

# Package: caMisc (via r-universe)

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**Type** Package

**Title** Different Functions

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**Description** More about what it does (maybe more than one line)

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**Imports** rmarkdown, MASS, jpeg, png, tiff, jsonlite, XML, RCurl, utils, methods

**LinkingTo** RcppEigen

**RxygenNote** 7.2.3

**Repository** <https://calbertsen.r-universe.dev>

**RemoteUrl** <https://github.com/calbertsen/caMisc>

**RemoteRef** HEAD

**RemoteSha** 51710160f4ece1d5cc36c8fbc66a5cc500743aee

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<b>addShade</b>	<i>Add shade to a color name</i>
-----------------	----------------------------------

---

## Description

Add shade to a color name

## Usage

```
addShade(name, shade = 0)
```

## Arguments

name	name of the color
shade	shade value (between 0 and 1). For zero, the same color is returned.

## Value

a new color name

## Author(s)

Christoffer Moesgaard Albertsen

---

**addTint** *Add tint to a color name*

---

### Description

Add tint to a color name

### Usage

```
addTint(name, tint = 0)
```

### Arguments

name	name of the color
tint	tint value (between 0 and 1). For 0, the same color is returned

### Value

a new color name

### Author(s)

Christoffer Moesgaard Albertsen

---

**addTone** *Add tone to a color name*

---

### Description

Add tone to a color name

### Usage

```
addTone(name, tone = 0)
```

### Arguments

name	name of the color
tone	tone value (between 0 and 1). For 0, the same color is returned.

### Value

a new color name

### Author(s)

Christoffer Moesgaard Albertsen

---

addTrans	<i>Add alpha value to a color name</i>
----------	--

---

**Description**

Add alpha value to a color name

**Usage**

```
addTrans(name, alpha = 1)
```

**Arguments**

name	name of the color
alpha	alpha value (between 0 and 1)

**Value**

a new color name

**Author(s)**

Christoffer Moesgaard Albertsen

---

axisInside	<i>Draw figure axis inside plot</i>
------------	-------------------------------------

---

**Description**

Draw figure axis inside plot

**Usage**

```
axisInside(side)
```

**Arguments**

side	side of plot to draw on
------	-------------------------

**Value**

Nothing, but plots as side effect

**Author(s)**

Christoffer Moesgaard Albertsen

---

beanplot

*Beanplot of something*

---

## Description

Beanplot of something

## Usage

`beanplot(x, ...)`

## Arguments

x	Some object
...	Other arguments

## Value

Creates a plot

## Author(s)

Christoffer Moesgaard Albertsen

---

---

bgtext

*Add text with background to plot*

---

## Description

Add text with background to plot

## Usage

```
bgtext(  
  x,  
  y,  
  labels,  
  ...,  
  cex = 1,  
  font = NULL,  
  bg = "white",  
  bgex = 1,  
  border = NA  
)
```

**Arguments**

x	x-coordinate
y	y-coordinate
labels	label to add
...	arguments passed to text
cex	size of text
font	font of text
bg	background color
bgex	background extend factor
border	border color of background

**Author(s)**

Christoffer Moesgaard Albertsen

`buildFromGithub`

*Download and build an R package from GitHub*

**Description**

Download and build an R package from GitHub

**Usage**

```
buildFromGithub(
  repo,
  ref,
  subdir = NULL,
  buildArgs = c("--no-build-vignettes")
)
```

**Arguments**

repo	GitHub user and repository separated by /
ref	Reference to commit or branch. Default is master
subdir	Path to subdir containing the package. Should be NULL if the package is in the top directory
buildArgs	Character vector of arguments passed to R CMD build. Default is "--no-build-vignettes"

**Value**

Nothing

**Author(s)**

Christoffer Moesgaard Albertsen

---

collapse	<i>Collapse vector to string</i>
----------	----------------------------------

---

**Description**

Collapse vector to string

**Usage**

```
collapse(  
  x,  
  collap = c(rep(", ", length(x) - 2), ifelse(length(x) > 2, ", and ", " and "))  
)
```

**Arguments**

x	vector of values
collap	rule for collapsing

**Value**

a string

**Author(s)**

Christoffer Moesgaard Albertsen

---

displayNum	<i>Convert number to display text</i>
------------	---------------------------------------

---

**Description**

Convert number to display text

**Usage**

```
displayNum(  
  x,  
  capitalize = FALSE,  
  big.mark = ",",  
  decimal.mark = ".",  
  small.mark = "",  
  digits = 0  
)
```

**Arguments**

x	number
capitalize	Capitalize first letter?

**Value**

LaTeX code for the fraction

**Author(s)**

Text representation of number

---

dtucols

*DTU color palette*

---

**Description**

DTU color palette

**Usage**

`dtucols(x)`

**Arguments**

x	name of colors to return
---	--------------------------

**Value**

color codes

**Author(s)**

Christoffer Moesgaard Albertsen

---

```
dzslides_presentation  Convert to a DZSlides presentation
```

---

## Description

Format for converting from R Markdown to a DZSlides presentation.

## Usage

```
dzslides_presentation(  
  incremental = FALSE,  
  fig_width = 8,  
  fig_height = 6,  
  fig_retina = if (!fig_caption) 2,  
  fig_caption = FALSE,  
  dev = "png",  
  smart = TRUE,  
  self_contained = TRUE,  
  highlight = "default",  
  mathjax = "default",  
  template = "default",  
  css = NULL,  
  includes = NULL,  
  keep_md = FALSE,  
  lib_dir = NULL,  
  md_extensions = NULL,  
  pandoc_args = NULL,  
  ...  
)
```

## Arguments

incremental	should lists be incremental?
fig_width	figure width
fig_height	figure height
fig_retina	retina figure?
fig_caption	figure caption?
dev	figure device
smart	smart?
self_contained	self contained?
highlight	highlighting style
mathjax	mathjax
template	template path
css	extra css

<code>includes</code>	extra includes
<code>keep_md</code>	keep markdown file?
<code>lib_dir</code>	...
<code>md_extensions</code>	markdown extensions to use
<code>pandoc_args</code>	extra pandoc arguments
<code>...</code>	other arguments

**Value**

R Markdown output format to pass to `render`

**Author(s)**

Christoffer Moesgaard Albertsen

**Examples**

```
## Not run:

library(rmarkdown)

# simple invocation
render("pres.Rmd", dzslides_presentation())

# specify an option for incremental rendering
render("pres.Rmd", dzslides_presentation(incremental = TRUE))

## End(Not run)
##' @export
```

**Description**

Small wrapper for `formatC` to limit text needed

**Usage**

```
fd(x, digits = 1)
```

**Arguments**

<code>x</code>	number
<code>digits</code>	number of digits

**Value**

string

**Author(s)**

Christoffer Moesgaard Albertsen

---

formatDate

*Format date with locale*

---

**Description**

Format date with locale

**Usage**

```
formatDate(x, format = "", locale = Sys.getlocale("LC_TIME"), ...)
```

**Arguments**

x	date
format	format to use
locale	locale to use
...	passed to strftime

**Value**

formatted date string

**Author(s)**

Christoffer Moesgaard Albertsen

---

**gcovTable***Create coverage table from list of gcov objects*

---

**Description**

Create coverage table from list of gcov objects

**Usage**

```
gcovTable(x, ...)
```

**Arguments**

x	list of gcov objects
...	additional arguments passed to formatC

**Value**

A coverage table

**Author(s)**

Christoffer Moesgaard Albertsen

---

**getPixelMatrix***Get a pixel matrix from an image file*

---

**Description**

Get a pixel matrix from an image file

**Usage**

```
getPixelMatrix(file, grey = FALSE)
```

**Arguments**

file	Path to image file
grey	Should output be greyscale?

**Value**

A matrix of pixel values (0-255)

**Author(s)**

Christoffer Moesgaard Albertsen

---

<code>grad</code>	<i>Calculate gradient of a function</i>
-------------------	---

---

### Description

Calculate gradient of a function

### Usage

```
grad(
  func,
  x,
  h = abs(1e-04 * x) + 1e-04 * (abs(x) < sqrt(.Machine$double.eps/7e-07)),
  ...
)
```

### Arguments

<code>func</code>	function
<code>x</code>	parameter values
<code>h</code>	step size
<code>...</code>	passed to func

### Value

gradient vector

---

<code>greenBlindness</code>	<i>Emulate green blindness in colors</i>
-----------------------------	--

---

### Description

Based on <https://personal.sron.nl/~pault/data/colourschemes.pdf>

### Usage

```
greenBlindness(red, green, blue, alpha = 1, names = NULL, maxColorValue = 255)
```

### Arguments

<code>red</code>	Red RGB value (or color HEX code or name - alpha is ignored)
<code>green</code>	Green RGB value
<code>blue</code>	Blue RGB value
<code>alpha</code>	alpha value to use
<code>names</code>	Names for the resulting vector
<code>maxColorValue</code>	Maximum color value for red, green, blue

**Value**

New colors

**Author(s)**

Christoffer Moesgaard Albertsen

**imagePlot**

*Plot image from file*

**Description**

Plot image from file

**Usage**

```
imagePlot(
  x,
  objectFit = c("fill", "contain", "cover", "none", "scale-down"),
  halign = c("c", "l", "r"),
  valign = c("c", "t", "b"),
  maxWidthPct = 1,
  maxHeightPct = 1,
  add = FALSE,
  noMargin = TRUE,
  ...
)
```

**Arguments**

x	path to image
objectFit	How the image should fill the plot: "fill", "contain", "cover", "none", "scale-down"
halign	Horizontal alignment: c, l, r
valign	Vertical alignment: c, t, b
noMargin	Plot with oma and mar set to zero
...	other arguments

**Value**

Plots the image

**Author(s)**

Christoffer Moesgaard Albertsen

---

installDependencies    *Download, build, and install package dependencies*

---

## Description

Download, build, and install package dependencies

## Usage

```
installDependencies(  
  descriptionPath,  
  buildArgs = c("--no-build-vignettes"),  
  installArgs = c(),  
  dependencies = c("Depends", "Imports", "LinkingTo")  
)
```

## Arguments

descriptionPath	Path to a DESCRIPTION file
buildArgs	Character vector of arguments passed to R CMD build.
installArgs	Character vector of arguments passed to R CMD INSTALL.
dependencies	Character vector of dependency types to install ("Depends", "Imports", "LinkingTo", "Enhances", or "Suggests")

## Value

Nothing

## Author(s)

Christoffer Moesgaard Albertsen

---

installFromGithub    *Download, build, and install an R package from GitHub*

---

## Description

Download, build, and install an R package from GitHub

**Usage**

```
installFromGithub(
  repo,
  ref,
  subdir = NULL,
  buildArgs = c("--no-build-vignettes"),
  installArgs = c(),
  dependencies = c("Depends", "Imports", "LinkingTo"),
  https = TRUE
)
```

**Arguments**

<code>repo</code>	GitHub user and repository separated by /
<code>ref</code>	Reference to commit or branch. Default is master
<code>subdir</code>	Path to subdir containing the package. Should be NULL if the package is in the top directory
<code>buildArgs</code>	Character vector of arguments passed to R CMD build.
<code>installArgs</code>	Character vector of arguments passed to R CMD INSTALL.
<code>dependencies</code>	Character vector of dependency types to install ("Depends", "Imports", "LinkingTo", "Enhances", or "Suggests")

**Value**

Nothing

**Author(s)**

Christoffer Moesgaard Albertsen

**jacobian**

*Calculate jacobian of a function*

**Description**

Calculate jacobian of a function

**Usage**

```
jacobian(
  func,
  x,
  h = abs(1e-04 * x) + 1e-04 * (abs(x) < sqrt(.Machine$double.eps/7e-07)),
  ...
)
```

**Arguments**

func	function
x	parameter values
h	step size
...	passed to func

**Value**

jacobian matrix

---

makeMapLegend      *Add map legend for choropleth map*

---

**Description**

Add map legend for choropleth map

**Usage**

```
makeMapLegend(x, cols, txt)
```

**Arguments**

x	Variable values plotted
cols	Colors
txt	Title / description

**Author(s)**

Christoffer Moesgaard Albertsen

---

makeSquare      *Make a pixel matrix/array square*

---

**Description**

Make a pixel matrix/array square

**Usage**

```
makeSquare(p, value = 0, asp = 1)
```

**Arguments**

p	Pixel matrix/array
value	Value of added pixels

**Author(s)**

Christoffer Moesgaard Albertsen

packageSkeleton      *Create a package skeleton*

**Description**

Create a package skeleton

**Usage**

```
packageSkeleton(pkg, path = file.path("."))
```

**Arguments**

pkg	Name of the new package
path	(existing) Directory of package

**Value**

Used for side effects

**Author(s)**

Christoffer Moesgaard Albertsen

particlefilter      *Bootstrap particle filter*

**Description**

Function to run a bootstrap particle filter for user defined model.

**Usage**

```
particlefilter(N, T, G, M, F = NULL, envir = .GlobalEnv, seed = NULL)
```

## Arguments

N	Number of particles
T	Number of time steps
G	Function to simulate hidden states (of dimension p) given previous (See Details)
M	Function to calculate log-likelihood of data for a given particle
F	Do not use!
envir	Environment the functions should be evaluated in (containing data and parameters)
seed	Random seed to start the filter

## Details

The G function should be of the form  $G \leftarrow \text{function}(t, X)$  and return a numeric vector of simulate values from  $X_t | X_{t-1} = X$ . For  $t = 1$  the function will be called as  $G(1, \text{NULL})$ . The M function should be of the form  $M \leftarrow \text{function}(t, X)$  and return the density of the observation at time t given the latent state  $X_t$

## Value

A list with a  $p \times N \times T$  array X containing the simulated particles (X) and a vector of length T with the negative log-likelihood contribution at each time point.

## Author(s)

Christoffer Moesgaard Albertsen <cmoe@aqua.dtu.dk>

## Examples

```
## Univariate example
env <- new.env()
env$pars <- list(sdobs = 0.4, sdstate = 0.3)
local({xx <- cumsum(rnorm(100,0,pars$sdstate))},env)
local({dat <- xx + rnorm(100,0,pars$sdobs)},env)
G <- function(t,X){
  if(t == 1){
    return(rnorm(1,0,pars$sdstate))
  }else{
    return(rnorm(1,X,pars$sdstate))
  }
}
M <- function(t,X){
  return(dnorm(dat[t],X,pars$sdobs, TRUE))
}
pest <- particlefilter(N = 1000, T = 100,
                       G=G,M=M,
                       envir=env, seed=1)
## Not run:
plot(env$dat)
lines(apply(pest$X[,],2,mean),col="red")
```

```

lines(apply(pest$X[, , 2], mean) + 2 * apply(pest$X[, , 2], sd), col="red", lty=2)
lines(apply(pest$X[, , 2], mean) - 2 * apply(pest$X[, , 2], sd), col="red", lty=2)
lines(env$xx)

## End(Not run)

## Bivariate example
env <- new.env()
env$pars <- list(sdobs = 0.4, sdstate = 0.3)
local({xx <- cbind(cumsum(rnorm(100, 0, pars$sdstate)),
                    cumsum(rnorm(100, 0, pars$sdstate))), env})
local({dat <- xx + matrix(rnorm(2 * 100, 0, pars$sdobs), ncol=2)}, env)
G <- function(t, X){
  if(t == 1){
    return(rnorm(2, 0, pars$sdstate))
  }else{
    return(rnorm(2, X, pars$sdstate))
  }
}
M <- function(t, X){
  return(sum(dnorm(dat[t, ], X, pars$sdobs, TRUE)))
}
pest <- particlefilter(N = 1000, T = 100,
                        G=G, M=M,
                        envir=env, seed=1)

## Not run:
layout(cbind(1, c(2, 3)))
plot(env$dat)
lines(env$xx)
lines(apply(pest$X, c(3, 1), mean), col="red")
plot(env$dat[, 1])
lines(apply(pest$X[, , 2], mean), col="red")
lines(apply(pest$X[, , 2], mean) + 2 * apply(pest$X[, , 2], sd), col="red", lty=2)
lines(apply(pest$X[, , 2], mean) - 2 * apply(pest$X[, , 2], sd), col="red", lty=2)
lines(env$xx[, 1])
plot(env$dat[, 2])
lines(apply(pest$X[, , 2], mean), col="red")
lines(apply(pest$X[, , 2], mean) + 2 * apply(pest$X[, , 2], sd), col="red", lty=2)
lines(apply(pest$X[, , 2], mean) - 2 * apply(pest$X[, , 2], sd), col="red", lty=2)
lines(env$xx[, 2])

## End(Not run)

```

**Description**

Read content of .gcov file

**Usage**

```
read_gcov(file)
```

**Arguments**

file                path to .gcov file

**Value**

a gcov object

**Author(s)**

Christoffer Moesgaard Albertsen

---

read\_massif                *Read valgrind Massif output*

---

**Description**

Read valgrind Massif output

**Usage**

```
read_massif(file, keep_details = FALSE)
```

**Arguments**

file                file to read  
keep\_details      keep details?

**Value**

massif S3 object

**Author(s)**

Christoffer Moesgaard Albertsen

<code>solarPosition</code>	<i>Calculate solar angles</i>
----------------------------	-------------------------------

### Description

Calculate solar angles

### Usage

```
solarPosition(date, lat, lon)
```

### Arguments

<code>date</code>	date (UTC) in the format "YYYY-mm-dd HH:MM:SS"
<code>lat</code>	latitude of observer
<code>lon</code>	longitude of observer

### Value

list of values

### Author(s)

Christoffer Moesgaard Albertsen Modified from <https://doi.org/10.1016/j.renene.2021.03.047>

<code>tofrac</code>	<i>Print number as (LaTeX) fraction</i>
---------------------	---

### Description

Print number as (LaTeX) fraction

### Usage

```
tofrac(x, dollar = TRUE)
```

### Arguments

<code>x</code>	number
<code>dollar</code>	

### Value

LaTeX code for the fraction

### Author(s)

Christoffer Moesgaard Albertsen

---

<code>toGreyscale</code>	<i>Turn pixel into grey scale</i>
--------------------------	-----------------------------------

---

**Description**

Turn pixel into grey scale

**Usage**

`toGreyscale(p)`

**Arguments**

<code>p</code>	Pixel matrix/array
----------------	--------------------

**Value**

A grey scale pixel matrix/array

**Author(s)**

Christoffer Moesgaard Albertsen

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