

# Package ‘aakmisc’

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**Type** Package

**Title** Miscellany for the King Lab

**Version** 0.27-1

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**Maintainer** Aaron A. King <kingaa@umich.edu>

**Description** Miscellaneous tools, collected in one place.

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

**Contact** <kingaa@umich.edu>

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

**Depends** R(>= 3.4.0)

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

**Suggests** scales, magrittr

**License** GPL-3

**LazyLoad** true

**LazyData** true

**Encoding** UTF-8

**Collate** 'db.R'

'tunnel.R'

'aaa.R'

'lazyload.R'

'matrix2latex.R'

'plotMatrix.R'

'random.R'

'scinot.R'

'trnc.R'

**RoxygenNote** 6.1.0

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db	<i>Interface with databases</i>
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### Description

Interface with project databases.

### Usage

```
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), ...)
```

```
getQuery(statement, host = getOption("aakmisc.dbhost", "localhost"),
  dbname = getOption("aakmisc.dbname", NULL),
  port = getOption("aakmisc.port", 5432),
  user = getOption("aakmisc.user", NULL), ...)
```

```
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), ...)
```

```
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), ...)

listScripts(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), ...)

```

### Arguments

name, value	Name and contents of table to create.
overwrite, append, row.names	See <a href="#">dbWriteTable</a> .
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.
...	Additional arguments will be passed to <a href="#">dbConnect</a> .
statement	SQL statement passed to <a href="#">dbGetQuery</a> .
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 <a href="#">cat</a> .

### Author(s)

Aaron A. King

### Examples

```

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

## End(Not run)

```

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`lazyload`*Functions for lazy-loading knitr caches.*

---

## Description

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

## 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>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 <a href="#">list.files</a> .
<code>...</code>	additional arguments passed to <a href="#">list.files</a>
<code>labels</code>	character vector; chunk labels to load
<code>filter</code>	optional function; passed to <a href="#">lazyLoad</a> . 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.

## Details

Use `lazyload_cache_dir` to load a whole directory of cached objects.

Use `lazyload_cache_labels` to load and explicit set of cached chunks.

## Value

Both functions return NULL, invisibly.

## Author(s)

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

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matrix2latex	<i>matrix2latex</i>
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**Description**

Format a matrix for latex.

**Usage**

```
matrix2latex(x, type = "pmatrix")
```

**Arguments**

x	matrix
type	latex matrix environment

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plotMatrix	<i>A scatterplot matrix with densities on the diagonal.</i>
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**Description**

A special scatterplot matrix.

**Usage**

```
plotMatrix(data, ...)

## 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 'data.frame'
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**

<code>data</code>	Data to plot.
<code>...</code>	optional arguments, passed to <a href="#">hist</a> .
<code>marg.exp</code>	Fraction by which to expand the plot at the margins.
<code>labels</code>	Names of variables plotted.
<code>alpha, pch, size</code>	Refer to the plotted points in the scatterplots.
<code>x</code>	<code>plotMatrix</code> object to display.
<code>newpage</code>	logical; if TRUE, <code>grid.newpage()</code> will be called before the graphics are drawn.
<code>vp</code>	viewport to use. See <a href="#">viewport</a> .

**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)
```

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random

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*Functions for generating and working with truly random integers.*


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**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.

**Details**

random.org gets seeds from [random.org](http://www.random.org).

urandom gets seeds locally from `/dev/urandom` on \*nix systems.

rngControl is a function to control RNG for the evaluation of an expression.

rngSeeds generates RNG seeds using [runif](#). It is included for situations when neither [random.org](#) nor [urandom](#) is available.

**Value**

integers suitable for use as RNG seeds

**Author(s)**

Aaron A. King

**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)

```

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scinot

*Scientific notation.*


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## 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. format="expression" results in an R expression. format="latex" results in a latex expression. format="math" is like "latex" but wraps the text in "\$".
simplify	logical. If simplify=TRUE, then $1 \times 10^n$ is simplified to $10^n$ .

## Author(s)

Aaron A. King

## 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)

```



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trnc	<i>Truncation of plots.</i>
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**Description**

Truncate to the specified window.

**Usage**

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

**Arguments**

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

**Details**

trnc is a function for truncating data to a specified window. It is suitable for use in `scale_{x,y}_{continuous,discrete}`, for example.

**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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