or applying functions by grids on data.tables is supported. In addition, workspace objects can be listed or removed more user friendly and multiple RData objects can be loaded into a list at once. For faster computation, some functions import multicore support from the parallel package. Also C++ is used for performance included by the package Rcpp. In the following examples, all available functions will be presented including explanations of their usage. Find the source code online at github.

2 Installation

To install the package call the command

```
install.packages("miscset")
```

within the R console or run R CMD INSTALL /path/to/miscset from a terminal in Linux/OSX. To use the most recent version install with:

```
if (!require(devtools))
  install.packages("devtools")
library(devtools)
install_github(repo='setempler/miscset')
```

To use it, the library or require function load the package.

```
library(miscset)
```

Part I

Usage

3 Numeric Functions

3.1 Generate triangular numbers - ntri

The function generates a series of triangular numbers of length n according to oeis.org.

```
ntri(12)
## [1] 0 1 3 6 10 15 21 28 36 45 55 66
```

3.2 Scale numeric vectors - scale0, scaler

The function scale0 scales all values in a numeric vector from 0 to 1 according to their minimum and maximum values. scaler provides an argument (r) which lets you set the range to scale a vector to. Additionally, the minimum and maximum border can be specified with the argument b, so that vectors with different ranges can be compared on the same scale.

```
do.call(rbind, list(
   scale0 = scale0(0:5),
   scale0_shift = scale0(-2:3),
```

```
scaler = scaler(0:5),
 scaler_newrange = scaler(0:5, r = 1:2),
 scaler_newbord = scaler(0:5, b = c(0, 10)))
##
                  [,1] [,2] [,3] [,4] [,5] [,6]
## scale0
                      0.2 0.4
                                0.6
                                     0.8
## scale0_shift
                    0 0.2
                            0.4
                                0.6 0.8 1.0
## scaler
                    0 0.2 0.4
                                0.6 0.8 1.0
## scaler_newrange
                    1 1.2 1.4
                                1.6 1.8 2.0
                    0 0.1 0.2 0.3 0.4 0.5
## scaler_newbord
```

3.3 p values to *** converting - p2star

With this function, one can quickly convert a vector with p values to symbols reflecting ranges of significance. symbols and breaks can be modified. See the example:

```
p2star(c(1e-5,.1,.7))
## [1] "***" "n.s."
```

4 Data and System

4.1 Apply a function on a data frame by a grid - gapply

To apply a function on a subset of a dataset, all named columns are used to create a grid for which each unique combination is used to extract the rows in the data.frame. Multicore support is implemented by mclapply. The grid can be extracted by the function levels and a row binding of elements that can be coerced to data.frames is implemented in the method as.data.frame. If a data.table is preferred, the method as.data.table is implemented and works only if the package is installed.

```
f <- function (x) c(conc.diff = diff(range(x$conc)), uptake.sum=sum(x$uptake))
d <- gapply(CO2, c('Type', 'Treatment'), f)</pre>
levels(d)
##
            Type Treatment
## 1
          Quebec nonchilled
## 2 Mississippi nonchilled
          Quebec
                    chilled
## 4 Mississippi
                     chilled
head(as.data.frame(d))
##
     conc.diff uptake.sum
                                  Type Treatment
## 1
           905
                    742.0
                                Quebec nonchilled
## 2
           905
                     545.0 Mississippi nonchilled
## 3
           905
                     666.8
                                Quebec
                                           chilled
## 4
           905
                     332.1 Mississippi
                                           chilled
```

4.2 Transform to squared matrix - squarematrix

The function squarematrix can generate a symmetric (square) matrix from an unsymmetric matrix by using the column and row names and filling empty pairs with NA.

```
M <- matrix(1:6, 2, dimnames = list(2:3,1:3))
M

## 1 2 3
## 3 2 4 6

squarematrix(M)

## 1 2 3
## 1 NA NA NA
## 2 1 3 5
## 3 2 4 6</pre>
```

4.3 Generate a pairwise list - enpaire

The function enpaire creates a pairwise list of matrix values. The result is a data.frame that contains a column for the names of each dimension and the upper and lower triangle values. Unsymmetric matrices are transformed by squarematrix (see previous section).

```
M <- matrix(letters[1:9], 3, 3, dimnames = list(1:3,1:3))
Μ
##
## 1 "a" "d" "g"
## 2 "b" "e" "h"
## 3 "c" "f" "i"
enpaire(M)
    row col lower upper
##
## 1
     1 2 b
## 2
          3
                С
                       g
          3
```

4.4 Create a latex document containing a table - textable

This function enhances the functionality of the xtable function from the similar named package. The output of xtable is captured, processed and then written to a file. The file may contain also latex header for an A4 portrait or landscape article. The function is called with the following syntax: textable(d, file, caption = NULL, label = NULL, align = NULL, rownames = FALSE, topcapt = TRUE, digits = NULL, as.document = FALSE, landscape = FALSE, margin = 2, pt.size = 10, cmd = NULL)

file is a character string with the name to the file of the function output. caption is a character string with the table's title. It is aligned to the top of the table, when topcapt is TRUE, otherwise to the table bottom. rownames is logical and allows to switch printing of row names on and off. when as.document is TRUE, a document header is added. Then, landscape defines the orientation of the page, pt.size the size of the characters and margin the table borders in cm. digits sets the number of digits to print for numeric values. With align the column alignements can be set. It is either one of 'r', 'c', 'l', or a vector of those elements for alignemt to the right, center or left, repsectively (use rep() or strplit() as support). label allows to supply a latex label for reference in form of a vector of length one.

```
## % Tue Feb 24 19:02:41 2015
##
## \documentclass[a4paper,10pt]{article}
## \usepackage[a4paper,margin=2cm]{geometry}
## \begin{document}
##
## \begin{table}[ht]
## \centering
## \caption{R dataset "trees".}
## \begin{tabular}{llrr}
     \hline
##
##
   & Girth & Height & Volume \\
##
    \hline
## 1 & 8.3000 & 70.0000 & 10.3000 \\
    2 & 8.6000 & 65.0000 & 10.3000 \\
##
     3 & 8.8000 & 63.0000 & 10.2000 \\
      \hline
##
## \end{tabular}
## \label{tab:trees}
## \end{table}
##
## \end{document}
```

In addition, a system command can be provided as a character string with cmd to create a pdf for example. An example therefore might be cmd = "pdflatex".

4.5 List details from and remove all objects - 1sall, rmall

With lsall(envir, ...) all object names, their length, class, mode and size is returned in a data.frame from a specified environment. rmall(...) removes the complete list of objects at the global environment.

```
lsall()
## Environment: R_GlobalEnv
## Objects:
     Name Length
                    Class
                               Mode Size Unit
## 1
                               list 4.9
        d
              4
                   gapply
               1 function function 2.5
## 2
        f
                                           Kb
## 3
                  matrix character 1.3
rmall()
lsall()
## Environment: R_GlobalEnv
## Objects:
## NULL
```

4.6 Load objects from R data files into a list - lload

11oad provides a way to load R objects from multiple R data files and stores all objects in a list. Thereby, the list names are respective to the R data files. Each entry consists of a sublist with as many entries as objects were loaded from the according R data file. The sublist names are the same as the object names that are stored. If simplify=TRUE is given, all object (sublist) names are checked for duplicates, and if none are found, the list is reduced in one level, dropping the file level.

5 Strings and Patterns

5.1 Prepend zeroes to unify number lengths - leading0

The function leading 0 aims to create e.g. index names with a common string length. It creates character strings from numeric values while attaching 0 in front of the number up to a certain length of total digits of each string.

```
paste0("page", leading0(8:10, 3))
## [1] "page008" "page009" "page010"
```

5.2 Extract substrings by pattern - strextr

The function strextr splits strings in a character vector by sep and extracts all substrings matching a given pattern.

```
s <- c("a1 b1 c1", "a2 b2", "aa a1", "aa", "b1 a1", "bb ab a1")
strextr(s, "^[ab][[:digit:]]$")
## [1] NA
          NA
                 "a1" NA
strextr(s, "^[ab][[:digit:]]$", mult = T)
## [[1]]
## [1] "a1" "b1"
##
## [[2]]
## [1] "a2" "b2"
##
## [[3]]
## [1] "a1"
##
## [[4]]
## [1] NA
##
## [[5]]
## [1] "b1" "a1"
##
## [[6]]
## [1] "a1"
strextr(s, "^[ab][[:digit:]]$", mult = T, unlist = T)
## [1] "a1" "b1" "a2" "b2" "a1" NA
                                     "b1" "a1" "a1"
strextr(s, "^[c][[:digit:]]$")
## [1] "c1" NA NA NA
                         NA
                                NA
```

5.3 Extract substrings by splitting - strpart

Similar to strextr the function strpart supplies a method to extract a substring, but by defining the nth part of the string split by the separator given in sep.

```
strpart(s, " ", 2)
## [1] "b1" "b2" "a1" NA "a1" "ab"
```

5.4 Reverse strings - strrev

With strrev you can create the reversed version of strings.

5.5 Multiple pattern replacement - msub, mgsub

msub and mgsub behave like sub and gsub but they replace multiple patterns. Replacement is done in order of the pattern input, and multicore support is enabled by mclapply from the parallel package.

```
## [1] "a1 b1 c1" "a2 b2"
                                          "aa"
                                                     "b1 a1"
                              "aa a1"
                                                                 "bb ab a1"
msub("A", "X", s)
## [1] "a1 b1 c1" "a2 b2"
                              "aa a1"
                                          "aa"
                                                     "b1 a1"
                                                                 "bb ab a1"
mgsub("A", "X", s)
## [1] "a1 b1 c1" "a2 b2"
                              "aa a1"
                                          "aa"
                                                     "b1 a1"
                                                                 "bb ab a1"
```

5.6 Get index of expression - gregexprind

5.7 Multiple pattern search - mgrepl

With mgrepl(patterns, text, ...) you can search for more than one regular expression, and use a logical function to combine the results for each single expression. It returns a logical vector, or when use.which = TRUE an integer vector with the indices as in the base function which.

```
## [1] "a1 b1 c1" "a2 b2" "aa a1" "aa" "b1 a1" "bb ab a1"

mgrepl(c("a","b"), s, any)

## [1] TRUE TRUE TRUE TRUE TRUE

mgrepl(c("a","b"), s, all)

## [1] TRUE TRUE FALSE FALSE TRUE TRUE

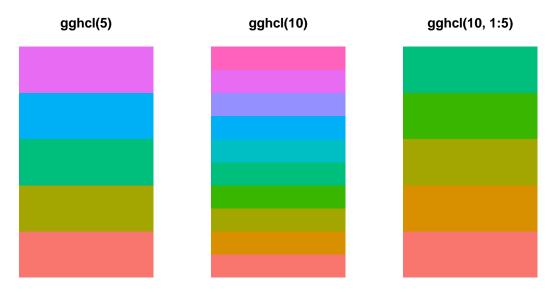
mgrepl(c("a","b"), s, all, use.which = TRUE)

## [1] 1 2 5 6
```

6 Graphics

6.1 Create a color palette - gghcl

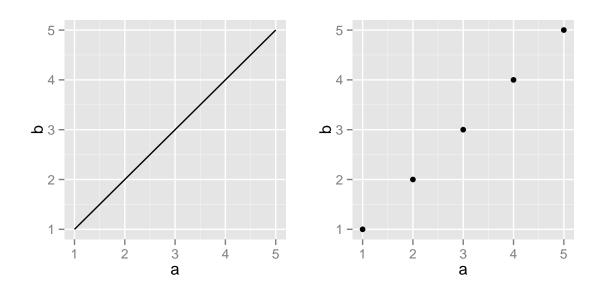
gghcl() creates color palettes. It enhances the hcl function. See some examples:



6.2 Arrange a list of ggplots in a grid - ggplotGrid, ggplotGridA4

ggplotGrid arranges your ggplot objects in a grid and optionally sends them to a graphics device such as pdf/ps/svg/png/eps for export to a file.

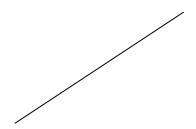
```
library(ggplot2)
d <- data.frame(a=1:5, b=1:5)
gg1 <- ggplot(d, aes(x=a, y=b)) + geom_line()
gg2 <- ggplot(d, aes(x=a, y=b)) + geom_point()
ggplotGrid(list(gg1, gg2), ncol = 2)</pre>
```



6.3 Create an empty plot - plotn

To produce nothing but a plot, use plotn:

```
plotn(1)
abline(0,1)
```



7 Deprecated

7.1 Arrange a list of ggplots in a grid - ggplotlist

The function ggplotlist is deprecated and will be dropped in future releases. Please use ggplotGrid as replacement.