Package 'aakmisc'

January 17, 2017

Type Package

Title Miscellany for the King Lab

Version 0.22-3
Date 2017-01-17
Maintainer Aaron A. King <kingaa@umich.edu></kingaa@umich.edu>
Description Miscellaneous tools, collected in one place.
<pre>URL http://github.com/kingaa/aakmisc</pre>
Contact kingaa at umich dot edu
BugReports http://github.com/kingaa/aakmisc/issues
Depends R(>= 3.0.0), plyr, reshape2, ggplot2(>= 1.0.1.9003), grid, magrittr
Imports methods, curl, DBI, RPostgreSQL
License GPL(>=3)
LazyLoad true
LazyData true
Collate aaa.R db.R tunnel.R random.R plotMatrix.R scinot.R lazy_load.R
R topics documented:
dbtools
Lazy loading of knitr caches
plotMatrix 5 random 6
scinot
Index 8

2 dbtools

dbtools

Interface with databases

Description

Interface with project databases.

Usage

```
getMLEs(host = getOption("aakmisc.dbhost", "localhost"),
        dbname = getOption("aakmisc.dbname", NULL),
        port = getOption("aakmisc.port",5432),
        user = getOption("aakmisc.user",NULL),
        ...)
recMLEs(mle, host = getOption("aakmisc.dbhost", "localhost"),
        dbname = getOption("aakmisc.dbname", NULL),
        port = getOption("aakmisc.port",5432),
        user = getOption("aakmisc.user", NULL),
        ...)
listScripts(host = getOption("aakmisc.dbhost","localhost"),
            dbname = getOption("aakmisc.dbname", NULL),
            port = getOption("aakmisc.port",5432),
            user = getOption("aakmisc.user", NULL),
            ...)
recScript(files, host = getOption("aakmisc.dbhost", "localhost"),
          dbname = getOption("aakmisc.dbname", NULL),
          port = getOption("aakmisc.port",5432),
          user = getOption("aakmisc.user", NULL),
dropScript(script, host = getOption("aakmisc.dbhost","localhost"),
           dbname = getOption("aakmisc.dbname", NULL),
           port = getOption("aakmisc.port",5432),
           user = getOption("aakmisc.user", NULL),
catScript(script, file = "", host = getOption("aakmisc.dbhost","localhost"),
          dbname = getOption("aakmisc.dbname", NULL),
          port = getOption("aakmisc.port",5432),
          user = getOption("aakmisc.user", NULL),
          ...)
getQuery(statement,
         host = getOption("aakmisc.dbhost","localhost"),
         dbname = getOption("aakmisc.dbname", NULL),
         port = getOption("aakmisc.port",5432),
         user = getOption("aakmisc.user", NULL),
writeDBTable(name, value, overwrite = FALSE, append = FALSE,
             row.names = FALSE,
```

dbtools 3

Arguments

host Hostname on which to connect to the PostgreSQL server.

dbname Name of PostgreSQL database.

port Port on which to connect to PostgreSQL database. If NULL, a random port

number will be used.

user Username to use in conneting to PostgreSQL database. If NULL, Sys.getenv("USER")

will be used.

mle A data-frame of MLEs to be recorded.

files Files containing R scripts to be recorded.

script Name of script.

file File to which the script will be written. See cat.

statement SQL statement passed to dbGetQuery.

name, value Name and contents of table to create.

overwrite, append, row.names

See dbWriteTable.

remotehost Hostname of PostgreSQL server. An ssh tunnel to this host will be created.

pid ID of ssh tunnel process. Set automatically by startTunnel.

sleep Time in seconds to sleep after initiating the ssh tunnel.
... Additional arguments will be passed to dbConnect.

Author(s)

Aaron A. King <kingaa at umich dot edu>

```
## Not run:
startTunnel()
listScripts()
stopTunnel()
## End(Not run)
```

Lazy loading of knitr caches

Functions for lazy-loading knitr caches.

Description

These functions are helpful for loading cached chunks into an interactive R session.

Use lazyload_cache_dir to load a whole directory of cached objects.

Use lazyload_cache_labels to load and explicit set of cached chunks.

Usage

Arguments

path	the path to the cache directory
labels	character vector; chunk labels to load
envir	the environment to load the objects into
ask	if TRUE, interactively ask whether to load each database discovered in path
verbose	if TRUE, display the names of chunk labels being loaded
full.names	use the full name, i.e., include the path, for the chunk label? This argument is passed to list.files.
filter	optional function; passed to lazyLoad. When called on a character vector of object names, this function should return a logical vector: objects for which this is TRUE will be loaded.
	additional arguments passed to list.files

Value

Both functions return NULL, invisibly.

Author(s)

Peter DeWitt (https://github.com/dewittpe).

plotMatrix 5

plotMatrix

A scatterplot matrix with densities on the diagonal.

Description

A special scatterplot matrix.

Usage

```
## S3 method for class 'data.frame'
plotMatrix(data, marg.exp = 0.02, labels = names(data),
    alpha = 1, pch = 16, size = unit(2,"mm"), ...)
## S3 method for class 'list'
plotMatrix(data, marg.exp = 0.02, labels = names(data),
    alpha = 1, pch = 16, size = unit(2,"mm"), ...)
## S3 method for class 'aakplot'
print(x, newpage = is.null(vp), vp = NULL, ...)
```

Arguments

```
data Data to plot.

marg.exp Fraction by which to expand the plot at the margins.

labels Names of variables plotted.

alpha, pch, size

Refer to the plotted points in the scatterplots.

optional arguments, passed to hist.

x plotMatrix object to display.

newpage logical; if TRUE, grid.newpage() will be called before the graphics are drawn.

vp viewport to use. See viewport.
```

Author(s)

Aaron A. King <kingaa at umich dot edu>

6 random

random

Functions for generating and working with truly random integers.

Description

Functions for generating and working with truly random seeds.

Usage

```
random.org(n = 10, rnd = "new")
urandom(n = 10)
rngControl(expr, seed = NULL)
rngSeeds(n, seed = NULL)
```

Arguments

n Number of integers required.

rnd random.org parameter

expr Expression to be evaluated with RNG control.

seed RNG seed.

Author(s)

```
Aaron A. King <kingaa at umich dot edu>
```

References

```
http://www.random.org
```

```
## Not run:
random.org(n=5)
seed <- urandom(n=1)
seeds <- rngSeeds(5,seed=seed)
set.seed(seed)
runif(5)</pre>
```

scinot 7

```
rngControl(runif(5), seed=seed[1])
rngControl(runif(5), seed=seed[1])
runif(5)
set.seed(seed)
runif(5)
runif(5)
## End(Not run)
```

scinot

Scientific notation.

Description

Format using scientific notation.

Usage

Arguments

x Number to format.
digits Number of significant digits in mantissa.
format Format specification. type="expression" results in an R expression. type="latex" results in a latex expression. type="math" is like "latex" but wraps the text in "\$".
simplify logical. If simplify=TRUE, then 1×10^n is simplified to 10^n .

Author(s)

Aaron A. King <kingaa at umich dot edu>

```
x <- c(0.0309595,8577676.441)
scinot(x[2],4)
scinot(x[1],2,"latex")
sapply(x,scinot,digits=3,format='math')</pre>
```

Index

```
*Topic programming
    Lazy loading of knitr caches, 4
cat. 3
catScript (dbtools), 2
dbConnect, 3
dbGetQuery, 3
dbtools, 2
dbWriteTable, 3
dropScript (dbtools), 2
getMLEs (dbtools), 2
getQuery (dbtools), 2
hist, 5
Lazy loading of knitr caches, 4
lazyLoad, 4
lazyload_cache_dir(Lazy loading of
        knitr caches), 4
lazyload_cache_labels(Lazy loading of
        knitr caches), 4
list.files, 4
listScripts (dbtools), 2
plotMatrix, 5
print.aakplot(plotMatrix), 5
random, 6
recMLEs (dbtools), 2
recScript (dbtools), 2
rngControl (random), 6
rngSeeds (random), 6
scinot, 7
startTunnel (dbtools), 2
stopTunnel (dbtools), 2
urandom (random), 6
viewport, 5
writeDBTable (dbtools), 2
```