

Package ‘aakmisc’

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Type Package

Title Miscellany for the King Lab

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Description Miscellaneous tools, collected in one place.

URL <http://github.com/kingaa/aakmisc>

Contact kingaa at umich dot edu

BugReports <http://github.com/kingaa/aakmisc/issues>

Depends R(>= 3.3.1)

Imports methods, grid, curl, DBI, RPostgreSQL, plyr, ggplot2

Suggests scales, magrittr, reshape2, readr

License GPL-3

LazyLoad true

LazyData true

Collate aaa.R db.R tunnel.R random.R plotMatrix.R scinot.R trnc.R lazy_load.R

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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,

```

```

        host = getOption("aakmisc.dbhost","localhost"),
        dbname = getOption("aakmisc.dbname",NULL),
        port = getOption("aakmisc.port",5432),
        user = getOption("aakmisc.user",NULL),
        ...)
startTunnel(port = NULL,
            remotehost = getOption("aakmisc.remotehost",NULL),
            user = getOption("aakmisc.user",NULL),
            sleep = 5)
stopTunnel(..., pid = getOption("aakmisc.tunnelpid",NULL))

```

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 connecting 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>

Examples

```

## 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 an explicit set of cached chunks.

Usage

```
lazyload_cache_dir(path = "./cache", envir = parent.frame(), ask = FALSE,  
                  verbose = getOption("verbose",FALSE), full.names = TRUE, ...)  
lazyload_cache_labels(labels, path = "./cache/", envir = parent.frame(),  
                     verbose = getOption("verbose",FALSE), filter,  
                     full.names = TRUE, ...)
```

Arguments

| | |
|-------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <code>path</code> | the path to the cache directory |
| <code>labels</code> | character vector; chunk labels to load |
| <code>envir</code> | the environment to load the objects into |
| <code>ask</code> | if TRUE, interactively ask whether to load each database discovered in path |
| <code>verbose</code> | if TRUE, display the names of chunk labels being loaded |
| <code>full.names</code> | use the full name, i.e., include the path, for the chunk label? This argument is passed to <code>list.files</code> . |
| <code>filter</code> | optional function; passed to <code>lazyLoad</code> . 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. |
| <code>...</code> | additional arguments passed to <code>list.files</code> |

Value

Both functions return NULL, invisibly.

Author(s)

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

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, <code>grid.newpage()</code> will be called before the graphics are drawn. |
| vp | viewport to use. See viewport . |

Author(s)

Aaron A. King <kingaa at umich dot edu>

Examples

```
## Not run:
x <- data.frame(a=rexp(n=1000,rate=1/3),b=rnorm(1000))
mutate(x,c=a+b^2,d=a-b^3) -> x

print(plotMatrix(x,alpha=0.2))

g <- plotMatrix(
  x[-2],
  labels=c(
    expression(alpha),
```

```

        expression(beta),
        expression(phi)
    ),
    alpha=0.3
)
print(g)

print(plotMatrix(as.list(x),alpha=0.2,breaks='scott'))

## End(Not run)

```

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>

Examples

```

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

```

```

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

```

scinot(x, digits = 2, format = c("expression", "latex", "math"),
       simplify = FALSE)

```

Arguments

| | |
|----------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| x | Number(s) 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>

See Also

[scientific](#)

Examples

```

x <- c(0.0309595, 8577676.441, 10000)
scinot(x[2], 4)
scinot(x[1], 2, "latex")
sapply(x, scinot, digits=3, format='math')
scinot(x, digits=0, simplify=FALSE)
scinot(x, digits=0, simplify=TRUE)

```

`trnc`*Truncation of plots.*

Description

Truncate to the specified window.

Usage

```
trnc(x, range=c(0,1), only.finite=TRUE)
```

Arguments

| | |
|--------------------------|-----------------------------------------------------------|
| <code>x</code> | Numeric vector of values to manipulate. |
| <code>range</code> | Numeric vector of length two giving desired output range. |
| <code>only.finite</code> | if TRUE (the default), will only modify finite values. |

Author(s)

Aaron A. King <kingaa at umich dot edu>

See Also

[censor](#)

Examples

```
trnc(c(-1,0.5,1,2,NA))
```


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