

# Package ‘R.utils’

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**Suggests** digest (>= 0.6.10)

**Title** Various Programming Utilities

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**Description** Utility functions useful when programming and developing R packages.

**License** LGPL (>= 2.1)

**LazyLoad** TRUE

**URL** <https://github.com/HenrikBengtsson/R.utils>

**BugReports** <https://github.com/HenrikBengtsson/R.utils/issues>

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## R topics documented:

R.utils-package	4
addFinalizerToLast	6
Arguments	7
as.character.binmode	8
Assert	9
attachLocally.list	10
callHooks	11
callHooks.function	13
capitalize	13
captureOutput	14
cmdArgs	16

colClasses . . . . .	17
compressFile . . . . .	18
compressPDF . . . . .	20
copyDirectory . . . . .	21
countLines . . . . .	22
createFileAtomically . . . . .	23
createLink . . . . .	25
createWindowsShortcut . . . . .	26
dataFrame . . . . .	28
detachPackage . . . . .	29
dimNA< - . . . . .	29
displayCode . . . . .	30
doCall . . . . .	31
downloadFile.character . . . . .	32
egsub . . . . .	34
env . . . . .	35
extract.array . . . . .	36
fileAccess . . . . .	38
filePath . . . . .	40
FileProgressBar . . . . .	42
finalizeSession . . . . .	43
findSourceTraceback . . . . .	44
gcat . . . . .	45
gcDLLs . . . . .	46
getAbsolutePath . . . . .	47
getParent . . . . .	48
getRelativePath . . . . .	49
GString . . . . .	50
gstring . . . . .	53
hasUrlProtocol . . . . .	54
hpaste . . . . .	55
inAnyInterval.numeric . . . . .	57
insert . . . . .	58
installPackages . . . . .	59
intervalsToSeq.matrix . . . . .	61
intToBin . . . . .	61
isAbsolutePath . . . . .	62
isDirectory . . . . .	63
isEof.connection . . . . .	63
isFile . . . . .	64
isOpen.character . . . . .	65
isPackageInstalled . . . . .	66
isPackageLoaded . . . . .	66
isReplicated . . . . .	67
isSingle . . . . .	69
isUrl . . . . .	69
isZero . . . . .	70
Java . . . . .	71

lastModified . . . . .	73
LComments . . . . .	74
listDirectory . . . . .	75
loadObject . . . . .	76
mapToIntervals.numeric . . . . .	77
mergeIntervals.numeric . . . . .	78
mkdirs . . . . .	79
mout . . . . .	80
mpager . . . . .	81
nullfile . . . . .	81
NullVerbose . . . . .	82
onGarbageCollect . . . . .	83
onSessionExit . . . . .	84
Options . . . . .	85
patchCode . . . . .	87
popBackupFile . . . . .	89
popTemporaryFile . . . . .	90
printf . . . . .	91
ProgressBar . . . . .	92
pushBackupFile . . . . .	93
pushTemporaryFile . . . . .	95
queryRCmdCheck . . . . .	97
readBinFragments . . . . .	98
readRdHelp . . . . .	100
readTable . . . . .	101
readTableIndex . . . . .	103
readWindowsShortcut . . . . .	104
removeDirectory . . . . .	105
resample . . . . .	106
saveObject . . . . .	107
seqToHumanReadable . . . . .	108
seqToIntervals . . . . .	109
setOption . . . . .	110
Settings . . . . .	110
shell.exec2 . . . . .	113
SmartComments . . . . .	114
sourceDirectory . . . . .	116
sourceTo . . . . .	117
splitByPattern . . . . .	119
stext . . . . .	120
subplots . . . . .	121
System . . . . .	122
systemR . . . . .	123
TextStatusBar . . . . .	124
TimeoutException . . . . .	126
touchFile . . . . .	127
toUrl . . . . .	129
unwrap.array . . . . .	129

useRepos . . . . .	130
VComments . . . . .	131
Verbose . . . . .	133
withCapture . . . . .	137
withLocale . . . . .	139
withOptions . . . . .	140
withRepos . . . . .	142
withSeed . . . . .	143
withSink . . . . .	144
withTimeout . . . . .	145
wrap.array . . . . .	148
writeBinFragments . . . . .	151
writeDataFrame.data.frame . . . . .	152

<b>Index</b>	<b>153</b>
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R.utils-package	<i>Package R.utils</i>
-----------------	------------------------

---

## Description

Utility functions useful when programming and developing R packages.

*Warning:* The Application Programming Interface (API) of the classes and methods in this package may change. Classes and methods are considered either to be stable, or to be in beta or alpha (pre-beta) stage. See list below for details.

The main reason for publishing this package on CRAN although it lacks a stable API, is that its methods and classes are used internally by other packages on CRAN that the author has published.

For package history, see `showHistory(R.utils)`.

## Requirements

This package requires the **R.oo** package [1].

## Installation and updates

To install this package do:

```
install.packages("R.utils")
```

## To get started

- [Arguments](#)[alpha] Methods for common argument processing.
- [Assert](#)[alpha] Methods for assertion of values and states.
- [GString](#)[alpha] A character string class with methods for simple substitution.
- [Java](#)[beta] Reads and writes Java streams.
- [Options](#)[alpha] Tree-structured options queried in a file-system like manner.

- [Settings](#)[alpha] An Options class for reading and writing package settings.
- [ProgressBar](#)[beta] Text-based progress bar.
- [FileProgressBar](#)[beta] A ProgressBar that reports progress as file size.
- [System](#)[alpha] Methods for access to system.
- [Verbose](#)[alpha] A class for verbose and log output. Utilized by the VComments and LComments classes.
- [SmartComments](#), [VComments](#), [LComments](#)[alpha] Methods for preprocessing source code comments of certain formats into R code.

In addition to the above, there is a large set of function for file handling such as support for reading/following Windows Shortcut links, but also other standalone utility functions. See package index for a list of these. These should also be considered to be in alpha or beta stage.

### How to cite this package

Whenever using this package, please cite [1] as

Bengtsson, H. The R.oo package - Object-Oriented Programming with References Using Standard R Code, Proceedings of the 3rd International Workshop on Distributed Statistical Computing (DSC 2003), ISSN 1609-395X, Hornik, K.; Leisch, F. & Zeileis, A. (ed.), 2003

### Wishlist

Here is a list of features that would be useful, but which I have too little time to add myself. Contributions are appreciated.

- Write a TclTkProgressBar class.
- Improve/stabilize the GString class.
- Mature the SmartComments classes. Also add AComments and PComments for assertion and progress/status comments.

If you consider implement some of the above, make sure it is not already implemented by downloading the latest "devel" version!

### License

The releases of this package is licensed under LGPL version 2.1 or newer.

The development code of the packages is under a private licence (where applicable) and patches sent to the author fall under the latter license, but will be, if incorporated, released under the "release" license above.

### References

- 1 H. Bengtsson, *The R.oo package - Object-Oriented Programming with References Using Standard R Code*, In Kurt Hornik, Friedrich Leisch and Achim Zeileis, editors, Proceedings of the 3rd International Workshop on Distributed Statistical Computing (DSC 2003), March 20-22, Vienna, Austria. <http://www.ci.tuwien.ac.at/Conferences/DSC-2003/Proceedings/>

**Author(s)**

Henrik Bengtsson

---

addFinalizerToLast      *Modifies .Last() to call 'finalizeSession()*

---

**Description**

Modifies `.Last()` to call `'finalizeSession()` *before* calling the default `.Last()` function.

Note that `.Last()` is *not* guaranteed to be called when the R session finished. For instance, the user may quit R by calling `quit(runLast=FALSE)` or run R in batch mode.

Note that this function is called when the R.utils package is loaded.

**Usage**

```
## Default S3 method:  
addFinalizerToLast(...)
```

**Arguments**

...                      Not used.

**Value**

Returns (invisibly) `TRUE` if `.Last()` was modified, otherwise `FALSE`.

**Author(s)**

Henrik Bengtsson

**See Also**

[onSessionExit\(\)](#).

---

Arguments

*Static class to validate and process arguments*


---

**Description**

Package: R.utils

**Class Arguments**[Object](#)

~~|

~~+--Arguments

**Directly known subclasses:**public static class **Arguments**extends [Object](#)**Fields and Methods****Methods:**

<a href="#">getCharacter</a>	-
<a href="#">getCharacters</a>	Coerces to a character vector and validates.
<a href="#">getDirectory</a>	-
<a href="#">getDouble</a>	-
<a href="#">getDoubles</a>	Coerces to a double vector and validates.
<a href="#">getEnvironment</a>	Gets an existing environment.
<a href="#">getFilename</a>	Gets and validates a filename.
<a href="#">getIndex</a>	-
<a href="#">getIndices</a>	Coerces to a integer vector and validates.
<a href="#">getInstanceOf</a>	Gets an instance of the object that is of a particular class.
<a href="#">getInteger</a>	-
<a href="#">getIntegers</a>	Coerces to a integer vector and validates.
<a href="#">getLogical</a>	-
<a href="#">getLogicals</a>	Coerces to a logical vector and validates.
<a href="#">getNumeric</a>	-
<a href="#">getNumerics</a>	Coerces to a numeric vector and validates.
<a href="#">getReadablePath</a>	-
<a href="#">getReadablePathname</a>	Gets a readable pathname.
<a href="#">getReadablePathnames</a>	Gets a readable pathname.
<a href="#">getRegularExpression</a>	Gets a valid regular expression pattern.
<a href="#">getVector</a>	Validates a vector.
<a href="#">getVerbose</a>	Coerces to Verbose object.

```
getWritablePath      -
getWritablePathname Gets a writable pathname.
```

**Methods inherited from Object:**

\$, \$<-, [], [[<-, as.character, attach, attachLocally, clearCache, clearLookupCache, clone, detach, equals, extend, finalize, getEnvironment, getFieldModifier, getFieldModifiers, getFields, getInstantiationTime, getStaticInstance, hasField, hashCode, ll, load, names, objectSize, print, save

**Author(s)**

Henrik Bengtsson

---

as.character.binmode *Converts a binary/octal/hexadecimal number into a string*

---

**Description**

Converts a binary/octal/hexadecimal number into a string.

**Usage**

```
## S3 method for class 'binmode'
as.character(x, ...)
```

**Arguments**

```
x          Object to be converted.
...        Not used.
```

**Value**

Returns a [character](#).

**Author(s)**

Henrik Bengtsson

**See Also**

as.character.octmode(), cf. [octmode.intToBin\(\)](#) (incl. [intToOct\(\)](#) and [intToHex\(\)](#)).



---

 Assert

*The Assert class*


---

**Description**

Package: R.utils

**Class Assert**[Object](#)

~~|

~~+--Assert

**Directly known subclasses:**public static class **Assert**extends [Object](#)**Usage**

Assert(...)

**Arguments**

... Not used.

**Fields and Methods****Methods:**

<a href="#">check</a>	Static method asserting that a generic condition is true.
<a href="#">inherits</a>	-
<a href="#">inheritsFrom</a>	Static method asserting that an object inherits from of a certain class.
<a href="#">isMatrix</a>	Static method asserting that an object is a matrix.
<a href="#">isScalar</a>	Static method asserting that an object is a single value.
<a href="#">isVector</a>	Static method asserting that an object is a vector.

**Methods inherited from Object:**

[\\$](#), [\\$<-](#), [\[\[](#), [\[\[<-](#), [as.character](#), [attach](#), [attachLocally](#), [clearCache](#), [clearLookupCache](#), [clone](#), [detach](#), [equals](#), [extend](#), [finalize](#), [getEnvironment](#), [getFieldModifier](#), [getFieldModifiers](#), [getFields](#), [getInstantiationTime](#), [getStaticInstance](#), [hasField](#), [hashCode](#), [ll](#), [load](#), [names](#), [objectSize](#), [print](#), [save](#)

**Author(s)**

Henrik Bengtsson

---

attachLocally.list      *Assigns an objects elements locally*

---

**Description**

Assigns an objects elements locally.

**Usage**

```
## S3 method for class 'list'  
attachLocally(object, fields=NULL, excludeFields=NULL, overwrite=TRUE,  
  envir=parent.frame(), ...)
```

**Arguments**

object	An object with named elements such as an <a href="#">environment</a> , a <a href="#">list</a> , or a <a href="#">data.frame</a> .
fields	A <a href="#">character vector</a> specifying elements to be copied. If <code>NULL</code> , all elements are considered.
excludeFields	A <a href="#">character vector</a> specifying elements not to be copied. This has higher priority than <code>fields</code> .
overwrite	If <code>FALSE</code> , fields that already exists will not be copied.
envir	The <a href="#">environment</a> where elements are copied to.
...	Not used.

**Value**

Returns (invisibly) a [character vector](#) of the fields copied.

**Author(s)**

Henrik Bengtsson

**See Also**

[attachLocally\(\)](#) of class `Object`. [attach\(\)](#).

**Examples**

```
foo <- function(object) {
  cat("Local objects in foo():\n")
  print(ls())

  attachLocally(object)

  cat("\nLocal objects in foo():\n")
  print(ls())

  for (name in ls()) {
    cat("\nObject '", name, "':\n", sep="")
    print(get(name, inherits=FALSE))
  }
}

a <- "A string"
l <- list(a=1:10, msg="Hello world", df=data.frame(a=NA, b=2))
foo(l)
print(a)
```

---

callHooks

*Call hook functions by hook name*


---

**Description**

Call hook functions by hook name.

**Usage**

```
## Default S3 method:
callHooks(hookName, ..., removeCalledHooks=FALSE)
```

**Arguments**

hookName            A [character](#) string of the hook name.  
...                    Argument passed to each hook function.  
removeCalledHooks    If [TRUE](#), called hook functions are removed, otherwise not.

**Value**

Returns (invisibly) whatever [callHooks.list\(\)](#) returns.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally, after retrieving hook functions, `callHooks.list()` is called.

**Examples**

```
# -----
# Example 1
# -----
# First, clean up if called more than once
setHook("myFunction.onEnter", NULL, action="replace")
setHook("myFunction.onExit", NULL, action="replace")

runConference <- function(...) {
  callHooks("myFunction.onEnter")
  cat("Speaker A: Hello there...\n")
  callHooks("myFunction.onExit")
}

setHook("myFunction.onEnter", function(...) {
  cat("Chair: Welcome to our conference.\n")
})

setHook("myFunction.onEnter", function(...) {
  cat("Chair: Please welcome Speaker A!\n")
})

setHook("myFunction.onExit", function(...) {
  cat("Chair: Please thanks Speaker A!\n")
})

runConference()

# -----
# Example 2
# -----
setHook("randomNumber", NULL, action="replace")
setHook("randomNumber", rnorm)      # By function
setHook("randomNumber", "rexp")    # By name
setHook("randomNumber", "runiff")  # Non-existing name
setHook("randomNumber", .GlobalEnv) # Not a function

res <- callHooks("randomNumber", n=1)
str(res)
cat("Number of hooks: ", length(res), "\n");
isErroneous <- unlist(lapply(res, FUN=function(x) !is.null(x$exception)));
cat("Erroneous hooks: ", sum(isErroneous), "\n");
```

---

callHooks.function	<i>Call hook functions</i>
--------------------	----------------------------

---

**Description**

Call hook functions.

**Usage**

```
## S3 method for class 'function'  
callHooks(hooks, ...)
```

**Arguments**

hooks	A <a href="#">function</a> or a <a href="#">list</a> of hook <a href="#">functions</a> or names of such.
...	Argument passed to each hook function.

**Value**

Returns (invisibly) a [list](#) that is named with hook names, if possible. Each element in the list is in turn a [list](#) with three element: `fcn` is the hook function called, `result` is its return value, and `exception` is the exception caught or [NULL](#).

**Author(s)**

Henrik Bengtsson

**See Also**

See [callHooks\(\)](#) to call hook function by name.

---

capitalize	<i>Capitalizes/decapitalizes each character string in a vector</i>
------------	--

---

**Description**

Capitalizes/decapitalized (making the first letter upper/lower case) of each character string in a vector.

**Usage**

```
## Default S3 method:  
capitalize(str, ...)  
## Default S3 method:  
decapitalize(str, ...)
```

**Arguments**

`str`            A [vector](#) of [character](#) strings to be capitalized.  
`...`            Not used.

**Value**

Returns a [vector](#) of [character](#) strings of the same length as the input vector.

**Author(s)**

Henrik Bengtsson

**See Also**

[toCamelCase](#).

**Examples**

```
words <- strsplit("Hello wOrld", " ")[[1]];
cat(paste(toupper(words), collapse=" "), "\n")        # "HELLO WORLD"
cat(paste(tolower(words), collapse=" "), "\n")        # "hello world"
cat(paste(capitalize(words), collapse=" "), "\n")     # "Hello WOrld"
cat(paste(decapitalize(words), collapse=" "), "\n")   # "hello wOrld"

# Sanity checks
stopifnot(paste(toupper(words), collapse=" ") == "HELLO WORLD")
stopifnot(paste(tolower(words), collapse=" ") == "hello world")
stopifnot(paste(capitalize(words), collapse=" ") == "Hello WOrld")
stopifnot(paste(decapitalize(words), collapse=" ") == "hello wOrld")
```

---

captureOutput

*Evaluate an R expression and captures the output*

---

**Description**

Evaluate an R expression and captures the output.

**Usage**

```
captureOutput(expr, file=NULL, append=FALSE, collapse=NULL, envir=parent.frame())
```

## Arguments

expr	The R expression to be evaluated.
file	A file name or a <a href="#">connection</a> to where the output is directed. Alternatively, if <a href="#">NULL</a> the output is captured to and returned as a <a href="#">character vector</a> .
append	If <a href="#">TRUE</a> , the output is appended to the file or the (unopened) connection, otherwise it overwrites.
collapse	A <a href="#">character</a> string used for collapsing the captured rows. If <a href="#">NULL</a> , the rows are not collapsed.
envir	The <a href="#">environment</a> in which the expression is evaluated.

## Details

This method imitates [capture.output](#) with the major difference that it captures strings via a [raw](#) connection rather than via internal strings. The latter becomes exponentially slow for large outputs [1,2].

## Value

Returns captured output as a [character vector](#).

## Author(s)

Henrik Bengtsson

## References

- [1] R-devel thread 'capture.output(): Using a rawConnection() [linear] instead of textConnection() [exponential]?', 2014-02-03. <https://stat.ethz.ch/pipermail/r-devel/2014-February/068349.html> [2] JottR blog post 'PERFORMANCE: captureOutput() is much faster than capture.output()', 2015-05-26. <http://www.jottr.org/2014/05/captureOutput.html>

## See Also

Internally, [eval\(\)](#) is used to evaluate the expression. and [capture.output](#) to capture the output.

## Examples

```
# captureOutput() is much faster than capture.output()
# for large outputs when capturing to a string.
for (n in c(10e3, 20e3, 30e3, 40e3)) {
  printf("n=%d\n", n)

  x <- rnorm(n)

  t0 <- system.time({
    bfr0 <- capture.output(print(x))
  })
  print(t0)
```

```

t1 <- system.time({
  bfr <- captureOutput(print(x))
})
print(t1)
print(t1/t0)

bfr2n <- captureOutput(print(x), collapse="\n")
bfr2r <- captureOutput(print(x), collapse="\r")

stopifnot(identical(bfr, bfr0))
} # for (n ...)

```

---

cmdArgs

*Simple access to parsed command-line arguments*


---

### Description

Simple access to parsed command-line arguments.

### Usage

```

cmdArgs(args=NULL, names=NULL, unique=TRUE, ..., .args=NULL)
cmdArg(...)

```

### Arguments

args	A named <a href="#">list</a> of arguments.
names	A <a href="#">character vector</a> specifying the arguments to be returned. If <code>NULL</code> , all arguments are returned.
unique	If <code>TRUE</code> , only unique arguments are returned.
...	For <code>cmdArgs()</code> , additional arguments passed to <code>commandArgs()</code> , e.g. <code>defaults</code> and <code>always</code> . For <code>cmdArg()</code> , named arguments <code>name</code> and <code>default</code> , where <code>name</code> must be a <a href="#">character</a> string and <code>default</code> is an optional default value (if not given, it's <code>NULL</code> ). Alternatively, <code>name</code> and <code>default</code> can be given as a named argument (e.g. <code>n=42</code> ).
.args	(advanced/internal) A named <a href="#">list</a> of parsed command-line arguments.

### Value

`cmdArgs()` returns a named [list](#) with command-line arguments. `cmdArg()` return the value of the requested command-line argument.

### Coercing to non-character data types

The value of each command-line argument is returned as a [character](#) string, unless an argument share name with `ditto` in the (optional) arguments `always` and `default` in case the retrieved value is coerced to that of the latter. Finally, remaining character string command-line arguments are coerced to [numerics](#) (via `as.numeric()`), if possible, that is unless the coerced value becomes `NA`.



**Author(s)**

Henrik Bengtsson

**See Also**

Internally, `commandArgs()` is used.

**Examples**

```
args <- cmdArgs()
cat("User command-line arguments used when invoking R:\n")
str(args)

# Retrieve command line argument 'n', e.g. '-n 13' or '--n=13'
n <- cmdArg("n", 42L)
printf("Argument n=%d\n", n)

# Short version doing the same
n <- cmdArg(n=42L)
printf("Argument n=%d\n", n)
```

---

colClasses

*Creates a vector of column classes used for tabular reading*

---

**Description**

Creates a vector of column classes used for tabular reading based on a compact format string.

**Usage**

```
## Default S3 method:
colClasses(fmt, ...)
```

**Arguments**

`fmt` A [character](#) string specifying the column-class format. This string is first translated by `sprintf()`.

`...` Optional arguments for the `sprintf()` translation.

**Value**

Returns a [vector](#) of [character](#) strings.

**Author(s)**

Henrik Bengtsson

**See Also**

[read.table.](#)

**Examples**

```
# All predefined types
print(colClasses("-?cdfilnrzDP"))
## [1] "NULL"      "NA"         "character"  "double"
## [5] "factor"     "integer"    "logical"    "numeric"
## [9] "raw"        "complex"    "Date"       "POSIXct"

# A string in column 1, integers in column 4 and 5, rest skipped
print(colClasses("c--ii----"))
## [1] "character" "NULL"      "NULL"       "integer"
## [5] "integer"   "NULL"      "NULL"       "NULL"
## [9] "NULL"

# Repeats and custom column classes
c1 <- colClasses("3c{MyClass}3{foo}")
print(c1)
## [1] "character" "character" "character"  "MyClass"
## [5] "foo"        "foo"        "foo"

# Passing repeats and class names using sprintf() syntax
c2 <- colClasses("%dc{%s}%d{foo}", 3, "MyClass", 3)
stopifnot(identical(c1, c2))

# Repeats of a vector of column classes
c3 <- colClasses("3{MyClass,c}")
print(c3)
## [1] "MyClass"   "character" "MyClass"   "character"
## [4] "MyClass"   "character"

# Large number repeats
c4 <- colClasses("321{MyClass,c,i,d}")
c5 <- rep(c("MyClass", "character", "integer", "double"), times=321)
stopifnot(identical(c4, c5))
```

---

compressFile

*Compressing and decompressing files*

---

**Description**

Compressing and decompressing files such as gzip:ed and bzip2:ed files.

*NOTE: The default (remove=TRUE) behavior is that the input file is removed after that the output file is fully created and closed.*

**Usage**

```

## Default S3 method:
compressFile(filename, destname=sprintf("%s.%s", filename, ext), ext, FUN,
  temporary=FALSE, skip=FALSE, overwrite=FALSE, remove=TRUE, BFR.SIZE=1e+07, ...)
## Default S3 method:
decompressFile(filename, destname=gsub(sprintf("[.]%s$", ext), ""), filename,
  ignore.case = TRUE), ext, FUN, temporary=FALSE, skip=FALSE, overwrite=FALSE,
  remove=TRUE, BFR.SIZE=1e+07, ...)
## Default S3 method:
isCompressedFile(filename, method=c("extension", "content"), ext, fileClass, ...)
## Default S3 method:
bzip2(filename, ..., ext="bz2", FUN=bzfile)
## Default S3 method:
bunzip2(filename, ..., ext="bz2", FUN=bzfile)
## Default S3 method:
gzip(filename, ..., ext="gz", FUN=gzfile)
## Default S3 method:
gunzip(filename, ..., ext="gz", FUN=gzfile)

```

**Arguments**

filename	Pathname of input file.
destname	Pathname of output file.
temporary	If <b>TRUE</b> , the output file is created in a temporary directory.
skip	If <b>TRUE</b> and the output file already exists, the output file is returned as is.
overwrite	If <b>TRUE</b> and the output file already exists, the file is silently overwritten, otherwise an exception is thrown (unless skip is <b>TRUE</b> ).
remove	If <b>TRUE</b> , the input file is removed afterward, otherwise not.
BFR.SIZE	The number of bytes read in each chunk.
...	Passed to the underlying function or alternatively not used.
method	A <b>character</b> string specifying how to infer whether a file is compressed or not.
ext, fileClass, FUN	(internal) Filename extension, file class, and a connection <b>function</b> used to read from/write to file.

**Details**

Internally `bzfile()` and `gzfile()` (see [connections](#)) are used to read (write) files. If the process is interrupted before completed, the partially written output file is automatically removed.

**Value**

Returns the pathname of the output file. The number of bytes processed is returned as an attribute.

`isCompressedFile()`, `isGzipped()` and `isBzipped()` return a **logical**. Note that with `method = "extension"` (default), only the filename extension is used to infer whether the file is compressed or not. Specifically, it does not matter whether the file actually exists or not.

**Author(s)**

Henrik Bengtsson

**Examples**

```
## bzip2
cat(file="foo.txt", "Hello world!")
print(isBzipped("foo.txt"))
print(isBzipped("foo.txt.bz2"))

bzip2("foo.txt")
print(file.info("foo.txt.bz2"))
print(isBzipped("foo.txt"))
print(isBzipped("foo.txt.bz2"))

bunzip2("foo.txt.bz2")
print(file.info("foo.txt"))

## gzip
cat(file="foo.txt", "Hello world!")
print(isGzipped("foo.txt"))
print(isGzipped("foo.txt.gz"))

gzip("foo.txt")
print(file.info("foo.txt.gz"))
print(isGzipped("foo.txt"))
print(isGzipped("foo.txt.gz"))

gunzip("foo.txt.gz")
print(file.info("foo.txt"))

## Cleanup
file.remove("foo.txt")
```

---

compressPDF

*Compresses a PDF (into a new PDF)*

---

**Description**

Compresses a PDF (into a new PDF).

**Usage**

```
## Default S3 method:
compressPDF(filename, path=NULL, outFilename=basename(pathname),
  outPath="compressedPDFs", skip=FALSE, overwrite=FALSE, compression="gs(ebook)+qpdf",
  ...)
```

**Arguments**

filename, path The filename and (optional) path of the PDF to be compressed.  
 outFilename, outPath The generated PDF.  
 skip If **TRUE** and an existing output file, then it is returned.  
 overwrite If **FALSE**, an error is thrown if the output file already exists, otherwise not.  
 compression A **character vector** of compression methods to apply. This overrides any low-level arguments passed via ... that **compactPDF**.  
 ... Additional arguments passed to **compactPDF**, e.g. `gs_quality`.

**Value**

Returns the pathname of the generated PDF.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally **compactPDF** is utilized.

**Examples**

```
## Not run:
  pathnameZ <- compressPDF("report.pdf")

## End(Not run)
```

---

copyDirectory

*Copies a directory*

---

**Description**

Copies a directory.

**Usage**

```
## Default S3 method:
copyDirectory(from, to=".", ..., private=TRUE, recursive=TRUE)
```

**Arguments**

from	The pathname of the source directory to be copied.
to	The pathname of the destination directory.
...	Additional arguments passed to <code>file.copy()</code> , e.g. <code>overwrite</code> .
private	If <code>TRUE</code> , files (and directories) starting with a period is also copied, otherwise not.
recursive	If <code>TRUE</code> , subdirectories are copied too, otherwise not.

**Details**

Note that this method does *not* use `copyFile()` to copy the files, but `file.copy()`.

**Value**

Returns (invisibly) a `character vector` of pathnames copied.

**Author(s)**

Henrik Bengtsson

---

countLines	<i>Counts the number of lines in a text file</i>
------------	--

---

**Description**

Counts the number of lines in a text file by counting the number of occurrences of platform-independent newlines (CR, LF, and CR+LF [1]), including a last line with neither. An empty file has zero lines.

**Usage**

```
## Default S3 method:
countLines(file, chunkSize=5e+07, ...)
```

**Arguments**

file	A <code>connection</code> or a pathname.
chunkSize	The number of bytes read in each chunk.
...	Not used.

**Details**

Both compressed and non-compressed files are supported.

**Value**

Returns an non-negative [integer](#).

**Author(s)**

Henrik Bengtsson

**References**

[1] Page *Newline*, Wikipedia, July 2008. <http://en.wikipedia.org/wiki/Newline>

**Examples**

```
pathname <- system.file("NEWS", package="R.utils");
n <- countLines(pathname);
n2 <- length(readLines(pathname));
stopifnot(n == n2);
```

---

createFileAtomically *Creates a file atomically*

---

**Description**

Creates a file atomically by first creating and writing to a temporary file which is then renamed.

**Usage**

```
## Default S3 method:
createFileAtomically(filename, path=NULL, FUN, ..., skip=FALSE, overwrite=FALSE,
  backup=TRUE, verbose=FALSE)
```

**Arguments**

filename	The filename of the file to create.
path	The path to the file.
FUN	A <a href="#">function</a> that creates and writes to the pathname that is passed as the first argument. This pathname is guaranteed to be a non-existing temporary pathname.
...	Additional arguments passed to <a href="#">pushTemporaryFile()</a> and <a href="#">popTemporaryFile()</a> .
skip	If <a href="#">TRUE</a> and a file with the same pathname already exists, nothing is done/written.
overwrite	If <a href="#">TRUE</a> and a file with the same pathname already exists, the existing file is overwritten. This is also done atomically such that if the new file was not successfully created, the already original file is restored. If restoration also failed, the original file remains as the pathname with suffix ".bak" appended.
backup	If <a href="#">TRUE</a> and a file with the same pathname already exists, then it is backed up while creating the new file. If the new file was not successfully created, the original file is restored from the backup copy.
verbose	A <a href="#">logical</a> or <a href="#">Verbose</a> .

**Value**

Returns (invisibly) the pathname.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally, [pushTemporaryFile\(\)](#) and [popTemporaryFile\(\)](#) are used for working toward a temporary file, and [pushBackupFile\(\)](#) and [popBackupFile\(\)](#) are used for backing up and restoring already existing file.

**Examples**

```
# -----
# Create a file atomically
# -----
n <- 10
createFileAtomically("foobar.txt", FUN=function(pathname) {
  cat(file=pathname, "This file was created atomically.\n")
  cat(file=pathname, "Timestamp: ", as.character(Sys.time()), "\n", sep="")
  for (kk in 1:n) {
    cat(file=pathname, kk, "\n", append=TRUE)
    # Emulate a slow process
    if (interactive()) Sys.sleep(0.1)
  }
  cat(file=pathname, "END OF FILE\n", append=TRUE)
}, overwrite=TRUE)

bfr <- readLines("foobar.txt")
cat(bfr, sep="\n")

# -----
# Overwrite the file atomically (emulate write failure)
# -----
tryCatch({
  createFileAtomically("foobar.txt", FUN=function(pathname) {
    cat(file=pathname, "Trying to create a new file.\n")
    cat(file=pathname, "Writing a bit, but then an error...\n", append=TRUE)
    # Emulate write error
    stop("An error occurred while writing to the new file.")
    cat(file=pathname, "END OF FILE\n", append=TRUE)
  }, overwrite=TRUE)
}, error = function(ex) {
  print(ex$message)
})

# The original file was never overwritten
bfr2 <- readLines("foobar.txt")
```



```

cat(bfr2, sep="\n")
stopifnot(identical(bfr2, bfr))

# The partially temporary file remains
stopifnot(isFile("foobar.txt.tmp"))
bfr3 <- readLines("foobar.txt.tmp")
cat(bfr3, sep="\n")

file.remove("foobar.txt.tmp")

```

---

createLink	<i>Creates a link to a file or a directory</i>
------------	--

---

### Description

Creates a link to a file or a directory. This method tries to create a link to a file/directory on the file system, e.g. a symbolic link and Windows Shortcut links. It depends on operating and file system (and argument settings), which type of link is finally created, but all this is hidden internally so that links can be created the same way regardless of system.

### Usage

```

## Default S3 method:
createLink(link=".", target, skip=!overwrite, overwrite=FALSE,
  methods=getOption("createLink/args/methods", c("unix-symlink", "windows-ntfs-symlink",
    "windows-shortcut")), ...)

```

### Arguments

link	The path or pathname of the link to be created. If "." (or <code>NULL</code> ), it is inferred from the target argument, if possible.
target	The target file or directory to which the shortcut should point to.
skip	If <code>TRUE</code> and a file with the same name as argument link already exists, then the nothing is done.
overwrite	If <code>TRUE</code> , an existing link file is overwritten, otherwise not.
methods	A <code>character vector</code> specifying what methods (and in what order) should be tried for creating links.
...	Not used.

### Value

Returns (invisibly) the path or pathname to the link. If no link was created, `NULL` is returned.

**Required privileges on Windows**

In order for method="unix-symlink" (utilizing `file.symlink()`), method="windows-ntfs-symlink" (utilizing executable `mklink`), and/or method="windows-shortcut" (utilizing `createWindowsShortcut()`) to succeed on Windows, the client/R session must run with sufficient privileges (it has been reported that Administrative rights are necessary).

**Author(s)**

Henrik Bengtsson

**References**

Ben Garrett, *Windows File Junctions, Symbolic Links and Hard Links*, September 2009 [<http://goo.gl/R21AC>]

**See Also**

`createWindowsShortcut()` and `file.symlink()`

---

`createWindowsShortcut` *Creates a Microsoft Windows Shortcut (.lnk file)*

---

**Description**

Creates a Microsoft Windows Shortcut (.lnk file).

**Usage**

```
## Default S3 method:  
createWindowsShortcut(pathname, target, overwrite=FALSE, ...)
```

**Arguments**

pathname	The pathname (with file extension *.lnk) of the link file to be created.
target	The target file or directory to which the shortcut should point to.
overwrite	If <code>TRUE</code> , an existing link file is overwritten, otherwise not.
...	Not used.

**Value**

Returns (invisibly) the pathname.

**Required privileges on Windows**

In order for this method, which utilizes Windows Script Host a VBScript, to succeed on Windows, the client/R session must run with sufficient privileges (it has been reported that Administrative rights are necessary).

**Author(s)**

Henrik Bengtsson

**References**

[1] Create a windows shortcut (.LNK file), SS64.com, <http://ss64.com/nt/shortcut.html>

**See Also**

[readWindowsShortcut\(\)](#)

**Examples**

```
# Create Windows Shortcut links to a directory and a file
targets <- list(
  system.file(package="R.utils"),
  system.file("DESCRIPTION", package="R.utils")
)

for (kk in seq_along(targets)) {
  cat("Link #", kk, "\n", sep="")

  target <- targets[[kk]]
  cat("Target: ", target, "\n", sep="")

  # Name of *.lnk file
  pathname <- sprintf("%s.LNK", tempfile())

  tryCatch({
    # Will only work on Windows systems with support for VB scripting
    createWindowsShortcut(pathname, target=target)
  }, error = function(ex) {
    print(ex)
  })

  # Was it created?
  if (isFile(pathname)) {
    cat("Created link file: ", pathname, "\n", sep="")

    # Validate that it points to the correct target
    dest <- filePath(pathname, expandLinks="any")
    cat("Available target: ", dest, "\n", sep="")

    res <- all.equal(tolower(dest), tolower(target))
    if (!isTRUE(res)) {
      msg <- sprintf("Link target does not match expected target: %s != %s", dest, target)
      cat(msg, "\n")
      warning(msg)
    }
  }

  # Cleanup
```

```
    file.remove(pathname)
  }
}
```

---

dataFrame	<i>Allocates a data frame with given column classes</i>
-----------	---

---

### Description

Allocates a data frame with given column classes.

### Usage

```
## Default S3 method:
dataFrame(colClasses, nrow=1, ...)
```

### Arguments

colClasses	A <a href="#">character vector</a> of column classes, cf. <a href="#">read.table</a> .
nrow	An <a href="#">integer</a> specifying the number of rows of the allocated data frame.
...	Not used.

### Value

Returns an NxK [data.frame](#) where N equals nrow and K equals length(colClasses).

### See Also

[data.frame](#).

### Examples

```
df <- dataFrame(colClasses=c(a="integer", b="double"), nrow=10)
df[,1] <- sample(1:nrow(df))
df[,2] <- rnorm(nrow(df))
print(df)
```

---

detachPackage	<i>Detaches packages by name</i>
---------------	----------------------------------

---

**Description**

Detaches packages by name, if loaded.

**Usage**

```
## Default S3 method:  
detachPackage(pkgname, ...)
```

**Arguments**

pkgname	A <a href="#">character vector</a> of package names to be detached.
...	Not used.

**Value**

Returns (invisibly) a named [logical vector](#) indicating whether each package was detached or not.

**Author(s)**

Henrik Bengtsson

**See Also**

[detach\(\)](#).

---

dimNA< -	<i>Sets the dimension of an object with the option to infer one dimension automatically</i>
----------	---

---

**Description**

Sets the dimension of an object with the option to infer one dimension automatically. If one of the elements in the dimension [vector](#) is [NA](#), then its value is inferred from the length of the object and the other elements in the dimension vector. If the inferred dimension is not an [integer](#), an error is thrown.

**Usage**

```
## Default S3 replacement method:  
dimNA(x) <- value
```

**Arguments**

x	An R object.
value	NULL of a positive <a href="#">numeric vector</a> with one optional <a href="#">NA</a> .

**Value**

Returns (invisibly) what `dim<-()` returns (see [dim\(\)](#) for more details).

**Author(s)**

Henrik Bengtsson

**See Also**

[dim\(\)](#).

**Examples**

```
x <- 1:12
dimNA(x) <- c(2,NA,3)
stopifnot(dim(x) == as.integer(c(2,2,3)))
```

---

displayCode

*Displays the contents of a text file with line numbers and more*


---

**Description**

Displays the contents of a text file with line numbers and more.

**Usage**

```
## Default S3 method:
displayCode(con=NULL, code=NULL, numerate=TRUE, lines=-1, wrap=79, highlight=NULL,
  pager=getOption("pager"), ...)
```

**Arguments**

con	A <a href="#">connection</a> or a <a href="#">character</a> string filename. If code is specified, this argument is ignored.
code	A <a href="#">character vector</a> of code lines to be displayed.
numerate	If <a href="#">TRUE</a> , line are numbers, otherwise not.
lines	If a single <a href="#">numeric</a> , the maximum number of lines to show. If -1, all lines are shown. If a <a href="#">vector</a> of <a href="#">numeric</a> , the lines numbers to display.
wrap	The (output) column <a href="#">numeric</a> where to wrap lines.
highlight	A <a href="#">vector</a> of line number to be highlighted.

pager	If "none", code is not displayed in a pager, but only returned. For other options, see <a href="#">file.show()</a> .
...	Additional arguments passed to <a href="#">file.show()</a> , which is used to display the formatted code.

**Value**

Returns (invisibly) the formatted code as a [character](#) string.

**Author(s)**

Henrik Bengtsson

**See Also**

[file.show\(\)](#).

**Examples**

```
file <- system.file("DESCRIPTION", package="R.utils")
cat("Displaying: ", file, "\n", sep="")
displayCode(file)

file <- system.file("NEWS", package="R.utils")
cat("Displaying: ", file, "\n", sep="")
displayCode(file, numerate=FALSE, lines=100:110, wrap=65)

file <- system.file("NEWS", package="R.utils")
cat("Displaying: ", file, "\n", sep="")
displayCode(file, lines=100:110, wrap=65, highlight=c(101,104:108))
```

---

doCall

*Executes a function call with option to ignore unused arguments*

---

**Description**

Executes a function call with option to ignore unused arguments.

**Usage**

```
## Default S3 method:
doCall(.fcn, ..., args=NULL, alwaysArgs=NULL, .functions=list(.fcn),
       .ignoreUnusedArgs=TRUE, envir=parent.frame())
```

**Arguments**

<code>.fcn</code>	A <a href="#">function</a> or a <a href="#">character</a> string specifying the name of a <a href="#">function</a> to be called.
<code>...</code>	Named arguments to be passed to the function.
<code>args</code>	A <a href="#">list</a> of additional named arguments that will be appended to the above arguments.
<code>alwaysArgs</code>	A <a href="#">list</a> of additional named arguments that will be appended to the above arguments and that will <i>never</i> be ignore.
<code>.functions</code>	A <a href="#">list</a> of <a href="#">function</a> :s or names of functions. This can be used to control which arguments are passed.
<code>.ignoreUnusedArgs</code>	If <a href="#">TRUE</a> , arguments that are not accepted by the function, will not be passed to it. Otherwise, all arguments are passed.
<code>envir</code>	An <a href="#">environment</a> in which to evaluate the call.

**Author(s)**

Henrik Bengtsson

**See Also**

[do.call\(\)](#).

**Examples**

```
doCall("plot", x=1:10, y=sin(1:10), col="red", dummyArg=54,
      alwaysArgs=list(xlab="x", ylab="y"),
      .functions=c("plot", "plot.xy"))
```

---

downloadFile.character

*Downloads a file*

---

**Description**

Downloads a file.

**Usage**

```
## S3 method for class 'character'
downloadFile(url, filename=basename(url), path=NULL, skip=TRUE, overwrite=!skip, ...,
  username=NULL, password=NULL, binary=TRUE, dropEmpty=TRUE, verbose=FALSE)
```



## Arguments

url	A <a href="#">character</a> string specifying the URL to be downloaded.
filename, path	(optional) <a href="#">character</a> strings specifying the local filename and the path of the downloaded file.
skip	If <a href="#">TRUE</a> , an already downloaded file is skipped.
overwrite	If <a href="#">TRUE</a> , an already downloaded file is overwritten, otherwise an error is thrown.
...	Additional arguments passed to <a href="#">download.file</a> .
username, password	<a href="#">character</a> strings specifying the username and password for authenticated downloads. The alternative is to specify these via the URL.
binary	If <a href="#">TRUE</a> , the file is downloaded exactly "as is", that is, byte by byte (recommended).
dropEmpty	If <a href="#">TRUE</a> and the downloaded file is empty, the file is ignored and <a href="#">NULL</a> is returned.
verbose	A <a href="#">logical</a> , <a href="#">integer</a> , or a <a href="#">Verbose</a> object.

## Details

Currently arguments `username` and `password` are only used for downloads via URL protocol 'https'. The 'https' protocol requires that either of 'curl' or 'wget' are available on the system.

## Value

Returns the local pathname to the downloaded filename, or [NULL](#) if no file was downloaded.

## Author(s)

Henrik Bengtsson

## See Also

Internally [download.file](#) is used. That function may generate an empty file if the URL is not available.

## Examples

```
## Not run:
pathname <- downloadFile("http://www.r-project.org/index.html", path="www.r-project.org/")
print(pathname)

## End(Not run)
```

---

egsub

*Global substitute of expression using regular expressions*

---

## Description

Global substitute of expression using regular expressions.

## Usage

```
egsub(pattern, replacement, x, ..., value=TRUE, envir=parent.frame(), inherits=TRUE)
```

## Arguments

pattern	A <a href="#">character</a> string with the regular expression to be matched, cf. <a href="#">gsub()</a> .
replacement	A <a href="#">character</a> string of the replacement to use when there is a match, cf. <a href="#">gsub()</a> .
x	The <a href="#">expression</a> or a <a href="#">function</a> to be modified.
...	Additional arguments passed to <a href="#">gsub()</a>
value	If <a href="#">TRUE</a> , the value of the replacement itself is used to look up a variable with that name and then using that variable's value as the replacement. Otherwise the replacement value is used.
envir, inherits	An <a href="#">environment</a> from where to find the variable and whether the search should also include enclosing frames, cf. <a href="#">get()</a> . Only use if <a href="#">value</a> is <a href="#">TRUE</a> .

## Value

Returns an [expression](#).

## Author(s)

Henrik Bengtsson

## Examples

```
# Original expression
expr <- substitute({
  res <- foo.bar.yaa(2)
  print(res)
  R.utils::use("R.oo")
  x <- .b.
})

# Some predefined objects
foo.bar.yaa <- function(x) str(x)
a <- 2
b <- a
```

```

# Substitute with variable name
expr2 <- egsub("^.[.]([a-zA-Z0-9_]+)[.]$", "\\1", expr, value=FALSE)
print(expr2)
## {
##   res <- foo.bar.yaa(2)
##   print(res)
##   R.utils::use("R.oo")
##   x <- b
## }

# Substitute with variable value
expr3 <- egsub("^.[.]([a-zA-Z0-9_]+)[.]$", "\\1", expr, value=TRUE)
print(expr3)
## {
##   res <- foo.bar.yaa(2)
##   print(res)
##   R.utils::use("R.oo")
##   x <- 2
## }

# Substitute the body of a function
warnifnot <- egsub("stop", "warning", stopifnot, value=FALSE)
print(warnifnot)
warnifnot(pi == 3.14)

```

---

env	<i>Creates a new environment, evaluates an expression therein, and returns the environment</i>
-----	--

---

## Description

Creates a new environment, evaluates an expression therein, and returns the environment.

## Usage

```
env(..., hash=FALSE, parent=parent.frame(), size=29L)
```

## Arguments

... Arguments passed to `evalq()`, particularly a [expression](#) to be evaluated inside the newly created [environment](#).

hash, parent, size Arguments passed to `new.env()`.

## Value

Returns an [environment](#).

## Author(s)

Henrik Bengtsson

## References

[1] R-devel thread 'Create an environment and assign objects to it in one go?' on March 9-10, 2011.

## See Also

Internally `new.env()` and `evalq()` are used.

## Examples

```
x <- list();

x$case1 <- env({
  # Cut'n'pasted from elsewhere
  a <- 1;
  b <- 2;
});

x$case2 <- env({
  # Cut'n'pasted from elsewhere
  foo <- function(x) x^2;
  a <- foo(2);
  b <- 1;
  rm(foo); # Not needed anymore
});

# Turn into a list of lists
x <- lapply(x, FUN=as.list);

str(x);
```

---

extract.array	<i>Extract a subset of an array, matrix or a vector with unknown dimensions</i>
---------------	---

---

## Description

Extract a subset of an array, matrix or a vector with unknown dimensions.

This method is useful when you do not know the number of dimensions of the object your wish to extract values from, cf. example.

## Usage

```
## S3 method for class 'array'
extract(x, ..., indices=list(...), dims=names(indices), drop=FALSE)
```

**Arguments**

x	An <a href="#">array</a> or a <a href="#">matrix</a> .
...	These arguments are by default put into the indices <a href="#">list</a> .
indices	A <a href="#">list</a> of index <a href="#">vectors</a> to be extracted.
dims	An <a href="#">vector</a> of dimensions - one per element in indices - which will be coerced to <a href="#">integers</a> . If <code>NULL</code> , it will default to <code>seq_along(indices)</code> .
drop	If <code>TRUE</code> , dimensions of length one are dropped, otherwise not.

**Value**

Returns an [array](#).

**Author(s)**

Henrik Bengtsson

**See Also**

[slice.index\(\)](#)

**Examples**

```
# -----
# Example using an array with a random number of dimensions
# -----
maxdim <- 4
dim <- sample(3:maxdim, size=sample(2:maxdim, size=1), replace=TRUE)
ndim <- length(dim)
dimnames <- list()
for (kk in 1:ndim)
  dimnames[[kk]] <- sprintf("%s%d", letters[kk], 1:dim[kk])
x <- 1:prod(dim)
x <- array(x, dim=dim, dimnames=dimnames)

cat("\nArray 'x':\n")
print(x)

cat("\nExtract 'x[2:3,...]':\n")
print(extract(x, "1"=2:3))

cat("\nExtract 'x[3,2:3,...]':\n")
print(extract(x, "1"=3,"2"=2:3))

cat("\nExtract 'x[... ,2:3]':\n")
print(extract(x, indices=2:3, dims=length(dim(x))))

# -----
```

```

# Assertions
# -----
y <- array(1:24, dim=c(2,3,4))
yA <- y[, ,2:3]
yB <- extract(y, indices=list(2:3), dims=length(dim(y)))
stopifnot(identical(yB, yA))

yA <- y[,2:3,2]
yB <- extract(y, indices=list(2:3,2), dims=c(2,3), drop=TRUE)
stopifnot(identical(yB, yA))

```

---

fileAccess

*Checks the permission of a file or a directory*


---

### Description

Checks the permission of a file or a directory.

### Usage

```

## Default S3 method:
fileAccess(pathname, mode=0, safe=TRUE, ...)

```

### Arguments

pathname	A <a href="#">character</a> string of the file or the directory to be checked.
mode	An <a href="#">integer</a> (0,1,2,4), cf. <a href="#">file.access()</a> .
safe	If <a href="#">TRUE</a> , the permissions are tested more carefully, otherwise <a href="#">file.access()</a> is used.
...	Not used.

### Details

In R there is [file.access\(\)](#) for checking whether the permission of a file. Unfortunately, that function cannot be 100% trusted depending on platform used and file system queried, cf. [1].

### Value

Returns an [integer](#); 0 if the permission exists, -1 if not.

### Symbolic links

This function follows symbolic links (also on Windows) and returns a value based on the link target (rather than the link itself).

### Author(s)

Henrik Bengtsson

## References

- [1] R-devel thread *file.access() on network (mounted) drive on Windows Vista?* on Nov 26, 2008. <https://stat.ethz.ch/pipermail/r-devel/2008-December/051461.html>
- [2] Filesystem permissions, Wikipedia, 2010. [http://en.wikipedia.org/wiki/Filesystem\\_permissions](http://en.wikipedia.org/wiki/Filesystem_permissions)

## See Also

[file.access\(\)](#)

## Examples

```
# - - - - -
# Current directory
# - - - - -
path <- "."

# Test for existence
print(fileAccess(path, mode=0))
# Test for execute permission
print(fileAccess(path, mode=1))
# Test for write permission
print(fileAccess(path, mode=2))
# Test for read permission
print(fileAccess(path, mode=4))

# - - - - -
# A temporary file
# - - - - -
pathname <- tempfile()
cat(file=pathname, "Hello world!")

# Test for existence
print(fileAccess(pathname, mode=0))
# Test for execute permission
print(fileAccess(pathname, mode=1))
# Test for write permission
print(fileAccess(pathname, mode=2))
# Test for read permission
print(fileAccess(pathname, mode=4))

file.remove(pathname)

# - - - - -
# The 'base' package directory
# - - - - -
path <- system.file(package="base")

# Test for existence
```

```

print(fileAccess(path, mode=0))
# Test for execute permission
print(fileAccess(path, mode=1))
# Test for write permission
print(fileAccess(path, mode=2))
# Test for read permission
print(fileAccess(path, mode=4))

# -----
# The 'base' package DESCRIPTION file
# -----
pathname <- system.file("DESCRIPTION", package="base")

# Test for existence
print(fileAccess(pathname, mode=0))
# Test for execute permission
print(fileAccess(pathname, mode=1))
# Test for write permission
print(fileAccess(pathname, mode=2))
# Test for read permission
print(fileAccess(pathname, mode=4))

```

---

filePath	<i>Construct the path to a file from components and expands Windows Shortcuts along the pathname from root to leaf</i>
----------	--

---

## Description

Construct the path to a file from components and expands Windows Shortcuts along the pathname from root to leaf. This function is backward compatible with `file.path()` when argument `removeUps=FALSE` and `expandLinks="none"`, except that a (character) `NA` is return if any argument is `NA`.

This function exists on all platforms, not only Windows systems.

## Usage

```

## Default S3 method:
filePath(..., fsep=.Platform$file.sep, removeUps=TRUE,
  expandLinks=c("none", "any", "local", "relative", "network"), unmap=FALSE,
  mustExist=FALSE, verbose=FALSE)

```

## Arguments

...	Arguments to be pasted together to a file path and then be parsed from the root to the leaf where Windows shortcut files are recognized and expanded according to argument which in each step.
fsep	the path separator to use.



removeUps	If <b>TRUE</b> , relative paths, for instance "foo/bar/../" are shortened into "foo/", but also "../" are removed from the final pathname, if possible.
expandLinks	A <b>character</b> string. If "none", Windows Shortcut files are ignored. If "local", the absolute target on the local file system is used. If "relative", the relative target is used. If "network", the network target is used. If "any", first the local, then the relative and finally the network target is searched for.
unmap	If <b>TRUE</b> , paths on mapped Windows drives are "followed" and translated to their corresponding "true" paths.
mustExist	If <b>TRUE</b> and if the target does not exist, the original pathname, that is, argument pathname is returned. In all other cases the target is returned.
verbose	If <b>TRUE</b> , extra information is written while reading.

### Details

If `expandLinks != "none"`, each component, call it *parent*, in the absolute path is processed from the left to the right as follows: 1. If a "real" directory of name *parent* exists, it is followed. 2. Otherwise, if Microsoft Windows Shortcut file with name *parent.lnk* exists, it is read. If its local target exists, that is followed, otherwise its network target is followed. 3. If no valid existing directory was found in (1) or (2), the expanded this far followed by the rest of the pathname is returned quietly. 4. If all of the absolute path was expanded successfully the expanded absolute path is returned.

### Value

Returns a **character** string.

### On speed

Internal `file.exists()` is call while expanding the pathname. This is used to check if there exists a Windows shortcut file named 'foo.lnk' in 'path/foo/bar'. If it does, 'foo.lnk' has to be followed, and in other cases 'foo' is ordinary directory. The `file.exists()` is unfortunately a bit slow, which is why this function appears slow if called many times.

### Author(s)

Henrik Bengtsson

### See Also

[readWindowsShellLink\(\)](#). [readWindowsShortcut\(\)](#). [file.path\(\)](#).

### Examples

```
# Default
print(filePath("foo", "bar", "..", "name")) # "foo/bar/../name"

# Shorten pathname, if possible
print(filePath("foo", "bar", "..", "name")) # "foo/name"
print(filePath("foo/bar/../name"))       # "foo/name"
```

```
# Recognize Windows Shortcut files along the path, cf. Unix soft links
filename <- system.file("data-ex/HISTORY.LNK", package="R.utils")
print(filename)
filename <- filePath(filename, expandLinks="relative")
print(filename)
```

---

FileProgressBar      *A progress bar that sets the size of a file accordingly*

---

## Description

Package: R.utils

**Class FileProgressBar**

Object

```
~~|
~~+--ProgressBar
~~~~~|
~~~~~+--FileProgressBar
```

**Directly known subclasses:**

```
public static class FileProgressBar
extends ProgressBar
```

## Usage

```
FileProgressBar(pathname=NULL, ...)
```

## Arguments

pathname      The pathname of the output file.  
 ...            Other arguments accepted by the [ProgressBar](#) constructor.

## Details

A progress bar that sets the size of a file accordingly. This class useful to check the progress of a batch job by just querying the size of a file, for instance, via ftp.

## Fields and Methods

**Methods:**

`cleanup` Removes the progress file for a file progress bar.  
`update` Updates file progress bar.

**Methods inherited from ProgressBar:**

`as.character`, `getBarString`, `increase`, `isDone`, `reset`, `setMaxValue`, `setProgress`, `setStepLength`, `setTicks`, `setValue`, `update`

**Methods inherited from Object:**

`$`, `$<-`, `[[`, `[[<-`, `as.character`, `attach`, `attachLocally`, `clearCache`, `clearLookupCache`, `clone`, `detach`, `equals`, `extend`, `finalize`, `getEnvironment`, `getFieldModifier`, `getFieldModifiers`, `getFields`, `getInstantiationTime`, `getStaticInstance`, `hasField`, `hashCode`, `ll`, `load`, `names`, `objectSize`, `print`, `save`

**Author(s)**

Henrik Bengtsson

**Examples**

```
## Not run:

# Creates a progress bar (of length 100) that displays it self as a file.
pb <- FileProgressBar("~/progress.simulation")
reset(pb)
while (!isDone(pb)) {
  x <- rnorm(3e4)
  increase(pb)
  # Emulate a slow process
  if (interactive()) Sys.sleep(0.1)
  Sys.sleep(0.01)
}

## End(Not run)
```

---

`finalizeSession`      *Function to call for finalizing the R session*

---

**Description**

Function to call for finalizing the R session. When called, all registered "onSessionExit" hooks (functions) are called. To define such hooks, use the `onSessionExit()` function.

This method should not be used by the user.

**Usage**

```
## Default S3 method:
finalizeSession(...)
```

**Arguments**

... Not used.

**Value**

Returns (invisibly) the hooks successfully called.

**Author(s)**

Henrik Bengtsson

**See Also**

[onSessionExit\(\)](#).

---

findSourceTraceback *Finds all 'srcfile' objects generated by source() in all call frames*

---

**Description**

Finds all 'srcfile' objects generated by source() in all call frames. This makes it possible to find out which files are currently scripted by [source\(\)](#).

**Usage**

```
## Default S3 method:  
findSourceTraceback(...)
```

**Arguments**

... Not used.

**Value**

Returns a named list of [srcfile\(\)](#) objects and/or [character](#) strings. The names of the list entries corresponds to the 'filename' value of each corresponding 'srcfile' object. The returned list is empty if [source\(\)](#) was not called.

**Author(s)**

Henrik Bengtsson

**See Also**

See also [sourceutils](#).

## Examples

```

# -----
# Create two R script files where one source():s the other
# and both lists the traceback of filenames source():d.
# -----
path <- tempdir();
pathnameA <- Arguments$getWritablePathname("foo.R", path=path);
pathnameB <- Arguments$getWritablePathname("bar.R", path=path);

code <- 'cat("BEGIN foo.R\n")';
code <- c(code, 'print(findSourceTraceback());');
code <- c(code, sprintf('source("%s");', pathnameB));
code <- c(code, 'cat("END foo.R\n")');
code <- paste(code, collapse="\n");
cat(file=pathnameA, code);

code <- 'cat("BEGIN bar.R\n")';
code <- c(code, 'x <- findSourceTraceback();');
code <- c(code, 'print(x);');
code <- c(code, 'cat("END bar.R\n")');
code <- paste(code, collapse="\n");
cat(file=pathnameB, code);

# -----
# Source the first file
# -----
source(pathnameA, echo=TRUE);

```

---

gcat

*Parses, evaluates and outputs a GString*


---

## Description

Parses, evaluates and outputs a GString.

## Usage

```

## Default S3 method:
gcat(..., file="", append=FALSE, envir=parent.frame())

```

## Arguments

...	character strings passed to <code>gstring()</code> .
file	A <a href="#">connection</a> , or a pathname where to direct the output. If "", the output is sent to the standard output.
append	Only applied if file specifies a pathname. If <code>TRUE</code> , then the output is appended to the file, otherwise the files content is overwritten.
envir	The <a href="#">environment</a> in which the <code>GString</code> is evaluated.

**Value**

Returns (invisibly) a `character` string.

**Author(s)**

Henrik Bengtsson

**See Also**

`gstring()`.

---

gcDLLs

*Identifies and removes DLLs of packages already unloaded*

---

**Description**

Identifies and removes DLLs of packages already unloaded. When packages are unloaded, they are ideally also unloading any DLLs (also known as a dynamic shared object or library) they have loaded. Unfortunately, not all package do this resulting in "stray" DLLs still being loaded and occupying R's limited registry. These functions identifies and removes such DLLs.

**Usage**

```
gcDLLs(gc=TRUE, quiet=TRUE)
```

**Arguments**

<code>gc</code>	If <code>TRUE</code> , if there are stray DLLs, then the garbage collector is run before unloading those DLLs. This is done in order to trigger any finalizers, of which some may need those DLLs, to be called.
<code>quiet</code>	If <code>FALSE</code> , a message is outputted for every stray DLL that is unloaded.

**Details**

If a library fails to unload, an informative warning is generated.

**Value**

Returns (invisibly) the set of stray DLLs identified.

**How to unload DLLs in package (for package developers)**

To unload a package DLL whenever the package is unloaded, add the following to your package:

```
.onUnload <- function(libpath) {  
  ## (1) Force finalizers to be called before removing the DLL  
  ##      in case some of them need the DLL.  
  gc()  
  
  ## (2) Unload the DLL for this package  
  library.dynam.unload(.packageName, libpath)  
}
```

**Author(s)**

Henrik Bengtsson

**See Also**

[getLoadedDLLs\(\)](#).

---

getAbsolutePath	<i>Gets the absolute pathname string</i>
-----------------	--

---

**Description**

Gets the absolute pathname string.

**Usage**

```
## Default S3 method:  
getAbsolutePath(pathname, workDirectory=getwd(), expandTilde=FALSE, ...)
```

**Arguments**

pathname	A <a href="#">character</a> string of the pathname to be converted into an absolute pathname.
workDirectory	A <a href="#">character</a> string of the current working directory.
expandTilde	If <a href="#">TRUE</a> , tilde (~) is expanded to the corresponding directory, otherwise not.
...	Not used.

**Details**

This method will replace replicated slashes ( '/') with a single one, except for the double forward slashes prefixing a Microsoft Windows UNC (Universal Naming Convention) pathname.

**Value**

Returns a [character](#) string of the absolute pathname.

**Author(s)**

Henrik Bengtsson

**See Also**

[isAbsolutePath\(\)](#).

---

getParent

*Gets the string of the parent specified by this pathname*

---

**Description**

Gets the string of the parent specified by this pathname. This is basically, by default the string before the last path separator of the absolute pathname.

**Usage**

```
## Default S3 method:  
getParent(pathname, depth=1L, fsep=.Platform$file.sep, ...)
```

**Arguments**

pathname	A <a href="#">character</a> string of the pathname to be checked.
depth	An <a href="#">integer</a> specifying how many generations up the path should go.
fsep	A <a href="#">character</a> string of the file separator.
...	Not used.

**Value**

Returns a [character](#) string if the parent exists, otherwise [NULL](#).

**Author(s)**

Henrik Bengtsson



---

getRelativePath	<i>Gets the relative pathname relative to a directory</i>
-----------------	---

---

### Description

Gets the relative pathname relative to a directory.

### Usage

```
## Default S3 method:  
getRelativePath(pathname, relativeTo=getwd(), caseSensitive=NULL, ...)
```

### Arguments

pathname	A <a href="#">character</a> string of the pathname to be converted into an relative pathname.
relativeTo	A <a href="#">character</a> string of the reference pathname.
caseSensitive	If <a href="#">TRUE</a> , the comparison is case sensitive, otherwise not. If <a href="#">NULL</a> , it is decided from the relative path.
...	Not used.

### Details

In case the two paths are on different file systems, for instance, C:/foo/bar/ and D:/foo/, the method returns pathname as is.

### Value

Returns a [character](#) string of the relative pathname.

### Non-case sensitive comparison

If `caseSensitive == NULL`, the relative path is used to decide if the comparison should be done in a case-sensitive mode or not. The current check is if it is a Windows path or not, that is, if the relative path starts with a device letter, then the comparison is non-case sensitive.

### Author(s)

Henrik Bengtsson

### See Also

[getAbsolutePath\(\)](#). [isAbsolutePath\(\)](#).

**Examples**

```

getRelativePath("foo", "foo")           # "."
getRelativePath("foo/bar", "foo")      # "bar"
getRelativePath("foo/bar", "foo/bar/yah") # ".."
getRelativePath("foo/bar/cool", "foo/bar/yah/sub/") # "../../cool"
getRelativePath("/tmp/bar/", "/bar/foo/") # "../../tmp/bar"

# Windows
getRelativePath("C:/foo/bar/", "C:/bar/") # "../foo/bar"
getRelativePath("C:/foo/bar/", "D:/bar/") # "C:/foo/bar"

```

GString

*Character string with advanced substitutions***Description**

Package: R.utils

**Class GString**

character

~~|

~~+--GString

**Directly known subclasses:**public static class **GString**

extends character

**Usage**

GString(..., sep="")

**Arguments**... one or more objects, to be coerced to [character](#) vectors.sep A [character](#) string to separate the terms.**Fields and Methods****Methods:**[as.character](#)

Gets the processed character string.

[evaluate](#)

Parses and evaluates a GString.

[gcat](#)

-

<code>getBuiltinDate</code>	Gets the current date.
<code>getBuiltinDatetime</code>	Gets the current date and time.
<code>getBuiltinHostname</code>	Gets the hostname of the system running R.
<code>getBuiltinOs</code>	Gets the operating system of the running machine.
<code>getBuiltinPid</code>	Gets the process id of the current R session.
<code>getBuiltinRhome</code>	Gets the path where R is installed.
<code>getBuiltinRversion</code>	Gets the current R version.
<code>getBuiltinTime</code>	Gets the current time.
<code>getBuiltinUsername</code>	Gets the username of the user running R.
<code>getRaw</code>	Gets the unprocessed GString.
<code>getVariableValue</code>	Gets a variable value given a name and attributes.
<code>gstring</code>	-
<code>parse</code>	Parses a GString.
<code>print</code>	Prints the processed GString.

**Methods inherited from character:**

Ops,nonStructure,vector-method, Ops,structure,vector-method, Ops,vector,nonStructure-method, Ops,vector,structure-method, all.equal, as.Date, as.POSIXlt, as.data.frame, as.raster, coerce,ANY,character-method, coerce,character,SuperClassM method, coerce,character,signature-method, coerce<-,ObjectsWithPackage,character-method, coerce<- ,signature,character-method, downloadFile, formula, getDLLRegisteredRoutines, isOpen, toAsciiRegExprPattern, toFileListTree, uses

**Author(s)**

Henrik Bengtsson

**See Also**

For convenience, see functions `gstring()` and `gcat()`.

**Examples**

```
# -----
# First example
# -----
who <- "world"

# Compare this...
cat(as.character(GString("Hello ${who}\n")))

# ...to this.
cat(GString("Hello ${who}\n"))

# Escaping
cat(as.character(GString("Hello \${who}\n")))

# -----
# Looping over vectors
```

```

# -----
x <- 1:5
y <- c("hello", "world")
cat(as.character(GString("(x,y)={x},{y}")), sep=" ", )
cat("\n")

cat(as.character(GString("(x,y)={x},{capitalize}{y}")), sep=" ", )
cat("\n")

# -----
# Predefined ("builtin") variables
# -----
cat(as.character(GString("Hello ${username} on host ${hostname} running ",
"R v${rversion} in process #${pid} on ${os}. R is installed in ${rhome}.")))

# Other built-in variables/functions...
cat(as.character(GString("Current date: ${date}\n")))
cat(as.character(GString("Current date: ${format='%d/%m/%y'}{date}\n")))
cat(as.character(GString("Current time: ${time}\n")))

# -----
# Evaluating inline R code
# -----
cat(as.character(GString("Simple calculation: 1+1=${1+1}\n")))
cat(as.character(GString("Alternative current date: ${`date()}`\n")))

# -----
# Function values
# -----
# Call function rnorm with arguments n=1, i.e. rnorm(n=1)
cat(as.character(GString("Random normal number: ${n=1}{rnorm}\n")))

# -----
# Global search-replace feature
# -----
# Replace all '-' with '.'
cat(as.character(GString("Current date: ${date/-/}\n")))
# Another example
cat(as.character(GString("Escaped string: 12*12=${`12*12`/1/}\n")))

# -----
# Defining new "builtin" function values
# -----
# Define your own builtin variables (functions)
setMethodS3("getBuiltinAletter", "GString", function(object, ...) {
  base::letters[runif(1, min=1, max=length(base::letters))]
})

```

```

cat(as.character(GString("A letter: ${aletter}\n")))
cat(as.character(GString("Another letter: ${aletter}\n")))

# Another example
setMethodS3("getBuiltinGString", "GString", function(object, ...) {
  # Return another GString.
  GString("${date} ${time}")
})

cat(as.character(GString("Advanced example: ${gstring}\n")))

# Advanced example
setMethodS3("getBuiltinRunif", "GString", function(object, n=1, min=0, max=1, ...) {
  formatC(runif(n=n, min=min, max=max), ...)
})

cat(as.character(GString("A random number: ${runif}\n")))
n <- 5
cat(as.character(GString("${n} random numbers: ")))
cat(as.character(GString("${n=n, format='f'}{runif}")))
cat("\n")

# Advanced options.
# Options are parsed as if they are elements in a list, e.g.
# list(n=runif(n=1,min=1,max=5), format='f')
cat(as.character(GString("$Random number of numbers: ")))
cat(as.character(GString("${n=runif(n=1,min=1,max=5), format='f'}{runif}")))
cat("\n")

```

---

gstring

*Parses and evaluates a GString into a regular string*


---

## Description

Parses and evaluates a GString into a regular string.

## Usage

```
## Default S3 method:
gstring(..., file=NULL, path=NULL, envir=parent.frame())
```

## Arguments

...	<a href="#">character</a> strings.
file, path	Alternatively, a file, a URL or a <a href="#">connection</a> from with the strings are read. If a file, the path is prepended to the file, iff given.
envir	The <a href="#">environment</a> in which the <a href="#">GString</a> is evaluated.

**Value**

Returns a [character](#) string.

**Author(s)**

Henrik Bengtsson

**See Also**

[gcat\(\)](#).

---

hasUrlProtocol	<i>Checks if one or several pathnames has a URL protocol</i>
----------------	--

---

**Description**

Checks if one or several pathnames has a URL protocol.

**Usage**

```
## Default S3 method:  
hasUrlProtocol(pathname, ...)
```

**Arguments**

pathname	A <a href="#">character vector</a> .
...	Not used.

**Value**

Returns a [logical vector](#).

**Author(s)**

Henrik Bengtsson

---

hpaste

*Concatenating vectors into human-readable strings*

---

## Description

Concatenating vectors into human-readable strings such as "1, 2, 3, ..., 10".

## Usage

```
## Default S3 method:
hpaste(..., sep="", collapse=", ", lastCollapse=NULL,
        maxHead=if (missing(lastCollapse)) 3 else Inf,
        maxTail=if (is.finite(maxHead)) 1 else Inf, abbreviate="...")
```

## Arguments

`...` Arguments to be pasted.

`sep` A [character](#) string used to concatenate the arguments in `...`, if more than one.

`collapse`, `lastCollapse`  
The [character](#) strings to collapse the elements together, where `lastCollapse` is specifying the collapse string used between the last two elements. If `lastCollapse` is `NULL` (default), it corresponds to using the default collapse.

`maxHead`, `maxTail`, `abbreviate`  
Non-negative [integers](#) (also [Inf](#)) specifying the maximum number of elements of the beginning and then end of the vector to be outputted. If `n = length(x)` is greater than `maxHead+maxTail+1`, then `x` is truncated to consist of `x[1:maxHead]`, `abbreviate`, and `x[(n-maxTail+1):n]`.

## Details

`hpaste(..., sep=" ", maxHead=Inf)` corresponds to `paste(..., sep=" ", collapse=", ")`.

## Value

Returns a [character](#) string.

## Author(s)

Henrik Bengtsson

## See Also

Internally [paste\(\)](#) is used.

**Examples**

```

# Some vectors
x <- 1:6
y <- 10:1
z <- LETTERS[x]

# -----
# Abbreviation of output vector
# -----
printf("x = %s.\n", hpaste(x))
## x = 1, 2, 3, ..., 6.

printf("x = %s.\n", hpaste(x, maxHead=2))
## x = 1, 2, ..., 6.

printf("x = %s.\n", hpaste(x, maxHead=3) # Default
## x = 1, 2, 3, ..., 6.

# It will never output 1, 2, 3, 4, ..., 6
printf("x = %s.\n", hpaste(x, maxHead=4))
## x = 1, 2, 3, 4, 5, 6.

# Showing the tail
printf("x = %s.\n", hpaste(x, maxHead=1, maxTail=2))
## x = 1, ..., 5, 6.

# Turning off abbreviation
printf("y = %s.\n", hpaste(y, maxHead=Inf))
## y = 10, 9, 8, 7, 6, 5, 4, 3, 2, 1

## ...or simply
printf("y = %s.\n", paste(y, collapse=", "))
## y = 10, 9, 8, 7, 6, 5, 4, 3, 2, 1

# -----
# Adding a special separator before the last element
# -----
# Change last separator
printf("x = %s.\n", hpaste(x, lastCollapse=", and "))
## x = 1, 2, 3, 4, 5, and 6.

# -----
# Backward compatibility with paste()
# -----
s1 <- hpaste(x, maxHead=Inf)
s2 <- paste(x, collapse=", ")
printf("s = %s.\n", s1);
stopifnot(identical(s1, s2))

```



```
s1 <- hpaste('<', x, '>', maxHead=Inf)
s2 <- paste('<', x, '>', sep="", collapse=", ")
printf("s = %s.\n", s1);
stopifnot(identical(s1, s2))

s1 <- hpaste(x, y, z, sep="/", maxHead=Inf)
s2 <- paste(x, y, z, sep="/", collapse=", ")
printf("s = %s.\n", s1);
stopifnot(identical(s1, s2))

s1 <- hpaste(x, collapse=NULL, maxHead=Inf)
s2 <- paste(x, collapse=NULL)
stopifnot(identical(s1, s2))
```

---

inAnyInterval.numeric *Checks if a set of values are inside one or more intervals*

---

## Description

Checks if a set of values are inside one or more intervals.

## Usage

```
## S3 method for class 'numeric'
inAnyInterval(...)
```

## Arguments

... Arguments passed to [\\*mapToIntervals\(\)](#).

## Value

Returns a [logical vector](#).

## Author(s)

Henrik Bengtsson

## See Also

[mapToIntervals\(\)](#).

---

insert	<i>Insert values to a vector at certain positions</i>
--------	---

---

**Description**

Insert values to a vector at certain positions.

**Usage**

```
## Default S3 method:
insert(x, ats, values=NA, useNames=TRUE, ...)
```

**Arguments**

x	The <a href="#">vector</a> of data values.
ats	The indices of x where the values should be inserted.
values	A <a href="#">list</a> or a <a href="#">vector</a> of the values to be inserted. Should be of same length as ats, unless if a single value when it is automatically extended without a warning.
useNames	If <a href="#">FALSE</a> , the names attribute is dropped/ignored, otherwise not. Only applied if argument x is named.
...	Not used.

**Author(s)**

Henrik Bengtsson

**See Also**

[append\(\)](#) takes argument after (a scalar). For example, `append(x, y, after=after) == insert(x, values=y, ats=after)`. Contrary to `append()`, `insert()` accepts a vector of insert indices.

**Examples**

```
# Insert NAs (default) between all values
y <- c(a=1, b=2, c=3)
print(y)
x <- insert(y, ats=2:length(y))
Ex <- c(y[1], NA_real_, y[2], NA_real_, y[3])
print(x)
stopifnot(identical(x,Ex))

# Insert at first position
y <- c(a=1, b=2, c=3)
print(y)
x <- insert(y, ats=1, values=rep(NA_real_,2))
Ex <- c(NA_real_,NA_real_,y)
print(x)
```

```
stopifnot(identical(x,Ex))

x <- insert(y, ats=1, values=rep(NA_real_,2), useNames=FALSE)
print(x)

# Insert at last position (names of 'values' are ignored
# because input vector has no names)
x <- insert(1:3, ats=4, values=c(d=2, e=1))
Ex <- c(1:3,2,1)
print(x)
stopifnot(identical(x,Ex))

# Insert in the middle of a vector
x <- insert(c(1,3,2,1), ats=2, values=2)
print(x)
stopifnot(identical(as.double(x),as.double(Ex)))

# Insert multiple vectors at multiple indices at once
x0 <- c(1:4, 8:11, 13:15)

x <- insert(x0, at=c(5,9), values=list(5:7,12))
print(x)
Ex <- 1:max(x)
stopifnot(identical(as.double(x),as.double(Ex)))

x <- insert(x0, at=c(5,9,12), values=list(5:7,12,16:18))
print(x)
Ex <- 1:max(x)
stopifnot(identical(as.double(x),as.double(Ex)))

# Insert missing indices
Ex <- 1:20
missing <- setdiff(Ex, x0)
x <- x0
for (m in missing)
  x <- insert(x, ats=m, values=m)
print(x)
stopifnot(identical(as.double(x),as.double(Ex)))
```

---

installPackages

*Install R packages by name or URL*

---

## Description

Install R packages by name or URL.

## Usage

```
## Default S3 method:  
installPackages(pkgs, types="auto", repos=getOption("repos"), ..., destPath=".",  
cleanup=TRUE)
```

## Arguments

pkgs	A <a href="#">character vector</a> specifying the names and/or the URLs of the R packages to be installed.
types	A <a href="#">character vector</a> of corresponding package types.
repos	A <a href="#">character vector</a> of package repository URLs.
...	Additional arguments passed to <a href="#">install.packages</a> .
destPath	Path where any downloaded files are saved.
cleanup	If <a href="#">TRUE</a> , downloaded and successfully installed package files are removed, otherwise not.

## Value

Returns nothing.

## Limitations

This method cannot install any packages that are already in use. Certain packages are always in use when calling this method, e.g. **R.methodsS3**, **R.oo**, and **R.utils**.

## Author(s)

Henrik Bengtsson

## Examples

```
## Not run:  
installPackages("R.rsp")  
installPackages("http://cran.r-project.org/src/contrib/Archive/R.rsp/R.rsp_0.8.2.tar.gz")  
installPackages("http://cran.r-project.org/bin/windows/contrib/r-release/R.rsp_0.9.17.zip")  
  
## End(Not run)
```

---

intervalsToSeq.matrix *Generates a vector of indices from a matrix of intervals*

---

**Description**

Generates a vector of indices from a matrix of intervals.

**Usage**

```
## S3 method for class 'matrix'  
intervalsToSeq(fromTo, sort=FALSE, unique=FALSE, ...)
```

**Arguments**

fromTo	An Nx2 <a href="#">integer matrix</a> .
sort	If <a href="#">TRUE</a> , the returned indices are ordered.
unique	If <a href="#">TRUE</a> , the returned indices are unique.
...	Not used.

**Author(s)**

Henrik Bengtsson

**See Also**

[seqToIntervals\(\)](#).

**Examples**

```
## Not run: See example(seqToIntervals)
```

---

intToBin *Converts an integer to a binary/octal/hexadecimal number*

---

**Description**

Converts an integer to a binary/octal/hexadecimal number.

**Usage**

```
intToBin(x)  
intToOct(x)  
intToHex(x)
```

**Arguments**

`x` A **numeric** vector of integers to be converted.

**Details**

For `length(x) > 1`, the number of characters in each of returned elements is the same and driven by the `x` element that requires the highest number of character - all other elements are padded with zeros (or ones for negative values). This is why we for instance get `intToHex(15) == "f"` but `intToHex(15:16) == c("0f", "10")`.

The supported range for `intToHex()`, `intToOct()`, and `intToBin()` is that of **R** integers, i.e. `[-.Machine$integer.max, +.Machine$integer.max]` where `.Machine$integer.max` is  $2^{31} - 1$ . This limitation is there such that negative values can be converted too.

**Value**

Returns a **character** string of length `length(x)`. For coercions out of range, `NA_character_` is returned for such elements.

**Author(s)**

Henrik Bengtsson

---

`isAbsolutePath`      *Checks if this pathname is absolute*

---

**Description**

Checks if this pathname is absolute.

**Usage**

```
## Default S3 method:
isAbsolutePath(pathname, ...)
```

**Arguments**

`pathname` A **character** string of the pathname to be checked.  
`...` Not used.

**Value**

Returns a **TRUE** if the pathname is absolute, otherwise **FALSE**.

**Author(s)**

Henrik Bengtsson

---

isDirectory	<i>Checks if the file specification is a directory</i>
-------------	--

---

**Description**

Checks if the file specification is a directory.

**Usage**

```
## Default S3 method:  
isDirectory(pathname, ...)
```

**Arguments**

pathname	A <a href="#">character</a> string of the pathname to be checked.
...	Not used.

**Value**

Returns [TRUE](#) if the file specification is a directory, otherwise [FALSE](#) is returned.

**Symbolic links**

This function follows symbolic links (also on Windows) and returns a value based on the link target (rather than the link itself).

**Author(s)**

Henrik Bengtsson

**See Also**

To check if it is a file see [isFile\(\)](#). Internally [file.info\(\)](#) is used. See also [file\\_test](#).

---

isEof.connection	<i>Checks if the current file position for a connection is at the 'End of File'</i>
------------------	---

---

**Description**

Checks if the current file position for a connection is at the 'End of File'.

**Usage**

```
## S3 method for class 'connection'  
isEof(con, ...)
```

**Arguments**

con	A <a href="#">connection</a> .
...	Not used.

**Details**

Internally [seek\(\)](#) is used, which according to the R help is discouraged on Windows. However, after many years of large-scale testing on various Windows versions and file systems we have yet to experience issues with using [seek\(\)](#) on Windows.

**Value**

Returns a [logical](#).

**Author(s)**

Henrik Bengtsson

**See Also**

For more information see [connection](#).

---

isFile	<i>Checks if the file specification is a file</i>
--------	---

---

**Description**

Checks if the file specification is a file.

**Usage**

```
## Default S3 method:  
isFile(pathname, ...)
```

**Arguments**

pathname	A <a href="#">character</a> string of the pathname to be checked.
...	Not used.

**Value**

Returns [TRUE](#) if the file specification is a file, otherwise [FALSE](#) is returned.

**Symbolic links**

This function follows symbolic links (also on Windows) and returns a value based on the link target (rather than the link itself).



**Author(s)**

Henrik Bengtsson

**See Also**

To check if it is a directory see [isDirectory\(\)](#). Internally [file.info\(\)](#) is used. See also [file\\_test](#).

---

isOpen.character	<i>Checks if there is an open connection to a file</i>
------------------	--

---

**Description**

Checks if there is an open connection to a file.

**Usage**

```
## S3 method for class 'character'  
isOpen(pathname, rw=c("read", "write"), ...)
```

**Arguments**

pathname	An <a href="#">character vector</a> .
rw	A <a href="#">character vector</a> . If "read", a file is considered to be open if there exist an open connection that can read from that file. If "write", a file is considered to be open if there exist an open connection that can write to that file. Both these values may be specified.
...	Not used.

**Value**

Returns a [logical vector](#) indicating for each file whether there exists an open file [connection](#) or not.

**Author(s)**

Henrik Bengtsson

**See Also**

See `isOpen()` in [connections.showConnections\(\)](#).

---

isPackageInstalled      *Checks if a package is installed or not*

---

**Description**

Checks if a package is installed or not.

**Usage**

```
## Default S3 method:  
isPackageInstalled(package, ...)
```

**Arguments**

package      A [character vector](#) of package names.  
...            Not used.

**Value**

Returns a [logical vector](#).

**Author(s)**

Henrik Bengtsson

**See Also**

[isPackageLoaded\(\)](#).

---

isPackageLoaded      *Checks if a package is loaded or not*

---

**Description**

Checks if a package is loaded or not. Note that, contrary to [require\(\)](#), this function does not load the package if not loaded.

**Usage**

```
## Default S3 method:  
isPackageLoaded(package, version=NULL, ...)
```

**Arguments**

package	The name of the package.
version	A <a href="#">character</a> string specifying the version to test for. If <code>NULL</code> , any version is tested for.
...	Not used.

**Value**

Returns a [logical](#).

**Author(s)**

Henrik Bengtsson

**See Also**

To check if a package is installed or not, see [isPackageInstalled\(\)](#).

---

isReplicated	<i>Identifies all entries with replicated values</i>
--------------	--

---

**Description**

Identifies all entries with replicated values, that is, with values that exist more than once.

**Usage**

```
isReplicated(x, ...)
replicates(x, ...)
```

**Arguments**

x	A <a href="#">vector</a> of length K.
...	Additional arguments passed to <a href="#">duplicated()</a> .

**Details**

Let `reps <- isReplicated(x)`. Then it always holds that:

- `reps == rev(isReplicated(rev(x)))`
- `reps == duplicated(x) | duplicated(x, fromLast=TRUE)`
- `reps == !is.element(x, setdiff(x, unique(x[duplicated(x)])))`

**Value**

A [logical vector](#) of length K, where `TRUE` indicates that the value exists elsewhere, otherwise not.

**Author(s)**

Henrik Bengtsson

**See Also**Internally `duplicated()` is used. See also `isSingle()`.**Examples**

```

x <- c(1,1,2,3,4,2,1)
x <- base::letters[x]
print(x)

# Identify entries with replicated values
reps <- isReplicated(x)
print(x[reps])
stopifnot(x[reps] == replicates(x))

# Identify entries with unique values
print(x[!reps])
stopifnot(x[!reps] == singles(x))

# -----
# Validation
# -----
x <- c(1,1,2,3,4,2,1)
x <- base::letters[x]
reps <- isReplicated(x)

stopifnot(all(table(x[reps]) > 1))
stopifnot(all(table(x[!reps]) == 1))
stopifnot(all(reps == rev(isReplicated(rev(x)))))
stopifnot(all(reps == duplicated(x) | duplicated(x, fromLast=TRUE)))
stopifnot(all(reps == !is.element(x, setdiff(x, unique(x[duplicated(x)]))))))
stopifnot(all(sort(c(singles(x), replicates(x))) == sort(x)))

# -----
# Benchmarking singles()
# -----
set.seed(0xBEEF)
n <- 1e6
x <- sample(1:(n/2), size=n, replace=TRUE)
t <- system.time({
  s <- isSingle(x)
})
print(sum(s))

t0 <- system.time({
  s0 <- !(x %in% x[duplicated(x)]);

```

```
})  
print(t/t0)  
stopifnot(all(s == s0))
```

---

isSingle	<i>Identifies all entries that exists exactly once</i>
----------	--

---

### Description

Identifies all entries that exists exactly once.

### Usage

```
isSingle(x, ...)  
singles(x, ...)
```

### Arguments

x	A <a href="#">vector</a> of length K.
...	Additional arguments passed to <a href="#">isReplicated()</a> .

### Value

A [logical vector](#) of length K, indicating whether the value is unique or not.

### Author(s)

Henrik Bengtsson

### See Also

Internally [isReplicated\(\)](#) is used.

---

isUrl	<i>Checks if one or several pathnames is URLs</i>
-------	---

---

### Description

Checks if one or several pathnames is URLs.

### Usage

```
## Default S3 method:  
isUrl(pathname, ...)
```

**Arguments**

pathname      A [character vector](#).  
 ...            Not used.

**Value**

Returns a [logical vector](#) of either `TRUE` or `FALSE`.

**Author(s)**

Henrik Bengtsson

---

isZero	<i>Checks if a value is (close to) zero or not</i>
--------	--

---

**Description**

Checks if a value (or a vector of values) is (close to) zero or not where "close" means if the absolute value is less than `neps*eps`. *Note that  $x == 0$  will not work in all cases.*

By default `eps` is the smallest possible floating point value that can be represented by the running machine, i.e. `.Machine$double.eps` and `neps` is one. By changing `neps` it is easy to adjust how close to zero "close" means without having to know the machine precision (or remembering how to get it).

**Usage**

```
## Default S3 method:
isZero(x, neps=1, eps=.Machine$double.eps, ...)
```

**Arguments**

`x`            A [vector](#) of values.  
`eps`            The smallest possible floating point.  
`neps`          A scale factor of `eps` specifying how close to zero "close" means. If `eps` is the smallest value such that  $1 + eps \neq 1$ , i.e. `.Machine$double.eps`, `neps` must be greater or equal to one.  
 ...            Not used.

**Value**

Returns a [logical vector](#) indicating if the elements are zero or not.

**Author(s)**

Henrik Bengtsson

**See Also**

[all.equal\(\). Comparison. .Machine.](#)

**Examples**

```
x <- 0
print(x == 0)      # TRUE
print(isZero(x))  # TRUE

x <- 1
print(x == 0)      # FALSE
print(isZero(x))  # FALSE

x <- .Machine$double.eps
print(x == 0)      # FALSE
print(isZero(x))  # FALSE

x <- 0.9*.Machine$double.eps
print(x == 0)      # FALSE
print(isZero(x))  # TRUE

# From help(Comparisons)
x1 <- 0.5 - 0.3
x2 <- 0.3 - 0.1
print(x1 - x2)
print(x1 == x2)    # FALSE on most machines
print(identical(all.equal(x1, x2), TRUE)) # TRUE everywhere
print(isZero(x1-x2)) # TRUE everywhere
```

---

 Java

*Static class for Java related methods*


---

**Description**

Package: R.utils

**Class Java**

**Object**

~~|

~~+--Java

**Directly known subclasses:**

```
public static class Java
  extends Object
```

Static class that provides methods for reading and writing Java data types. Currently the following data types are supported: byte, short and int. R character strings can be written as UTF-8 formatted strings, which can be read by Java. Currently on Java String's that contain ASCII characters can be imported into R. The reason for this is that other characters are translated into non-eight bits data, e.g. 16- and 24-bits, which the readChar() method currently does not support.

Furthermore, the Java class defines some static constants describing the minimum and maximum value of some of the common Java data types: `BYTE.MIN`, `BYTE.MAX`, `SHORT.MIN`, `SHORT.MAX`, `INT.MIN`, `INT.MAX`, `LONG.MIN`, and `LONG.MAX`.

## Usage

```
Java()
```

## Fields and Methods

### Methods:

<code>asByte</code>	Converts a numeric to a Java byte.
<code>asInt</code>	Converts an numeric to a Java integer.
<code>asLong</code>	Converts a numeric to a Java long.
<code>asShort</code>	Converts a numeric to a Java short.
<code>readByte</code>	Reads a Java formatted byte (8 bits) from a connection.
<code>readInt</code>	Reads a Java formatted int (32 bits) from a connection.
<code>readShort</code>	Reads a Java formatted short (16 bits) from a connection.
<code>readUTF</code>	Reads a Java (UTF-8) formatted string from a connection.
<code>writeByte</code>	Writes a byte (8 bits) to a connection in Java format.
<code>writeInt</code>	Writes a integer (32 bits) to a connection in Java format.
<code>writeShort</code>	Writes a short (16 bits) to a connection in Java format.
<code>writeUTF</code>	Writes a string to a connection in Java format (UTF-8).

### Methods inherited from Object:

`$`, `$<-`, `[]`, `[[<-`, `as.character`, `attach`, `attachLocally`, `clearCache`, `clearLookupCache`, `clone`, `detach`, `equals`, `extend`, `finalize`, `getEnvironment`, `getFieldModifier`, `getFieldModifiers`, `getFields`, `getInstantiationTime`, `getStaticInstance`, `hasField`, `hashCode`, `ll`, `load`, `names`, `objectSize`, `print`, `save`

## Author(s)

Henrik Bengtsson

## Examples

```
pathname <- tempfile()

# Open the temporary file for writing
out <- file(pathname, open="wb")
b <- -128:127
```



```

Java$writeByte(out, b)
s <- -32768:32767
Java$writeShort(out, s)
i <- c(-2147483648, -2147483647, -1, 0, +1, 2147483646, 2147483647);
Java$writeInt(out, i)
str <- c("This R string was written (using the UTF-8 format) using",
        "the static methods of the Java class in the R.io package.")
str <- paste(str, collapse="\n")
Java$writeUTF(out, str)
close(out)

# Open the temporary file for reading
inn <- file(pathname, open="rb")

bfr <- Java$readByte(inn, n=length(b))
cat("Read ", length(bfr), " bytes.\n", sep="")
if (!identical(bfr, b))
  throw("Failed to read the same data that was written.")

bfr <- Java$readShort(inn, n=length(s))
cat("Read ", length(bfr), " shorts.\n", sep="")
if (!identical(bfr, s))
  throw("Failed to read the same data that was written.")

bfr <- Java$readInt(inn, n=length(i))
cat("Read ", length(bfr), " ints.\n", sep="")
if (!identical(bfr, i))
  throw("Failed to read the same data that was written.")

bfr <- Java$readUTF(inn)
cat("Read ", nchar(bfr), " UTF characters:\n", "'", bfr, "'\n", sep="")

close(inn)

file.remove(pathname)

```

---

lastModified

*Gets the time when the file was last modified*


---

### Description

Gets the time when the file was last modified. The time is returned as a POSIXct object.

### Usage

```

## Default S3 method:
lastModified(pathname, ...)

```

**Arguments**

pathname      A [character](#) string of the pathname to be checked.  
 ...            Not used.

**Value**

Returns POSIXct object specifying when the file was last modified. If the file does not exist or it is a directory, 0 is returned.

**Symbolic links**

This function follows symbolic links (also on Windows) and returns a value based on the link target (rather than the link itself).

**Author(s)**

Henrik Bengtsson

**See Also**

Internally [file.info\(\)](#) is used.

---

 LComments

*The LComments class*


---

**Description**

Package: R.utils

**Class LComments****Object**

```

~|
~+--SmartComments
~~~~~|
~~~~~+--VComments
~~~~~|
~~~~~+--LComments

```

**Directly known subclasses:**

```

public static class LComments
  extends VComments

```

The LComments class.

This class, is almost identical to the super class, except that the constructor has different defaults.

**Usage**

```
LComments(letter="L", verboseName="log", ...)
```

**Arguments**

letter	The smart letter.
verboseName	The name of the verbose object.
...	Not used.

**Fields and Methods****Methods:**

*No methods defined.*

**Methods inherited from VComments:**

convertComment, reset, validate

**Methods inherited from SmartComments:**

compile, convertComment, parse, reset, validate

**Methods inherited from Object:**

\$, \$<-, [], [[<-, as.character, attach, attachLocally, clearCache, clearLookupCache, clone, detach, equals, extend, finalize, getEnvironment, getFieldModifier, getFieldModifiers, getFields, getInstantiationTime, getStaticInstance, hasField, hashCode, ll, load, names, objectSize, print, save

**Author(s)**

Henrik Bengtsson

---

listDirectory	<i>Gets the file names in the directory</i>
---------------	---

---

**Description**

Gets the file names in the directory.

Contrary to `list.files()`, this method guarantees to work recursively. Moreover, when subdirectories are processed recursively, directory names are also returned.

**Usage**

```
## Default S3 method:  
listDirectory(path=".", pattern=NULL, recursive=FALSE, allNames=FALSE, fullNames=FALSE,  
...)
```

**Arguments**

path	A path to be listed.
pattern	A <a href="#">character</a> string of the filename pattern passed. See <a href="#">list.files()</a> for more details.
recursive	If <a href="#">TRUE</a> , subdirectories are recursively processed, and not if <a href="#">FALSE</a> . Alternatively, the maximum recursive depth can be specified as a non-negative <a href="#">numeric</a> , where <a href="#">FALSE</a> corresponds to 0L depth and <a href="#">TRUE</a> corresponds to +Inf depth.
allNames	If <a href="#">TRUE</a> , also files starting with a period are returned.
fullNames	If <a href="#">TRUE</a> , the full path names are returned.
...	Not used.

**Value**

Returns a [vector](#) of file names.

**Recursive searching**

Recursive searching of directory structure is done breath-first in a lexicographic order.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally [list.files\(\)](#) is used.

---

loadObject	<i>Method to load object from a file or a connection</i>
------------	--

---

**Description**

Method to load object from a file or a connection, which previously have been saved using [saveObject\(\)](#).

**Usage**

```
## Default S3 method:
loadObject(file, path=NULL, format=c("auto", "xdr", "rds"), ...)
```

**Arguments**

file	A filename or <a href="#">connection</a> to read the object from.
path	The path where the file exists.
format	File format.
...	Not used.

**Details**

The main difference from this method and `load()` in the **base** package, is that this one returns the object read rather than storing it in the global environment by its default name. This makes it possible to load objects back using any variable name.

**Value**

Returns the saved object.

**Author(s)**

Henrik Bengtsson

**See Also**

`saveObject()` to save an object to file. Internally `load()` is used. See also `loadToEnv()`. See also `saveRDS()`.

---

mapToIntervals.numeric

*Maps values to intervals*

---

**Description**

Maps values to intervals by returning an index **vector** specifying the (first) interval that each value maps to, if any.

**Usage**

```
## S3 method for class 'numeric'
mapToIntervals(x, intervals, includeLower=TRUE, includeUpper=TRUE, ...)
```

**Arguments**

<code>x</code>	A <b>numeric vector</b> of K values to be matched.
<code>intervals</code>	The N intervals to be matched against. If an Nx2 <b>numeric matrix</b> , the first column should be the lower bounds and the second column the upper bounds of each interval. If a <b>numeric vector</b> of length 2N, each consecutive pair should be the lower and upper bounds of an interval.
<code>includeLower, includeUpper</code>	If <b>TRUE</b> , the lower (upper) bound of <i>each</i> interval is included in the test, otherwise not.
<code>...</code>	Not used.

**Value**

Returns an **integer vector** of length K. Values that do not map to any interval have return value **NA**.

**Author(s)**

Henrik Bengtsson

**See Also**[inAnyInterval\(\)](#). [match\(\)](#). [findInterval\(\)](#). [cut\(\)](#).

---

`mergeIntervals.numeric`*Merges intervals*

---

**Description**

Merges intervals by returning an index [vector](#) specifying the (first) interval that each value maps to, if any.

**Usage**

```
## S3 method for class 'numeric'
mergeIntervals(intervals, ...)
```

**Arguments**

<code>intervals</code>	The N intervals to be merged. If an Nx2 <a href="#">numeric matrix</a> , the first column should be the lower bounds and the second column the upper bounds of each interval. If a <a href="#">numeric vector</a> of length 2N, each consecutive pair should be the lower and upper bounds of an interval.
<code>...</code>	Not used.

**Details**

The upper and lower bounds are considered to be inclusive, that is, all intervals are interpreted to be of form [a,b]. There is currently no way to specify intervals with open bounds, e.g. (a,b).

Furthermore, the bounds are currently treated as real values. For instance, merging [0,1] and [2,3] will return the same intervals. Note, if integer intervals were treated specially, we would merge these intervals to integer interval [0,3] == {0,1,2,3}.

**Value**

Returns a [matrix](#) (or a [vector](#)) of M intervals, where  $M \leq N$ . The intervals are ordered by their lower bounds. The @mode of the returned intervals is the same as the mode of the input intervals.

**Author(s)**

Henrik Bengtsson

**See Also**

[inAnyInterval\(\). match\(\)](#).

---

mkdirs	<i>Creates a directory including any necessary but nonexistent parent directories</i>
--------	---

---

**Description**

Creates a directory including any necessary but nonexistent parent directories.

**Usage**

```
## Default S3 method:  
mkdirs(pathname, mustWork=FALSE, maxTries=5L, ...)
```

**Arguments**

pathname	A <a href="#">character</a> string of the pathname to be checked.
mustWork	If <a href="#">TRUE</a> and the directory does not already exist or is failed to be created, an error is thrown, otherwise not.
maxTries	A positive <a href="#">integer</a> specifying how many times the method should try to create a missing directory before giving up.
...	Not used.

**Value**

Returns [TRUE](#) if the directory was successfully created, otherwise [FALSE](#). Note that if the directory already exists, [FALSE](#) is returned.

**Slow file systems**

On very rare occasions, we have observed on a large shared file system that if one tests for the existence of a directory immediately after creating it with [dir.create\(\)](#), it may appear not to be created. We believe this is due to the fact that there is a short delay between creating a directory and that information being fully propagated on the file system. To minimize the risk for such false assertions on "slow" file systems, this method tries to create a missing directory multiple times (argument `maxTries`) (while waiting a short period of time between each round) before giving up.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally [dir.create\(\)](#) is used.

---

`mout`*Miscellaneous functions for outputting via message()*

---

## Description

Miscellaneous functions for outputting via `message()`. These "m\*" methods work analogously to their corresponding "\*" methods `print()`, `cat()`, `show`, `str`, and `printf()` but uses `message()` to output the content, which in turn outputs to standard error. The `mout()` method can be used for all other output methods, e.g. `mout(write(x, file=stdout()))`.

## Usage

```
mout(..., appendLF=FALSE)
```

## Arguments

... Arguments passed to the underlying output method.

appendLF A [logical](#) specifying whether to append a newline at the end or not.

## Value

Returns what the `message()` returns.

## Author(s)

Henrik Bengtsson

## Examples

```
print(letters[1:8])
mprint(letters[1:8])

cat(c(letters[1:8], "\n"))
mcat(c(letters[1:8], "\n"))

str(letters[1:8])
mstr(letters[1:8])

printf("x=%d\n", 1:3)
mprintf("x=%d\n", 1:3)
```



---

mpager	<i>A \"pager\" function that outputs to standard error</i>
--------	--

---

**Description**

A \"pager\" function that outputs to standard error and is compatible with [file.show\(\)](#).

**Usage**

```
mpager(files, header=NULL, title="R Information", delete.file=FALSE)
```

**Arguments**

files	A <a href="#">character vector</a> of K pathnames.
header	A <a href="#">character vector</a> of K headers.
title	A <a href="#">character</a> string.
delete.file	If <a href="#">TRUE</a> , the files are deleted after displayed, otherwise not.

**Value**

Returns nothing.

**Author(s)**

Henrik Bengtsson

**See Also**

[file.show\(\)](#) and argument pager.

---

nullfile	<i>Gets the pathname or a connection to the NULL device on the current platform</i>
----------	---

---

**Description**

Gets the pathname or a connection to the NULL device on the current platform.

**Usage**

```
nullfile()  
nullcon()
```

**Value**

nullfile() returns a [character](#) string, which is `"/dev/null"` except on Windows where it is `"nul:"`. nullcon() returns a *newly opened* (binary) [connection](#) to the NULL device - make sure to close it when no longer needed.

**Author(s)**

Henrik Bengtsson

**See Also**

In R ( $\geq 3.6.0$ ), there exists `base::nullfile()`, which is identical to `R.utils::nullfile()`.

---

NullVerbose

*A Verbose class ignoring everything*

---

**Description**

Package: R.utils

**Class NullVerbose**

**Object**

```
~~|
```

```
~~+---Verbose
```

```
~~~~~|
```

```
~~~~~+---NullVerbose
```

**Directly known subclasses:**

```
public static class NullVerbose
```

```
extends Verbose
```

A Verbose class ignoring everything.

**Usage**

```
NullVerbose(...)
```

**Arguments**

```
... Ignored.
```

**Fields and Methods**

**Methods:**

cat	-
enter	-
evaluate	-
exit	-
header	-
isOn	Checks if the output is on.
isVisible	Checks if a certain verbose level will be shown or not.
newline	-
print	-
printf	-
ruler	-
str	-
summary	-
writeRaw	All output methods.

**Methods inherited from Verbose:**

as.character, as.double, as.logical, capture, cat, enter, enterf, equals, evaluate, exit, getThreshold, getTimestampFormat, header, isOn, isVisible, less, more, newline, off, on, popState, print, printf, pushState, ruler, setDefaultLevel, setThreshold, setTimestampFormat, str, summary, timestamp, timestampOff, timestampOn, warnings, writeRaw

**Methods inherited from Object:**

\$, \$<-, [[, [[<-, as.character, attach, attachLocally, clearCache, clearLookupCache, clone, detach, equals, extend, finalize, getEnvironment, getFieldModifier, getFieldModifiers, getFields, getInstantiationTime, getStaticInstance, hasField, hashCode, ll, load, names, objectSize, print, save

**Author(s)**

Henrik Bengtsson

**Examples**

```
verbose <- Verbose()
cat(verbose, "A verbose messages")

verbose <- NullVerbose()
cat(verbose, "A verbose messages") # Ignored
```

---

onGarbageCollect	<i>Registers a function to be called when the R garbage collector is (detected to be) running</i>
------------------	---

---

**Description**

Registers a function to be called when the R garbage collector is (detected to be) running.

**Usage**

```
## Default S3 method:
onGarbageCollect(fcn, action=c("prepend", "append", "replace"), ...)
```

**Arguments**

fcn	A <a href="#">function</a> to be called without argument.
action	A <a href="#">character</a> string specifying how the hook function is added to list of hooks.
...	Not used.

**Value**

Returns (invisibly) the hooks successfully called.

**Author(s)**

Henrik Bengtsson

**Examples**

```
## Not run:
onGarbageCollect(function(...) {
  message("The R garbage collector is running!")
})

## End(Not run)
```

---

onSessionExit	<i>Registers a function to be called when the R session finishes</i>
---------------	--

---

**Description**

Registers a function to be called when the R session finishes.

**Usage**

```
## Default S3 method:
onSessionExit(fcn, action=c("prepend", "append", "replace"), ...)
```

**Arguments**

fcn	A <a href="#">function</a> to be called without argument.
action	A <a href="#">character</a> string specifying how the hook function is added to list of hooks.
...	Not used.

**Details**

Functions registered this way are called when `finalizeSession()` is called. Moreover, when this package is loaded, the `.Last()` function is modified such that `finalizeSession()` is called. However, note that `.Last()` is *not* guaranteed to be called when the R session finished. For instance, the user may quit R by calling `quit(callLast=FALSE)`. Moreover, when R is run in batch mode, `.Last()` is never called.

**Value**

Returns (invisibly) the hooks successfully called.

**Author(s)**

Henrik Bengtsson

**See Also**

`.Last()`, `finalizeSession()`.

**Examples**

```
## Not run:
  onSessionExit(function(...) {
    message("Bye bye world!")
  })

  quit()

## End(Not run)
```

---

Options

*The Options class*

---

**Description**

Package: R.utils

**Class Options**

**Object**

~~|

~~+--Options

**Directly known subclasses:**

[Settings](#)

public static class **Options**  
 extends [Object](#)

A class to set and get either options stored in a [list](#) tree structure.

Each option has a pathname. The format of a pathname is similar to a (Unix) filesystem pathname, e.g. "graphics/cex". See examples for more details.

### Usage

```
Options(options=list(), ...)
```

### Arguments

<code>options</code>	A tree <a href="#">list</a> structure of options.
<code>...</code>	Not used.

### Details

Note, this class and its methods do *not* operate on the global options structure defined in R ([options](#)).

### Value

The constructor returns an Options object.

### Fields and Methods

#### Methods:

<a href="#">as.character</a>	Returns a character string version of this object.
<a href="#">as.list</a>	Gets a list representation of the options.
<a href="#">equals</a>	Checks if this object is equal to another Options object.
<a href="#">getLeaves</a>	Gets all (non-list) options in a flat list.
<a href="#">getOption</a>	Gets an option.
<a href="#">hasOption</a>	Checks if an option exists.
<a href="#">names</a>	Gets the full pathname of all (non-list) options.
<a href="#">nbrOfOptions</a>	Gets the number of options set.
<a href="#">setOption</a>	Sets an option.
<a href="#">str</a>	Prints the structure of the options.

#### Methods inherited from Object:

`$`, `$<-`, `[[`, `[[<-`, `as.character`, `attach`, `attachLocally`, `clearCache`, `clearLookupCache`, `clone`, `detach`, `equals`, `extend`, `finalize`, `getEnvironment`, `getFieldModifier`, `getFieldModifiers`, `getFields`, `getInstantiationTime`, `getStaticInstance`, `hasField`, `hashCode`, `ll`, `load`, `names`, `objectSize`, `print`, `save`

### Author(s)

Henrik Bengtsson

**Examples**

```

local <- Options()

# Query a missing option
cex <- getOption(local, "graphics/cex")
cat("graphics/cex =", cex, "\n") # Returns NULL

# Query a missing option with default value
cex <- getOption(local, "graphics/cex", defaultValue=1)
cat("graphics/cex =", cex, "\n") # Returns NULL

# Set option and get previous value
oldCex <- setOption(local, "graphics/cex", 2)
cat("previous graphics/cex =", oldCex, "\n") # Returns NULL

# Set option again and get previous value
oldCex <- setOption(local, "graphics/cex", 3)
cat("previous graphics/cex =", oldCex, "\n") # Returns 2

# Query a missing option with default value, which is ignored
cex <- getOption(local, "graphics/cex", defaultValue=1)
cat("graphics/cex =", cex, "\n") # Returns 3

# Query multiple options with multiple default values
multi <- getOption(local, c("graphics/cex", "graphics/pch"), c(1,2))
print(multi);

# Check existence of multiple options
has <- hasOption(local, c("graphics/cex", "graphics/pch"))
print(has);

# Get a subtree of options
graphics <- getOption(local, "graphics")
print(graphics)

# Get the complete tree of options
all <- getOption(local)
print(all)

```

---

patchCode

*Patches installed and loaded packages and more*


---

**Description**

Patches installed and loaded packages and more.

**Usage**

```
## Default S3 method:
patchCode(paths=NULL, recursive=TRUE, suppressWarnings=TRUE,
          knownExtensions=c("R", "r", "S", "s"), verbose=FALSE, ...)
```

**Arguments**

paths	The path to the directory (and subdirectories) which contains source code that will patch loaded packages. If <code>NULL</code> , the patch path is given by the option <code>R_PATCHES</code> , If the latter is not set, the system environment with the same name is used. If neither is given, then <code>~/R-patches/</code> is used.
recursive	If <code>TRUE</code> , source code in subdirectories will also get loaded.
suppressWarnings	If <code>TRUE</code> , <code>warnings</code> will be suppressed, otherwise not.
knownExtensions	A <code>character vector</code> of filename extensions used to identify source code files. All other files are ignored.
verbose	If <code>TRUE</code> , extra information is printed while patching, otherwise not.
...	Not used.

**Details**

The method will look for source code files (recursively or not) that match known filename extensions. Each found source code file is then `source()`d.

If the search is recursive, subdirectories are entered if and only if either (1) the name of the subdirectory is the same as a *loaded* (and installed) package, or (2) if there is no installed package with that name. The latter allows common code to be organized in directories although it is still not assigned to packages.

Each of the directories given by argument `paths` will be processed one by one. This makes it possible to have more than one file tree containing patches.

To set an options, see `options()`. To set a system environment, see `Sys.setenv()`. The character `;` is interpreted as a separator. Due to incompatibility with Windows pathnames, `:` is *not* a valid separator.

**Value**

Returns (invisibly) the number of files sourced.

**Author(s)**

Henrik Bengtsson

**See Also**

`source()`. `library()`.



**Examples**

```
## Not run:
# Patch all source code files in the current directory
patchCode(".")

# Patch all source code files in R_PATCHES
options("R_PATCHES"=~~/R-patches/")
# alternatively, Sys.setenv("R_PATCHES"=~~/R-patches/")
patchCode()

## End(Not run)
```

---

popBackupFile	<i>Drops a backup suffix from the backup pathname</i>
---------------	---

---

**Description**

Drops a backup suffix from the backup pathname and, by default, restores an existing backup file accordingly by renaming it.

**Usage**

```
## Default S3 method:
popBackupFile(filename, path=NULL, suffix=".bak", isFile=TRUE,
  onMissing=c("ignore", "error"), drop=TRUE, ..., verbose=FALSE)
```

**Arguments**

filename	The filename of the backup file.
path	The path of the file.
suffix	The suffix of the filename to be dropped.
isFile	If <b>TRUE</b> , the backup file must exist and will be renamed. If <b>FALSE</b> , it is only the pathname string that will be modified. For details, see below.
onMissing	A <b>character</b> string specifying what to do if the backup file does not exist.
drop	If <b>TRUE</b> , the backup file will be dropped in case the original file already exists or was successfully restored.
...	Not used.
verbose	A <b>logical</b> or <b>Verbose</b> .

**Value**

Returns the pathname with the backup suffix dropped.

**Author(s)**

Henrik Bengtsson

**See Also**

See [pushBackupFile\(\)](#) for more details and an example.

---

popTemporaryFile      *Drops a temporary suffix from the temporary pathname*

---

**Description**

Drops a temporary suffix from the temporary pathname and, by default, renames an existing temporary file accordingly.

**Usage**

```
## Default S3 method:
popTemporaryFile(filename, path=NULL, suffix=".tmp", isFile=TRUE, ..., verbose=FALSE)
```

**Arguments**

filename	The filename of the temporary file.
path	The path of the temporary file.
suffix	The suffix of the temporary filename to be dropped.
isFile	If <b>TRUE</b> , the temporary file must exist and will be renamed. If <b>FALSE</b> , it is only the pathname string that will be modified. For details, see below.
...	Not used.
verbose	A <a href="#">logical</a> or <a href="#">Verbose</a> .

**Details**

If `isFile` is **FALSE**, the pathname where the suffix of the temporary pathname has been dropped is returned. If `isFile` is **TRUE**, the temporary file is renamed. Then, if the temporary file does not exist or it was not successfully renamed, an exception is thrown.

**Value**

Returns the pathname with the temporary suffix dropped.

**Author(s)**

Henrik Bengtsson

**See Also**

See [pushTemporaryFile\(\)](#) for more details and an example.

---

printf	<i>C-style formatted output</i>
--------	---------------------------------

---

**Description**

C-style formatted output.

**Usage**

```
## Default S3 method:  
printf(fmt, ..., sep="", file="")
```

**Arguments**

fmt	A <a href="#">character</a> vector of format strings. See same argument for <a href="#">sprintf()</a> .
...	Additional arguments <a href="#">sprintf()</a> .
sep	A <a href="#">character vector</a> of strings to append after each element.
file	A <a href="#">connection</a> , or a <a href="#">character</a> of a file to print to. See same argument for <a href="#">cat()</a> .

**Value**

Returns nothing.

**Author(s)**

Henrik Bengtsson

**See Also**

For C-style formatting of [character](#) strings, see [sprintf\(\)](#).

**Examples**

```
cat("Hello world\n")  
printf("Hello world\n")  
  
x <- 1.23  
cat(sprintf("x=%.2f\n", x))  
printf("x=%.2f\n", x)  
  
y <- 4.56  
cat(sprintf(c("x=%.2f\n", "y=%.2f\n"), c(x,y)), sep="")  
printf(c("x=%.2f\n", "y=%.2f\n"), c(x,y))
```

---

 ProgressBar

*Provides text based counting progress bar*


---

## Description

Package: R.utils

### Class ProgressBar

[Object](#)

~~|

~~+--ProgressBar

### Directly known subclasses:

[FileProgressBar](#)

public static class **ProgressBar**

extends [Object](#)

## Usage

```
ProgressBar(max=100, ticks=10, stepLength=1, newlineWhenDone=TRUE)
```

## Arguments

max	The maximum number of steps.
ticks	Put visual "ticks" every ticks step.
stepLength	The default length for each increase.
newlineWhenDone	If <b>TRUE</b> , a newline is outputted when bar is updated, when done, otherwise not.

## Fields and Methods

### Methods:

<a href="#">as.character</a>	Gets a string description of the progress bar.
<a href="#">getBarString</a>	Gets the progress bar string to be displayed.
<a href="#">increase</a>	Increases (steps) progress bar.
<a href="#">isDone</a>	Checks if progress bar is completed.
<a href="#">reset</a>	Reset progress bar.
<a href="#">setMaxValue</a>	Sets maximum value.
<a href="#">setProgress</a>	Sets current progress.
<a href="#">setStepLength</a>	Sets default step length.
<a href="#">setTicks</a>	Sets values for which ticks should be visible.
<a href="#">setValue</a>	Sets current value.

`update` Updates progress bar.

**Methods inherited from Object:**

\$, \$<-, [[, [[<-, as.character, attach, attachLocally, clearCache, clearLookupCache, clone, detach, equals, extend, finalize, getEnvironment, getFieldModifier, getFieldModifiers, getFields, getInstantiationTime, getStaticInstance, hasField, hashCode, ll, load, names, objectSize, print, save

**Author(s)**

Henrik Bengtsson

**Examples**

```
# A progress bar with default step length one.
pb <- ProgressBar(max=42)
reset(pb)
while (!isDone(pb)) {
  x <- rnorm(3e4)
  increase(pb)
  # Emulate a slow process
  if (interactive()) Sys.sleep(0.02)
}
cat("\n")

# A "faster" progress bar with default step length 1.4.
pb <- ProgressBar(max=42, stepLength=1.4)
reset(pb)
while (!isDone(pb)) {
  x <- rnorm(3e4)
  increase(pb)
  # Emulate a slow process
  if (interactive()) Sys.sleep(0.02)
}

cat("\n")
```

---

pushBackupFile

*Appends a backup suffix to the pathname*

---

**Description**

Appends a backup suffix to the pathname and, optionally, renames an existing file accordingly.

In combination with `popBackupFile()`, this method is useful for creating a backup of a file and restoring it.

**Usage**

```
## Default S3 method:
pushBackupFile(filename, path=NULL, suffix=".bak", isFile=TRUE,
  onMissing=c("ignore", "error"), copy=FALSE, overwrite=TRUE, ..., verbose=FALSE)
```

**Arguments**

filename	The filename of the file to backup.
path	The path of the file.
suffix	The suffix to be appended.
isFile	If <b>TRUE</b> , the file must exist and will be renamed on the file system. If <b>FALSE</b> , it is only the pathname string that will be modified. For details, see below.
onMissing	A <b>character</b> string specifying what to do if the file does not exist.
copy	If <b>TRUE</b> , an existing original file remains after creating the backup copy, otherwise it is dropped.
overwrite	If <b>TRUE</b> , any existing backup files are overwritten, otherwise an exception is thrown.
...	Not used.
verbose	A <b>logical</b> or <b>Verbose</b> .

**Value**

Returns the pathname with the suffix appended.

**Author(s)**

Henrik Bengtsson

**See Also**

[popBackupFile\(\)](#).

**Examples**

```
# Create a file
pathname <- "foobar.txt";
cat(file=pathname, "File v1\n");

# -----
# (a) Backup and restore a file
# -----
# Turn it into a backup file
pathnameB <- pushBackupFile(pathname, verbose=TRUE);
print(pathnameB);

# Restore main file from backup
pathnameR <- popBackupFile(pathnameB, verbose=TRUE);
```

```

print(pathnameR);

# -----
# (b) Backup, create a new file and frop backup file
# -----
# Turn it into a backup file
pathnameB <- pushBackupFile(pathname, verbose=TRUE);
print(pathnameB);

# Create a new file
cat(file=pathname, "File v2\n");

# Drop backup because a new main file was successfully created
pathnameR <- popBackupFile(pathnameB, verbose=TRUE);
print(pathnameR);

```

---

pushTemporaryFile      *Appends a temporary suffix to the pathname*

---

## Description

Appends a temporary suffix to the pathname and, optionally, renames an existing file accordingly.

In combination with [popTemporaryFile\(\)](#), this method is useful for creating a file/writing data to file *atomically*, by first writing to a temporary file which is then renamed. If for some reason the generation of the file was interrupted, for instance by a user interrupt or a power failure, then it is only the temporary file that is incomplete.

## Usage

```

## Default S3 method:
pushTemporaryFile(filename, path=NULL, suffix=".tmp", isFile=FALSE, ..., verbose=FALSE)

```

## Arguments

filename	The filename of the file.
path	The path of the file.
suffix	The suffix to be appended.
isFile	If <b>TRUE</b> , the file must exist and will be renamed on the file system. If <b>FALSE</b> , it is only the pathname string that will be modified. For details, see below.
...	Not used.
verbose	A <a href="#">logical</a> or <a href="#">Verbose</a> .

**Details**

If `isFile` is `FALSE`, the pathname where the suffix of the temporary pathname has been added is returned. If `isFile` is `TRUE`, the file is also renamed. Then, if the file does not exist or it was not successfully renamed, an exception is thrown.

**Value**

Returns the pathname with the suffix appended.

**Author(s)**

Henrik Bengtsson

**See Also**

[popTemporaryFile\(\)](#).

**Examples**

```
createAtomically <- function(pathname, ...) {
  cat("Pathname: ", pathname, "\n", sep="");

  # Generate a file atomically, i.e. the file will either be
  # complete or not created at all. If interrupted while
  # writing, only a temporary file will exist/remain.
  pathnameT <- pushTemporaryFile(pathname);
  cat("Temporary pathname: ", pathnameT, "\n", sep="");

  cat(file=pathnameT, "This file was created atomically:\n");
  for (kk in 1:10) {
    cat(file=pathnameT, kk, "\n", append=TRUE);
    # Emulate a slow process
    if (interactive()) Sys.sleep(0.1)
  }
  cat(file=pathnameT, "END OF FILE\n", append=TRUE);

  # Rename the temporary file
  pathname <- popTemporaryFile(pathnameT);

  pathname;
} # createAtomically()

pathname <- tempfile();

tryCatch({
  # Try to interrupt the process while writing...
  pathname <- createAtomically(pathname);
}, interrupt=function(intr) {
  str(intr);
})
```



```
# ...and this will throw an exception
bfr <- readLines(pathname);
cat(bfr, sep="\n");
```

---

queryRCmdCheck	<i>Gets the on R CMD check if the current R session was launched by it</i>
----------------	--

---

### Description

Gets the on R CMD check if the current R session was launched by it.

### Usage

```
queryRCmdCheck(...)
```

### Arguments

... Not used.

### Value

Returns `character` string "checkingTests" if 'R CMD check' runs one of the package tests, and "checkingExamples" if it runs one of the package examples. If the current R session was not launched by 'R CMD check', then "notRunning" is returned.

### Limitations

This function only works if the working directory has not been changed.

### Author(s)

Henrik Bengtsson

### Examples

```
status <- queryRCmdCheck()
if (status != "notRunning") {
  cat("The current R session was launched by R CMD check. Status: ", status, "\n")
} else {
  cat("The current R session was not launched by R CMD check.\n")
}

# Display how R was launched
print(base::commandArgs())

# Display loaded packages etc.
print(search())
```

```
# Display current working directory
print(getwd())
```

---

readBinFragments	<i>Reads binary data from disjoint sections of a connection or a file</i>
------------------	---

---

## Description

Reads binary data from disjoint sections of a connection or a file.

## Usage

```
## Default S3 method:
readBinFragments(con, what, idxs=1, origin=c("current", "start"), size=NA, ...,
  verbose=FALSE)
```

## Arguments

con	A <a href="#">connection</a> or the pathname of an existing file.
what	A <a href="#">character</a> string or an object specifying the the data type ( <a href="#">mode()</a> ) to be read.
idxs	A <a href="#">vector</a> of (non-duplicated) indices or a Nx2 <a href="#">matrix</a> of N from-to index intervals specifying the elements to be read. Positions are either relative to the start or the current location of the file/connection as given by argument origin.
origin	A <a href="#">character</a> string specify whether the indices in argument idxs are relative to the "start" or the "current" position of the file/connection.
size	The size of the data type to be read. If <a href="#">NA</a> , the natural size of the data type is used.
...	Additional arguments passed to <a href="#">readBin()</a> .
verbose	A <a href="#">logical</a> or a <a href="#">Verbose</a> object.

## Value

Returns a [vector](#) of the requested [mode\(\)](#).

## Author(s)

Henrik Bengtsson

## See Also

[writeBinFragments\(\)](#).

**Examples**

```

# -----
# Create a data file
# -----
data <- 1:255
size <- 2
pathname <- tempfile("exampleReadBinFragments")
writeBin(con=pathname, data, size=size)

# -----
# Read and write using index vectors
# -----
cat("Read file...\n")
# Read every 16:th byte in the file
idxs <- seq(from=1, to=255, by=16)
x <- readBinFragments(pathname, what="integer", size=size, signed=FALSE, idxs=idxs)
stopifnot(identical(x, data[idxs]))
print(x)
# Read every 16:th byte in a connection starting with the 6th.
idxs <- idxs + 5L;
x <- readBinFragments(pathname, what="integer", size=size, signed=FALSE, idxs=idxs)
stopifnot(identical(x, data[idxs]))
print(x)
cat("Read file...done\n")

cat("Write file...\n")
# Update every 16:th byte in the file
idxs <- seq(from=1, to=255, by=16)
x0 <- data[idxs]
writeBinFragments(pathname, idxs=idxs, rev(x0), size=size)
x <- readBinFragments(pathname, what="integer", size=size, signed=FALSE, idxs=idxs)
print(x)
stopifnot(identical(rev(x0), x))

# Update every 16:th byte in the file
idxs <- seq(from=1, to=255, by=16)
writeBinFragments(pathname, idxs=idxs, rev(x), size=size)
x <- readBinFragments(pathname, what="integer", size=size, signed=FALSE, idxs=idxs)
print(x)
stopifnot(identical(x0, x))

# Assert everything is as expected
# Read the complete file
x <- readBin(pathname, what="integer", size=size, signed=FALSE, n=length(data))
stopifnot(identical(x, data))
cat("Write file...done\n")

# -----
# Ditto but via a connection
# -----

```

```

cat("Read connection...\n")
# Read every 16:th byte in a connection
con <- file(pathname, open="rb")
idxs <- seq(from=1, to=255, by=16)
x <- readBinFragments(con, what="integer", size=size, signed=FALSE, idxs=idxs)
stopifnot(identical(x, data[idxs]))
print(x)

# Read every 16:th byte in a connection starting with the 6th.
idxs <- idxs + 5L;
x <- readBinFragments(con, what="integer", size=size, signed=FALSE, idxs=idxs, origin="start")
stopifnot(identical(x, data[idxs]))
print(x)
close(con)
cat("Read connection...done\n")

# Update every 16:th byte in a connection
cat("Write connection...\n")
con <- file(pathname, open="r+b")
idxs <- seq(from=1, to=255, by=16)
x0 <- data[idxs]
writeBinFragments(pathname, idxs=idxs, rev(x0), size=size)
x <- readBinFragments(pathname, what="integer", size=size, signed=FALSE, idxs=idxs)
print(x)
stopifnot(identical(rev(x0), x))

# Update every 16:th byte in the file
idxs <- seq(from=1, to=255, by=16)
writeBinFragments(pathname, idxs=idxs, rev(x), size=size)
x <- readBinFragments(pathname, what="integer", size=size, signed=FALSE, idxs=idxs, origin="start")
print(x)
stopifnot(identical(x0, x))

close(con)

# Assert everything is as expected
# Read the complete file
x <- readBin(pathname, what="integer", size=size, signed=FALSE, n=length(data))
stopifnot(identical(x, data))
cat("Write connection...done\n")

# -----
# Clean up
# -----
file.remove(pathname)

```

**Description**

Reads one or more Rd help files in a certain format.

**Usage**

```
## Default S3 method:
readRdHelp(..., format=c("text", "html", "latex", "rd"), drop=TRUE)
```

**Arguments**

... Arguments passed to [help](#).

format A [character](#) string specifying the return type.

drop If [FALSE](#) or more than one help entry is found, the result is returned as a [list](#).

**Value**

Returns a [list](#) of [character](#) strings or a single [character](#) string.

**Author(s)**

Henrik Bengtsson

---

readTable	<i>Reads a file in table format</i>
-----------	-------------------------------------

---

**Description**

Reads a file in table format and creates a data frame from it, with cases corresponding to lines and variables to fields in the file.

*WARNING: This method is very much in an alpha stage. Expect it to change.*

This method is an extension to the default [read.table](#) function in R. It is possible to specify a column name to column class map such that the column classes are automatically assigned from the column header in the file.

In addition, it is possible to read any subset of rows. The method is optimized such that only columns and rows that are of interest are parsed and read into R's memory. This minimizes memory usage at the same time as it speeds up the reading.

**Usage**

```
## Default S3 method:
readTable(file, colClasses=NULL, isPatterns=FALSE, defColClass=NA, header=FALSE, skip=0,
  nrows=-1, rows=NULL, col.names=NULL, check.names=FALSE, path=NULL, ...,
  stripQuotes=TRUE, method=c("readLines", "intervals"), verbose=FALSE)
```

**Arguments**

file	A <a href="#">connection</a> or a filename. If a filename, the path specified by path is added to the front of the filename. Unopened files are opened and closed at the end.
colClasses	Either a named or an unnamed <a href="#">character vector</a> . If unnamed, it specified the column classes just as used by <a href="#">read.table</a> . If it is a named vector, names(colClasses) are used to match the column names read (this requires that header=TRUE) and the column classes are set to the corresponding values.
isPatterns	If <a href="#">TRUE</a> , the matching of names(colClasses) to the read column names is done by regular expressions matching.
defColClass	If the column class map specified by a named colClasses argument does not match some of the read column names, the column class is by default set to this class. The default is to read the columns in an "as is" way.
header	If <a href="#">TRUE</a> , column names are read from the file.
skip	The number of lines (commented or non-commented) to skip before trying to read the header or alternatively the data table.
nrows	The number of rows to read of the data table. Ignored if rows is specified.
rows	An row index <a href="#">vector</a> specifying which rows of the table to read, e.g. row one is the row following the header. Non-existing rows are ignored. Note that rows are returned in the same order they are requested and duplicated rows are also returned.
col.names	Same as in <a href="#">read.table()</a> .
check.names	Same as in <a href="#">read.table()</a> , but default value is <a href="#">FALSE</a> here.
path	If file is a filename, this path is added to it, otherwise ignored.
...	Arguments passed to <a href="#">read.table</a> used internally.
stripQuotes	If <a href="#">TRUE</a> , quotes are stripped from values before being parse. This argument is only effective when method=="readLines".
method	If "readLines", (readLines()) is used internally to first only read rows of interest, which is then passed to <a href="#">read.table()</a> . If "intervals", contiguous intervals are first identified in the rows of interest. These intervals are the read one by one using <a href="#">read.table()</a> . The latter methods is faster and especially more memory efficient if the intervals are not too many, where as the former is preferred if many "scattered" rows are to be read.
verbose	A <a href="#">logical</a> or a <a href="#">Verbose</a> object.

**Value**

Returns a [data.frame](#).

**Author(s)**

Henrik Bengtsson

**See Also**

[readTableIndex\(\)](#). [read.table.colClasses\(\)](#).

---

readTableIndex	<i>Reads a single column from file in table format</i>
----------------	--

---

### Description

Reads a single column from file in table format, which can then be used as a index-to-row (look-up) map for fast access to a subset of rows using `readTable()`.

### Usage

```
## Default S3 method:  
readTableIndex(..., indexColumn=1, colClass="character", verbose=FALSE)
```

### Arguments

<code>indexColumn</code>	An single <a href="#">integer</a> of the index column.
<code>colClass</code>	A single <a href="#">character</a> specifying the class of the index column.
<code>...</code>	Arguments passed to <code>readTable()</code> used internally.
<code>verbose</code>	A <a href="#">logical</a> or a <a href="#">Verbose</a> object.

### Value

Returns a [vector](#).

### Author(s)

Henrik Bengtsson

### See Also

[readTable\(\)](#).

### Examples

```
## Not run:  
# File containing data table to be access many times  
filename <- "somefile.txt"  
  
# Create a look-up index  
index <- readTableIndex(filename)  
  
# Keys of interest  
keys <- c("foo", "bar", "wah")  
  
# Read only those keys and do it fast  
df <- readTable(filename, rows=match(keys, index))  
  
## End(Not run)
```

readWindowsShortcut     *Reads a Microsoft Windows Shortcut (.lnk file)*

---

### Description

Reads a Microsoft Windows Shortcut (.lnk file).

### Usage

```
## Default S3 method:  
readWindowsShortcut(con, verbose=FALSE, ...)
```

### Arguments

con	A <a href="#">connection</a> or a <a href="#">character</a> string (filename).
verbose	If <a href="#">TRUE</a> , extra information is written while reading.
...	Not used.

### Details

The MIME type for a Windows Shortcut file is `application/x-ms-shortcut`.

### Value

Returns a [list](#) structure.

### Author(s)

Henrik Bengtsson

### References

- [1] Wotsit's Format, <http://www.wotsit.org/>, 2005.
- [2] Hager J, *The Windows Shortcut File Format* (as reverse-engineered by), version 1.0.
- [3] Microsoft Developer Network, *IShellLink Interface*, 2008. <http://msdn2.microsoft.com/en-us/library/bb774950.aspx>
- [4] Andrews D, *Parsing Windows Shortcuts (lnk) files in java*, `comp.lang.java.help`, Aug 1999. [http://groups.google.com/group/comp.lang.java.help/browse\\_thread/thread/a2e147b07d5480a2/](http://groups.google.com/group/comp.lang.java.help/browse_thread/thread/a2e147b07d5480a2/)
- [5] Multiple authors, *Windows shell links* (in Tcl), Tccler's Wiki, April 2008. <http://wiki.tcl.tk/1844>
- [6] Daniel S. Bensen, *Shortcut File Format (.lnk)*, Stdlib.com, April 24, 2009. <https://web.archive.org/web/20110817051855/http://www.stdlib.com/art6-Shortcut-File-Format-lnk.html> (was <http://www.stdlib.com/art6-Shortcut-File-Format-lnk.html>)
- [7] [MS-SHLLINK]: Shell Link (.LNK) Binary File Format, Microsoft Inc., September 25, 2009.



**See Also**

[createWindowsShortcut\(\)](#) and [filePath\(\)](#)

**Examples**

```
pathname <- system.file("data-ex/HISTORY.LNK", package="R.utils")
lnk <- readWindowsShortcut(pathname)

# Print all information
print(lnk)

# Get the relative path to the target file
history <- filePath(dirname(pathname), lnk$relativePath)

# Alternatively, everything in one call
history <- filePath(pathname, expandLinks="relative")
```

---

removeDirectory	<i>Removes a directory</i>
-----------------	----------------------------

---

**Description**

Removes a directory, and if requested, also its contents.

**Usage**

```
## Default S3 method:
removeDirectory(path, recursive=FALSE, mustExist=TRUE, ...)
```

**Arguments**

path	A <a href="#">character</a> string specifying the directory to be removed.
recursive	If <a href="#">TRUE</a> , subdirectories and files are also removed. If <a href="#">FALSE</a> , and directory is non-empty, an exception is thrown.
mustExist	If <a href="#">TRUE</a> , and the directory does not exist, an exception is thrown.
...	Not used.

**Value**

Returns (invisibly) [TRUE](#), the directory was successfully removed, otherwise [FALSE](#), unless an exception is thrown.

**Symbolic links**

This function can also be used to remove symbolic links to directories without removing the target. Note that neither [file.remove\(\)](#) nor [unlink\(\)](#) is capable of remove symbolic *directory* links on Windows.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally [unlink\(\)](#) is used.

---

resample

*Sample values from a set of elements*

---

**Description**

Sample values from a set of elements. Contrary to [sample\(\)](#), this function also works as expected when there is only one element in the set to be sampled, cf. [1]. This function originates from the example code of [sample\(\)](#) as of R v2.12.0.

**Usage**

```
## Default S3 method:  
resample(x, ...)
```

**Arguments**

`x` A [vector](#) of any length and data type.  
`...` Additional arguments passed to [sample.int\(\)](#).

**Value**

Returns a sampled [vector](#) of the same data types as argument `x`.

**Author(s)**

Henrik Bengtsson

**References**

[1] Henrik Bengtsson, *Using [sample\(\)](#) to sample one value from a single value?*, R-devel mailing list, 2010-11-03.

**See Also**

Internally [sample.int\(\)](#) is used.

---

saveObject	<i>Saves an object to a file or a connection</i>
------------	--

---

### Description

Saves an object to a file or a connection.

### Usage

```
## Default S3 method:  
saveObject(object, file=NULL, path=NULL, format=c("auto", "xdr", "rds"), compress=TRUE,  
  ..., safe=TRUE)
```

### Arguments

object	The object to be saved.
file	A filename or <a href="#">connection</a> where the object should be saved. If <code>NULL</code> , the filename will be the hash code of the object plus ".xdr".
path	Optional path, if file is a filename.
format	File format.
compress	If <code>TRUE</code> , the file is compressed to, otherwise not.
...	Other arguments accepted by <code>save()</code> in the base package.
safe	If <code>TRUE</code> and file is a file, then, in order to lower the risk for incomplete files, the object is first written to a temporary file, which is then renamed to the final name.

### Value

Returns (invisibly) the pathname or the [connection](#).

### Author(s)

Henrik Bengtsson

### See Also

[loadObject\(\)](#) to load an object from file. [digest](#) for how hash codes are calculated from an object. See also [saveRDS\(\)](#).

---

seqToHumanReadable	<i>Gets a short human readable string representation of an vector of indices</i>
--------------------	--

---

### Description

Gets a short human readable string representation of an vector of indices.

### Usage

```
## Default S3 method:  
seqToHumanReadable(idx, tau=2L, delimiter="-", collapse=", ", ...)
```

### Arguments

idx	A <a href="#">vector</a> of <a href="#">integer</a> indices.
tau	A non-negative <a href="#">integer</a> specifying the minimum span of of a contiguous sequences for it to be collapsed to <from>-<to>.
delimiter	A <a href="#">character</a> string delimiter.
collapse	A <a href="#">character</a> string used to collapse subsequences.
...	Not used.

### Author(s)

Henrik Bengtsson

### See Also

Internally, [seqToIntervals\(\)](#) is used.

### Examples

```
print(seqToHumanReadable(1:2))           # "1, 2"  
print(seqToHumanReadable(1:2, tau=1))    # "1-2"  
print(seqToHumanReadable(1:10))         # "1-10"  
print(seqToHumanReadable(c(1:10, 15:18, 20))) # "1-10, 15-18, 20"
```

---

seqToIntervals	<i>Gets all contiguous intervals of a vector of indices</i>
----------------	---

---

### Description

Gets all contiguous intervals of a vector of indices.

### Usage

```
## Default S3 method:  
seqToIntervals(idx, ...)
```

### Arguments

idx	A <a href="#">vector</a> of N <a href="#">integer</a> indices.
...	Not used.

### Value

An Nx2 [integer matrix](#).

### Author(s)

Henrik Bengtsson

### See Also

[\\*intervalsToSeq\(\)](#). To identify sequences of *equal* values, see [rle\(\)](#).

### Examples

```
x <- 1:10  
y <- seqToIntervals(x)  
print(y) # [1 10]  
  
x <- c(1:10, 15:18, 20)  
y <- seqToIntervals(x)  
print(y) # [1 10; 15 18; 20 20]  
  
z <- intervalsToSeq(y)  
print(z)  
stopifnot(all.equal(x,z))
```

---

setOption	<i>Sets a option in R</i>
-----------	---------------------------

---

**Description**

Sets a option in R by specifying its name as a [character](#) string.

**Usage**

```
## Default S3 method:
setOption(x, value, ...)
```

**Arguments**

x	The name of the option to be set.
value	The new value of the option.
...	Not used.

**Value**

Returns (invisibly) the previous value of the option.

**Author(s)**

Henrik Bengtsson

**See Also**

See [getOption\(\)](#) and "base::options".

---

Settings	<i>Class for applicational settings</i>
----------	---

---

**Description**

Package: R.utils

**Class Settings****Object**

```
~~|
```

```
~~+--Options
```

```
~~~~~|
```

```
~~~~~+--Settings
```

**Directly known subclasses:**

```
public static class Settings
  extends Options
```

Class for applicational settings.

**Usage**

```
Settings(basename=NULL, ...)
```

**Arguments**

basename	A <a href="#">character</a> string of the basename of the settings file.
...	Arguments passed to constructor of superclass <a href="#">Options</a> .

**Fields and Methods****Methods:**

<a href="#">findSettings</a>	Searches for the settings file in one or several directories.
<a href="#">getLoadedPathname</a>	Gets the pathname of the settings file loaded.
<a href="#">isModified</a>	Checks if settings has been modified compared to whats on file.
<a href="#">loadAnywhere</a>	Loads settings from file.
<a href="#">promptAndSave</a>	Prompt user to save modified settings.
<a href="#">saveAnywhere</a>	Saves settings to file.

**Methods inherited from Options:**

as.character, as.list, equals, getLeaves, getOption, hasOption, names, nbrOfOptions, setOption, str

**Methods inherited from Object:**

\$, \$<-, [[, [[<-, as.character, attach, attachLocally, clearCache, clearLookupCache, clone, detach, equals, extend, finalize, getEnvironment, getFieldModifier, getFieldModifiers, getFields, getInstantiationTime, getStaticInstance, hasField, hashCode, ll, load, names, objectSize, print, save

**Load settings with package and save on exit**

Here is a generic `.First.lib()` function for loading settings with package. It also (almost) assures that the package is detached when R finishes. See `onSessionExit()` why it is not guaranteed!

The almost generic `.Last.lib()` function, which will prompt user to save settings, is called when a package is detached.

It is custom to put these functions in a file named `zzz.R`.

**.First.lib():**

```
.First.lib <- function(libname, pkgname) {
```

```

# Write a welcome message when package is loaded
pkg <- Package(pkgname)
assign(pkgname, pkg, pos=getPosition(pkg))

# Read settings file "<pkgname>Settings" and store it in package
# variable '<pkgname>Settings'.
varname <- paste(pkgname, "Settings")
basename <- paste(".", varname, sep="")
settings <- Settings$loadAnywhere(basename, verbose=TRUE)
if (is.null(settings))
  settings <- Settings(basename)
assign(varname, settings, pos=getPosition(pkg))

# Detach package when R finishes, which will save package settings too.
onSessionExit(function(...) detachPackage(pkgname))

packageStartupMessage(getName(pkg), " v", getVersion(pkg),
  " (" , getDate(pkg), ") successfully loaded. See ?", pkgname,
  " for help.\n", sep="")
} # .First.lib()

```

**.Last.lib():**

```

.Last.lib <- function(libpath) {
  pkgname <- "<package name>"

  # Prompt and save package settings when package is detached.
  varname <- paste(pkgname, "Settings", sep="")
  if (exists(varname)) {
    settings <- get(varname)
    if (inherits(settings, "Settings"))
      promptAndSave(settings)
  }
} # .Last.lib()

```

**Author(s)**

Henrik Bengtsson

**Examples**

```

# Load settings from file, or create default settings
basename <- "some.settings"
settings <- Settings$loadAnywhere(basename)
if (is.null(settings))
  settings <- Settings(basename)

# Set default options, if missing.

```



```
setOption(settings, "graphics/verbose", TRUE, overwrite=FALSE)
setOption(settings, "io/verbose", Verbose(threshold=-1), overwrite=FALSE)

# Save and reload settings
path <- tempdir()
saveAnywhere(settings, path=path)
settings2 <- Settings$loadAnywhere(basename, paths=path)

# Clean up
file.remove(getLoadedPathname(settings2))

# Assert correctness
stopifnot(equals(settings, settings2))
```

---

shell.exec2

*Open a file or URL using Windows File Associations*

---

### Description

Open a file or URL using Windows File Associations using `shell.exec()` but makes some tweaks to filenames to make them more likely to be opened properly.

*This function is only applicable on Windows systems.*

### Usage

```
shell.exec2(file)
```

### Arguments

`file` A [character](#) string specifying a file or an URL.

### Details

Before passing a *file* on the file system to `shell.exec()`, this function: (i) unmaps any mapped drive letters used in the pathname (e.g. `'X:/foo.bar.html'` to `'C:/Users/Joe/bar.html'`), (ii) and replaces any forward slashed with backward ones (e.g. `'C:/Users/Joe/bar.html'` to `'C:\Users\Joe\bar.html'`). URLs are passed as is to `shell.exec()`.

The reason for (i) is that some web browsers (e.g. Google Chrome) will not open files on mapped drives. The reason for (ii) is that if forward slashes are used, then `shell.exec()` will give an error that the file was not found (at least with the default Windows shell).

### Value

Returns nothing.

**Setting on startup**

The intended usage of this function is to set it as the default browser for [browseURL](#). Just add the following to your [.Rprofile](#) file:

```
if (.Platform$OS.type == "windows")
  options(browser=function(...) R.utils::shell.exec2(...))
```

This will only load (not attach) the **R.utils** package when the browser function is actual used.

**Author(s)**

Henrik Bengtsson

---

SmartComments

*Abstract class SmartComments*

---

**Description**

Package: R.utils

**Class SmartComments**

[Object](#)

~~|

~~+--SmartComments

**Directly known subclasses:**

[LComments](#), [VComments](#)

public abstract static class **SmartComments**  
 extends [Object](#)

Abstract class SmartComments.

**Usage**

```
SmartComments(letter=NA, ...)
```

**Arguments**

letter            A single [character](#).

...                Not used.

## Details

A "smart" source-code comment is an R comment, which start with a '#', but is followed by a single letter, then a single symbol and a second '#' and then an option character string, and there must not be any code before the comment on the same line. In summary, a smart comment line has format: <white spaces>#<letter><symbol># <some text>.

Example code with two smart comments (VComments):

```
x <- 2
#V1# threshold=-1
#Vc# A v-comment log message
cat("Hello world")
```

which after compilation becomes

```
x <- 2
verbose <- Verbose(threshold=-1)
if (verbose) { cat(verbose, "A v-comment log message"); }
cat("Hello world")
```

## Fields and Methods

### Methods:

<a href="#">compile</a>	Preprocess a vector of code lines.
<a href="#">convertComment</a>	Converts a single smart comment to R code.
<a href="#">parse</a>	Parses one single smart comment.
<a href="#">reset</a>	Resets a SmartComments compiler.
<a href="#">validate</a>	Validates the compiled lines.

### Methods inherited from Object:

\$, \$<-, [[, [[<-, as.character, attach, attachLocally, clearCache, clearLookupCache, clone, detach, equals, extend, finalize, getEnvironment, getFieldModifier, getFieldModifiers, getFields, getInstantiationTime, getStaticInstance, hasField, hashCode, ll, load, names, objectSize, print, save

### Author(s)

Henrik Bengtsson

### See Also

[VComments](#).

---

sourceDirectory	<i>Sources files recursively to either local or global environment</i>
-----------------	--

---

### Description

Sources files recursively to either local or global environment.

### Usage

```
## Default S3 method:
sourceDirectory(path, pattern=".*[.](r|R|s|S|q)([.](lnk|LNK))*$", recursive=TRUE,
  envir=parent.frame(), onError=c("error", "warning", "skip"), modifiedOnly=TRUE, ...,
  verbose=FALSE)
```

### Arguments

path	A path to a directory to be sourced.
pattern	A regular expression file name pattern to identify source code files.
recursive	If <b>TRUE</b> , subdirectories are recursively sourced first, otherwise not.
envir	An <b>environment</b> in which the code should be evaluated.
onError	If an error occurs, the error may stop the job, give a warning, or silently be skipped.
modifiedOnly	If <b>TRUE</b> , only files that are modified since the last time they were sourced are sourced, otherwise regardless.
...	Additional arguments passed to <code>sourceTo()</code> .
verbose	A <b>logical</b> or a <b>Verbose</b> object.

### Value

Returns a **vector** of the full pathnames of the files sourced.

### Details

Subdirectories and files in each (sub-)directory are sourced in lexicographic order.

### Hooks

This method does not provide hooks, but the internally used `sourceTo()` does.

### Author(s)

Henrik Bengtsson

### See Also

`sourceTo()` and compare to `source()`.

---

sourceTo	<i>Parses and evaluates code from a file or a connection</i>
----------	--

---

### Description

Parses and evaluates code from a file or a connection. This has the same effect as if `source(..., local=TRUE)` would have been called from within the given environment. This is useful when setting up a new local working environment.

### Usage

```
## Default S3 method:  
sourceTo(file, path=NULL, chdir=FALSE, ..., local=TRUE, envir=parent.frame(),  
         modifiedOnly=FALSE)
```

### Arguments

file	A <a href="#">connection</a> or a <a href="#">character</a> string giving the pathname of the file or URL to read from.
path	An optional <a href="#">character</a> string giving the path to the file. Ignored if file is a connection.
chdir	If <a href="#">TRUE</a> and file is a pathname, the R working directory is temporarily changed to the directory containing file for evaluating.
...	Arguments to <a href="#">source()</a> . If argument file is not explicitly given, the first argument is assumed to be the file argument. This argument is converted into a string by <code>as.character()</code> .
local	If <a href="#">FALSE</a> , evaluation is done in the global environment, otherwise in the calling environment.
envir	An <a href="#">environment</a> in which <a href="#">source()</a> should be called. If <a href="#">NULL</a> , the global environment is used.
modifiedOnly	If <a href="#">TRUE</a> , the file is sourced only if modified since the last time it was sourced, otherwise regardless.

### Value

Return the result of [source\(\)](#).

### Hooks

This methods recognizes the hook `sourceTo/onPreprocess`, which is called after the lines in file has been read, but before they have been parsed by the R parser, cf. [parse\(\)](#). An `onPreprocess` hook function should take a [character vector](#) of code lines and return a [character vector](#) of code lines. This can for instance be used to pre-process R source code with special directives such as [VComments](#).

Note that only one hook function can be used for this function, otherwise an error is generated.

**Author(s)**

Henrik Bengtsson

**See Also**[sourceDirectory\(\)](#), [sys.source\(\)](#) and [source\(\)](#).**Examples**

```

# -----
# Example 1
# -----
cat("=== Example 1 =====\n")

foo <- function(file, ...) {
  cat("Local objects before calling sourceTo():\n")
  print(ls())

  res <- sourceTo(file, ...)

  cat("Local objects after calling sourceTo():\n")
  print(ls())
}

cat("Global objects before calling foo():\n")
lsBefore <- NA
lsBefore <- ls()
foo(file=textConnection(c('a <- 1', 'b <- 2')))

cat("Global objects after calling foo():\n")
stopifnot(length(setdiff(ls(), lsBefore)) == 0)

# -----
# Example 2 - with VComments preprocessor
# -----
cat("=== Example 2 =====\n")

preprocessor <- function(lines, ...) {
  cat("-----\n")
  cat("Source code before preprocessing:\n")
  displayCode(code=lines, pager="console")
  cat("-----\n")
  cat("Source code after preprocessing:\n")
  lines <- VComments$compile(lines)
  displayCode(code=lines, pager="console")
  cat("-----\n")
  lines
}

oldHooks <- getHook("sourceTo/onPreprocess")
setHook("sourceTo/onPreprocess", preprocessor, action="replace")

```

```
code <- c(
  'x <- 2',
  '#V1# threshold=-1',
  '#Vc# A v-comment log message',
  'print("Hello world")'
)
fh <- textConnection(code)
sourceTo(fh)
setHook("sourceTo/onPreprocess", oldHooks, action="replace")
```

---

**splitByPattern***Splits a single character string by pattern*

---

### Description

Splits a single character string by pattern. The main difference compared to `strsplit()` is that this method also returns the part of the string that matched the pattern. Also, it only takes a single character string.

### Usage

```
## Default S3 method:
splitByPattern(str, pattern, ...)
```

### Arguments

<code>str</code>	A single <a href="#">character</a> string to be split.
<code>pattern</code>	A regular expression <a href="#">character</a> string.
<code>...</code>	Not used.

### Value

Returns a named [character vector](#) with names equal to "TRUE" if element is a pattern part and "FALSE" otherwise.

### Author(s)

Henrik Bengtsson

### See Also

Compare to [strsplit\(\)](#).

### Examples

```
rspCode <- "<body>Hello <%=\"world\"%></body>"
rspParts <- splitByPattern(rspCode, pattern="<%.*%>")
cat(rspCode, "\n")
print(rspParts)
```

---

stext	<i>Writes text in the margin along the sides of a plot</i>
-------	--

---

### Description

Writes text in the margin along the sides of a plot.

### Usage

```
## Default S3 method:  
stext(text, side=1, line=0, pos=0.5, margin=c(0.2, 0.2),  
      charDim=c(strwidth("M", cex = cex), strheight("M", cex = cex)), cex=par("cex"), ...)
```

### Arguments

text	The text to be written. See <a href="#">mtext</a> for details.
side	An <a href="#">integer</a> specifying which side to write the text on. See <a href="#">mtext</a> for details.
line	A <a href="#">numeric</a> specifying on which line to write on.
pos	A <a href="#">numeric</a> , often in [0,1], specifying the position of the text relative to the left and right edges.
margin	A <a href="#">numeric vector</a> length two specifying the text margin.
charDim	A <a href="#">numeric vector</a> length two specifying the size of a typical symbol.
cex	A <a href="#">numeric</a> specifying the character expansion factor.
...	Additional arguments passed to <a href="#">mtext</a> .

### Value

Returns what [mtext](#) returns.

### Author(s)

Henrik Bengtsson

### See Also

Internally [mtext](#) is used.



---

subplots *Creates a grid of subplots*

---

### Description

Creates a grid of subplots in the current figure. If arguments `nrow` and `ncol` are given a `nrow`-by-`ncol` grid of subplots are created. If only argument `n` is given then a `r`-by-`s` grid is created where  $|r-s| \leq 1$ , i.e. a square or almost a square of subplots is created. If `n` and `nrow` is given then a grid with `nrow` rows and at least `n` subplots are created. Similar if `n` and `ncol` is given. The argument `byrow` specifies if the order of the subplots should be rowwise (`byrow=TRUE`) or columnwise.

### Usage

```
## Default S3 method:
subplots(n=1, nrow=NULL, ncol=NULL, byrow=TRUE, ...)
```

### Arguments

<code>n</code>	If given, the minimum number of subplots.
<code>nrow</code>	If given, the number of rows the grid of subplots should contain.
<code>ncol</code>	If given, the number of columns the grid of subplots should contain.
<code>byrow</code>	If <code>TRUE</code> , the panels are ordered row by row in the grid, otherwise column by column.
<code>...</code>	Not used.

### Value

Returns the `matrix` containing the order of plots.

### Author(s)

Henrik Bengtsson

### See Also

[layout](#) and `layout.show()`.

### Examples

```
subplots(nrow=2, ncol=3) # 2-by-3 grid of subplots
subplots(n=6, nrow=2)   # 2-by-3 grid of subplots
subplots(n=5, ncol=2)   # 3-by-2 grid of subplots
subplots(1)             # (Reset) to a 1-by-1 grid of subplots
subplots(2)             # 1-by-2 grid of subplots
subplots(3)             # 2-by-2 grid of subplots
l <- subplots(8)        # 3-by-3 grid of subplots
layout.show(length(l))
```

---

System

*Static class to query information about the system*


---

**Description**

Package: R.utils

**Class System**[Object](#)

~~|

~~+--System

**Directly known subclasses:**public static class **System**extends [Object](#)

The System class contains several useful class fields and methods. It cannot be instantiated.

**Fields and Methods****Methods:**

<a href="#">currentTimeMillis</a>	Get the current time in milliseconds.
<a href="#">findGhostscript</a>	Searches for a Ghostview executable on the current system.
<a href="#">findGraphicsDevice</a>	Searches for a working PNG device.
<a href="#">getHostname</a>	Retrieves the computer name of the current host.
<a href="#">getMappedDrivesOnWindows</a>	-
<a href="#">getUsername</a>	Retrieves the name of the user running R.
<a href="#">mapDriveOnWindows</a>	-
<a href="#">openBrowser</a>	Opens an HTML document using the OS default HTML browser.
<a href="#">parseDebian</a>	Parses a string, file or connection for Debian formatted parameters.
<a href="#">unmapDriveOnWindows</a>	-

**Methods inherited from Object:**

\$, \$<-, [[, [[<-, as.character, attach, attachLocally, clearCache, clearLookupCache, clone, detach, equals, extend, finalize, getEnvironment, getFieldModifier, getFieldModifiers, getFields, getInstantiationTime, getStaticInstance, hasField, hashCode, ll, load, names, objectSize, print, save

**Author(s)**

Henrik Bengtsson

---

`systemR`*Launches another R process from within R*

---

## Description

Launches another R process from within R via `system()` by automatically locating the R executable, cf [1].

## Usage

```
## Default S3 method:  
systemR(command="", ..., Rcommand="R", verbose=FALSE)
```

## Arguments

<code>command</code>	A <a href="#">character</a> string be appended to the <code>system()</code> call. If a <a href="#">vector</a> , then the strings are concatenated separated with a space.
<code>...</code>	Additional arguments passed to <code>system()</code> .
<code>Rcommand</code>	A <a href="#">character</a> string specifying the basename of the R executable.
<code>verbose</code>	A <a href="#">logical</a> or a <a href="#">Verbose</a> object.

## Value

Returns what `system()` returns.

## Author(s)

Henrik Bengtsson

## References

[1] R-devel thread 'Best way to locate R executable from within R?', May 22, 2012.

## See Also

The R executable is located using `R.home()`, which is then launched using `system()`.

## Examples

```
res <- systemR("--slave -e cat(runif(1))", intern=TRUE)  
cat("A random number: ", res, "\n", sep="")
```

---

TextStatusBar                    *A status bar at the R prompt that can be updated*

---

## Description

Package: R.utils

**Class TextStatusBar**

[Object](#)

~~|

~~+--TextStatusBar

### Directly known subclasses:

public static class **TextStatusBar**

extends [Object](#)

A status bar at the R prompt that can be updated.

## Usage

```
TextStatusBar(fmt=paste("%-",getOption("width") - 1, "s", sep = ""), ...)
```

## Arguments

`fmt`                    A [character](#) format string to be used by [sprintf\(\)](#). Default is a left-aligned string of full width.

`...`                    Named arguments to be passed to [sprintf\(\)](#) together with the format string.

## Details

A label with name `hfill` can be used for automatic horizontal filling. It must be [numeric](#) and be immediate before a string label such that a `hfill` label and the following string label together specifies an `sprintf` format such as `"%*-s"`. The value of `hfill` will be set such that the resulting status bar has width equal to `getOption("width")-1` (the reason for the `-1` is to prevent the text status bar from writing into the next line). If more than one `hfill` label is used their widths will be uniformly distributed. Left over spaces will be distributed between `hfill` labels with initial values of one.

## Fields and Methods

### Methods:

[flush](#)                    Flushes the output.

<code>getLabel</code>	Gets the current value of a label.
<code>newline</code>	Writes a newline.
<code>popMessage</code>	Adds a message above the status bar.
<code>setLabel</code>	Sets the value of a label.
<code>setLabels</code>	Sets new values of given labels.
<code>update</code>	Updates the status bar (visually).
<code>updateLabels</code>	Sets the new values of given labels and updates the status bar.

**Methods inherited from Object:**

`$`, `$<-`, `[[`, `[[<-`, `as.character`, `attach`, `attachLocally`, `clearCache`, `clearLookupCache`, `clone`, `detach`, `equals`, `extend`, `finalize`, `getEnvironment`, `getFieldModifier`, `getFieldModifiers`, `getFields`, `getInstantiationTime`, `getStaticInstance`, `hasField`, `hashCode`, `ll`, `load`, `names`, `objectSize`, `print`, `save`

**Author(s)**

Henrik Bengtsson

**Examples**

```
# -----
# Read all HTML files in the base package
# -----
path <- system.file(package="base")
files <- list.files(path, recursive=TRUE, full.names=TRUE)
files <- files[sapply(files, FUN=isFile)]
nfiles <- length(files)

cat(sprintf("Reading %d files in %s:\n", nfiles, path))

# Create a status bar with four labels
sb <- TextStatusBar("File: %-*s [%3.0f%% %7.0f bytes %-8s]",
                    hfill=1, file="", progress=0, nbytes=0L, time="")

nbytes <- 0L
for (kk in seq_len(nfiles)) {
  file <- files[kk]

  # Update the status bar
  if (sb) {
    setLabel(sb, "progress", 100*kk/nfiles)
    if (kk %% 10 == 1 || kk == nfiles)
      setLabel(sb, "file", substr(basename(file), 1, 44))

    size <- file.info(file)$size
    # popMessage() calls update() too
    popMessage(sb, sprintf("Processing %s (%.2fkB)",
                           basename(file), size/1024))

    flush(sb)
  }
}
```

```

# Read the file
bfr <- readBin(file, what="raw", n=size)
nbytes <- nbytes + size

# Emulate a slow process
if (interactive()) Sys.sleep(rexp(1, rate=60))

# Update the status bar
if (sb) {
  setLabel(sb, "nbytes", nbytes)
  setLabel(sb, "time", format(Sys.time(), "%H:%M:%S"))
  update(sb)
}
}
setLabel(sb, "file", "<done>")
update(sb)
cat("\n")

```

---

TimeoutException

*TimeoutException represents timeout errors*


---

## Description

Package: R.utils

### Class TimeoutException

#### Object

```

~|
~+--try-error
~~~~~|
~~~~~+--condition
~~~~~|
~~~~~+--error
~~~~~|
~~~~~+--simpleError
~~~~~|
~~~~~+--Exception
~~~~~|
~~~~~+--TimeoutException

```

#### Directly known subclasses:

```

public static class TimeoutException
extends Exception

```

TimeoutException represents timeout errors occurring when a set of R expressions executed did not finish in time.

**Usage**

```
TimeoutException(..., cpu=NA, elapsed=NA)
```

**Arguments**

... Any arguments accepted by [Exception](#).

cpu, elapsed The maximum time the R expressions were allowed to be running before the timeout occurred as measured in CPU time and (physically) elapsed time.

**Fields and Methods****Methods:**

[getMessage](#) Gets the message of the exception.

**Methods inherited from Exception:**

as.character, getCall, getCalls, getLastException, getMessage, getStackTrace, getWhen, print, printStackTrace, throw

**Methods inherited from error:**

as.character, throw

**Methods inherited from condition:**

abort, as.character, conditionCall, conditionMessage, print

**Methods inherited from Object:**

\$, \$<-, [[, [[<-, as.character, attach, attachLocally, clearCache, clearLookupCache, clone, detach, equals, extend, finalize, getEnvironment, getFieldModifier, getFieldModifiers, getFields, getInstantiationTime, getStaticInstance, hasField, hashCode, ll, load, names, objectSize, print, save

**Author(s)**

Henrik Bengtsson

**See Also**

For detailed information about exceptions see [Exception](#).

---

touchFile

*Updates the timestamp of a file*

---

**Description**

Updates the timestamp of a file. Currently, it is only possible to change the timestamp specifying when the file was last modified, and time can only be set to the current time.

**Usage**

```
## Default S3 method:  
touchFile(pathname, ...)
```

**Arguments**

```
pathname      A character vector specifying files to be updated.  
...          Not used.
```

**Value**

Returns (invisibly) a **vector** of the old timestamps.

**Author(s)**

Henrik Bengtsson

**References**

[1] R-devel mailing list thread *Unix-like touch to update modification timestamp of file?*, started on 2008-02-26. <http://stat.ethz.ch/pipermail/r-devel/2008-February/048542.html>

**See Also**

Internally, `Sys.setFileTime()` (iff available) and `file.info()` are utilized.

**Examples**

```
# 1. Create a file  
pathname <- tempfile()  
cat(file=pathname, "Hello world!")  
md5a <- digest::digest(pathname, file=TRUE)  
  
# 2. Current time stamp  
ta <- file.info(pathname)$mtime  
print(ta)  
  
# 3. Update time stamp  
Sys.sleep(1.2)  
touchFile(pathname)  
tb <- file.info(pathname)$mtime  
print(tb)  
  
# 4. Verify that the timestamp got updated  
stopifnot(tb > ta)  
  
# 5. Verify that the contents did not change  
md5b <- digest::digest(pathname, file=TRUE)  
stopifnot(identical(md5a, md5b))
```



---

toUrl	<i>Converts a pathname into a URL</i>
-------	---------------------------------------

---

**Description**

Converts a pathname into a URL starting with file://.

**Usage**

```
## Default S3 method:  
toUrl(pathname, safe=TRUE, ...)
```

**Arguments**

pathname	A <a href="#">character vector</a> of pathnames to be made into URLs.
safe	If <a href="#">TRUE</a> , certain "unsafe" characters are escaped.
...	Not used.

**Value**

Returns a [character vector](#).

**Author(s)**

Henrik Bengtsson

**See Also**

[URLencode](#).

---

unwrap.array	<i>Unwrap an array, matrix or a vector to an array of more dimensions</i>
--------------	---

---

**Description**

Unwrap an array, matrix or a vector to an array of more dimensions. This is done by splitting up each dimension into several dimension based on the names of that dimension.

**Usage**

```
## S3 method for class 'array'  
unwrap(x, split=rep("[]", length(dim(x))), drop=FALSE, ...)
```

**Arguments**

x	An <a href="#">array</a> or a <a href="#">matrix</a> .
split	A <a href="#">list</a> or a <a href="#">character vector</a> . If a <a href="#">list</a> , it should contain <a href="#">functions</a> that takes a <a href="#">character vector</a> as the first argument and optional ... arguments. Each function should split the <a href="#">vector</a> into a <a href="#">list</a> of same length and where all elements contains the same number of parts. If a <a href="#">character vector</a> , each element split[i] is replaced by a <a href="#">function</a> call <code>function(names, ...) strsplit(names, split=split[i])</code> .
drop	If <a href="#">TRUE</a> , dimensions of of length one are dropped, otherwise not.
...	Arguments passed to the <a href="#">split functions</a> .

**Details**

Although not tested thoroughly, `unwrap()` should be the inverse of `wrap()` such that `identical(unwrap(wrap(x)), x)` holds.

**Value**

Returns an [array](#).

**Author(s)**

Henrik Bengtsson

**See Also**

[\\*wrap\(\)](#).

**Examples**

```
## Not run: See ?wrap.array for an example
```

---

useRepos

*Sets package repositories*

---

**Description**

Sets package repositories.

**Usage**

```
useRepos(repos=NULL, where=c("before", "after", "replace"), unique=TRUE, fallback=TRUE,
...)
```

**Arguments**

repos	A <a href="#">character vector</a> of repositories to use. If <code>NULL</code> , nothing is replaced.
where	A <a href="#">character</a> string specifying how to add them to the current set of repositories.
unique	If <code>TRUE</code> , only unique repositories are set.
fallback	If <code>TRUE</code> , any remaining non-specified repository value of format <code>'...@'</code> (e.g. <code>@CRAN@</code> ) that could not be recovered by other means, will be assigned to a pre-defined known value, if possible. If so, then an informative warning is given.
...	Not used.

**Value**

Returns a [list](#) with element 'repos' reflecting `options("repos")` as the options where prior to calling this function.

**Author(s)**

Henrik Bengtsson

**See Also**

[withRepos\(\)](#).

---

VComments

*The VComments class*


---

**Description**

Package: R.utils

**Class VComments****Object**

```

~~|
~~+--SmartComments
~~~~~|
~~~~~+--VComments

```

**Directly known subclasses:**

[LComments](#)

```

public static class VComments
extends SmartComments

```

The VComments class.

**Usage**

```
VComments(letter="V", verboseName="verbose", ...)
```

**Arguments**

letter	The smart letter.
verboseName	The name of the verbose object.
...	Not used.

**Details**

The 'v' in VComments stands for 'verbose', because of its relationship to the [Verbose](#) class. Here is a list of VComments and the R code that replaces each of them by the compiler:

**Constructors**

- `\#V0\#[<args>]` - `NullVerbose(<args>)`
- `\#V1\#[<args>]` - `Verbose(<args>)`

**Controls**

- `\#V=\#[<variable>]` - Sets the name of the <verbose> object. Default is 'verbose'.
- `\#V^\#[<threshold>]` - `setThreshold(<verbose>, <threshold>)`
- `\#V?\#[<expression>]` - `if (isVisible(<verbose>)) { <expression> }`
- `\#V@\#[<level>]` - `setDefaultLevel(<verbose>, <level>)`
- `\#Vm\#[<method> <args>]` - `<method>(<verbose>, <args>)`

**Enters and exits**

- `\#V+\#[<message>]` - `enter(<verbose>, <message>)`
- `\#V-\#[<message>]` - `exit(<verbose>, <message>)`
- `\#V!\#[<message>]` - `pushState(<verbose>)`  
`on.exit(popState(<verbose>))`  
`If <message>, enter(<verbose>, <message>)`

**Simple output**

- `\#Vn\#[<ignored>]` - `newline(<verbose>)`
- `\#Vr\#[<ignored>]` - `ruler(<verbose>)`
- `\#Vt\#[<ignored>]` - `timestamp(<verbose>)`
- `\#Vw\#[<title>]` - `warnings(<verbose>, <title>)`

**Output messages**

- `\#Vc\#[<message>]` - `cat(<verbose>, <message>)`
- `\#Ve\#[<expression>]` - `eval(<verbose>, <expression>)`
- `\#Vh\#[<message>]` - `header(<verbose>, <message>)`
- `\#Vp\#[<object>]` - `print(<verbose>, <object>)`
- `\#Vs\#[<object>]` - `summary(<verbose>, <object>)`
- `\#Vz\#[<object>]` - `str(<verbose>, <object>)`

## Fields and Methods

### Methods:

<code>convertComment</code>	Converts a verbose comment to R code.
<code>reset</code>	Resets a VComments compiler.
<code>validate</code>	Validates the compiled lines.

### Methods inherited from SmartComments:

`compile`, `convertComment`, `parse`, `reset`, `validate`

### Methods inherited from Object:

`$`, `$<-`, `[[`, `[[<-`, `as.character`, `attach`, `attachLocally`, `clearCache`, `clearLookupCache`, `clone`, `detach`, `equals`, `extend`, `finalize`, `getEnvironment`, `getFieldModifier`, `getFieldModifiers`, `getFields`, `getInstantiationTime`, `getStaticInstance`, `hasField`, `hashCode`, `ll`, `load`, `names`, `objectSize`, `print`, `save`

## Author(s)

Henrik Bengtsson

## Examples

```
filename <- system.file("data-ex/exampleVComments.R", package="R.utils")
lines <- readLines(filename)

cat("Code before preprocessing:\n")
displayCode(code=lines, pager="console")

lines <- VComments$compile(lines)

cat("Code after preprocessing:\n")
displayCode(code=lines, pager="console")
```

**Description**

Package: R.utils

**Class Verbose**

[Object](#)

~~|

~~+--Verbose

**Directly known subclasses:**

[MultiVerbose](#), [NullVerbose](#)

public static class **Verbose**

extends [Object](#)

Class to writing verbose messages to a connection or file.

**Usage**

```
Verbose(con=stderr(), on=TRUE, threshold=0, asGString=TRUE, timestamp=FALSE,
        removeFile=TRUE, core=TRUE, ...)
```

**Arguments**

con	A <a href="#">connection</a> or a <a href="#">character</a> string filename.
on	A <a href="#">logical</a> indicating if the writer is on or off.
threshold	A <a href="#">numeric</a> threshold that the level argument of any write method has to be equal to or larger than in order to the message being written. Thus, the lower the threshold is the more and more details will be outputted.
timestamp	If <a href="#">TRUE</a> , each output is preceded with a timestamp.
removeFile	If <a href="#">TRUE</a> and con is a filename, the file is first deleted, if it exists.
asGString	If <a href="#">TRUE</a> , all messages are interpreted as <a href="#">GString</a> before being output, otherwise not.
core	Internal use only.
...	Not used.

**Fields and Methods****Methods:**

<a href="#">as.character</a>	Returns a character string version of this object.
<a href="#">as.double</a>	Gets a numeric value of this object.
<a href="#">as.logical</a>	Gets a logical value of this object.
<a href="#">capture</a>	Captures output of a function.
<a href="#">cat</a>	Concatenates and prints objects if above threshold.

<code>enter</code>	Writes a message and indents the following output.
<code>enterf</code>	-
<code>equals</code>	Checks if this object is equal to another.
<code>evaluate</code>	Evaluates a function and prints its results if above threshold.
<code>exit</code>	Writes a message and unindents the following output.
<code>getThreshold</code>	Gets current verbose threshold.
<code>getTimestampFormat</code>	Gets the default timestamp format.
<code>header</code>	Writes a header.
<code>isOn</code>	Checks if the output is on.
<code>isVisible</code>	Checks if a certain verbose level will be shown or not.
<code>less</code>	Creates a cloned instance with a higher threshold.
<code>more</code>	Creates a cloned instance with a lower threshold.
<code>newline</code>	Writes one or several empty lines.
<code>off</code>	Turn off the output.
<code>on</code>	Turn on the output.
<code>popState</code>	-
<code>print</code>	Prints objects if above threshold.
<code>printf</code>	Formats and prints object if above threshold.
<code>pushState</code>	Pushes the current indentation state of the Verbose object.
<code>ruler</code>	Writes a ruler.
<code>setDefaultLevel</code>	Sets the current default verbose level.
<code>setThreshold</code>	Sets verbose threshold.
<code>setTimestampFormat</code>	Sets the default timestamp format.
<code>str</code>	Prints the structure of an object if above threshold.
<code>summary</code>	Generates a summary of an object if above threshold.
<code>timestamp</code>	Writes a timestamp.
<code>timestampOff</code>	-
<code>timestampOn</code>	Turns automatic timestamping on and off.
<code>warnings</code>	Outputs any warnings recorded.
<code>writeRaw</code>	Writes objects if above threshold.

#### Methods inherited from Object:

`$`, `$<-`, `[]`, `[[<-`, `as.character`, `attach`, `attachLocally`, `clearCache`, `clearLookupCache`, `clone`, `detach`, `equals`, `extend`, `finalize`, `getEnvironment`, `getFieldModifier`, `getFieldModifiers`, `getFields`, `getInstantiationTime`, `getStaticInstance`, `hasField`, `hashCode`, `ll`, `load`, `names`, `objectSize`, `print`, `save`

#### Output levels

As a guideline, use the following levels when outputting verbose/debug message using the Verbose class. For a message to be shown, the output level must be greater than (not equal to) current threshold. Thus, the lower the threshold is set, the more messages will be seen.

- `<= -100` Only for debug messages, i.e. messages containing all necessary information for debugging purposes and to find bugs in the code. Normally these messages are so detailed so they will be a pain for the regular user, but very useful for bug reporting and bug tracking by the developer.

- -99 – -11 Detailed verbose messages. These will typically be useful for the user to understand what is going on and do some simple debugging fixing problems typically due to themselves and not due to bugs in the code.
- -10 – -1 Verbose messages. For example, these will typically report the name of the file to be read, the current step in a sequence of analysis steps and so on. These message are not very useful for debugging.
- 0 Default level in all output methods and default threshold. Thus, by default, messages at level 0 are not shown.
- $\geq +1$  Message that are always outputted (if threshold is kept at 0). We recommend not to output message at this level, because methods should be quiet by default (at the default threshold 0).

### A compatibility trick and a speed-up trick

If you want to include calls to `Verbose` in a package of yours in order to debug code, but not use it otherwise, you might not want to load `R.utils` all the time, but only for debugging. To achieve this, the value of a reference variable to a `Verbose` class is always set to `TRUE`, cf. typically an Object reference has value `NA`. This makes it possible to use the reference variable as a first test before calling `Verbose` methods. Example:

```
foo <- function(..., verbose=FALSE) {
  # enter() will never be called if verbose==FALSE, thus no error.
  verbose && enter(verbose, "Loading")
}
```

Thus, `R.utils` is not required for `foo()`, but for `foo(verbose==Verbose(level=-1))` it is.

Moreover, if using the `NullVerbose` class for ignoring all verbose messages, the above trick will indeed speed up the code, because the value of a `NullVerbose` reference variable is always `FALSE`.

### Extending the `Verbose` class

If extending this class, make sure to output messages via `*writeRaw()` or one of the other output methods (which in turn all call the former). This guarantees that `*writeRaw()` has full control of the output, e.g. this makes it possible to split output to standard output and to file.

### Author(s)

Henrik Bengtsson

### See Also

[NullVerbose](#).



**Examples**

```

verbose <- Verbose(threshold=-1)

header(verbose, "A verbose writer example", padding=0)

enter(verbose, "Analysis A")
for (kk in 1:10) {
  printf(verbose, "step %d\n", kk)
  if (kk == 2) {
    cat(verbose, "Turning ON automatic timestamps")
    timestampOn(verbose);
  } else if (kk == 4) {
    timestampOff(verbose);
    cat(verbose, "Turned OFF automatic timestamps")
    cat(verbose, "Turning OFF verbose messages for steps ", kk, "-6")
    off(verbose)
  } else if (kk == 6) {
    on(verbose)
    cat(verbose, "Turned ON verbose messages just before step ", kk+1)
  }

  if (kk %in% c(5,8)) {
    enterf(verbose, "Sub analysis #d", kk)
    for (jj in c("i", "ii", "iii")) {
      cat(verbose, "part ", jj)
    }
    exit(verbose)
  }
}
cat(verbose, "All steps completed!")
exit(verbose)

ruler(verbose)
cat(verbose, "Demo of some other methods:")
str(verbose, c(a=1, b=2, c=3))
print(verbose, c(a=1, b=2, c=3))
summary(verbose, c(a=1, b=2, c=3))
evaluate(verbose, rnorm, n=3, mean=2, sd=3)

ruler(verbose)
newline(verbose)

```

---

withCapture

*Evaluates an expression and captures the code and/or the output*


---

**Description**

Evaluates an expression and captures the code and/or the output.

**Usage**

```
withCapture(expr, replace=getOption("withCapture/substitute", ".x."), code=TRUE,
  output=code, ..., max.deparse.length=getOption("max.deparse.length", 10000), trim=TRUE,
  newline=getOption("withCapture/newline", TRUE), collapse="\n", substitute=replace,
  envir=parent.frame())
```

**Arguments**

expr	The R expression to be evaluated.
replace	An optional named <b>list</b> used for substituting symbols with other strings.
code	If <b>TRUE</b> , the deparsed code of the expression is echoed.
output	If <b>TRUE</b> , the output of each evaluated subexpression is echoed.
...	Additional arguments passed to <b>sourceTo</b> which in turn passes arguments to <b>source()</b> .
max.deparse.length	A positive <b>integer</b> specifying the maximum length of a deparsed expression, before truncating it.
trim	If <b>TRUE</b> , the captured rows are trimmed.
newline	If <b>TRUE</b> and collapse is non- <b>NULL</b> , a newline is appended at the end.
collapse	A <b>character</b> string used for collapsing the captured rows. If <b>NULL</b> , the rows are not collapsed.
substitute	(to be deprecated) use replace instead.
envir	The <b>environment</b> in which the expression is evaluated.

**Value**

Returns a **character** string class 'CapturedEvaluation'.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally, **eval()** is used to evaluate the expression.

**Examples**

```
print(withCapture({
  n <- 3;
  n;

  for (kk in 1:3) {
    printf("Iteration #%d\n", kk);
  }

  print(Sys.time());
```

```
    type <- "horse";
    type;
  )))

## > n <- 3
## > n
## [1] 3
## > for (kk in 1:3) {
## +   printf("Iteration #%d\n", kk)
## + }
## Iteration #1
## Iteration #2
## Iteration #3
## > print(Sys.time())
## [1] "2011-11-06 11:06:32 PST"
## > type <- "horse"
## > type
## [1] "horse"

# Automatic "variable" substitute
# (disable with relabel=NULL)
a <- 2
b <- "Hello world!"

print(withCapture({
  x <- .a.
  s <- .b.
  x
  s
}))

## > x <- 2
## > s <- "Hello world!"
## > x
## [1] 2
## > s
## [1] "Hello world!"
```

---

withLocale

*Evaluate an R expression with locale set temporarily*

---

### Description

Evaluate an R expression with locale set temporarily.

### Usage

```
withLocale(expr, category, locale, ..., substitute=TRUE, envir=parent.frame())
```

**Arguments**

expr	The R expression to be evaluated.
category	A <a href="#">character</a> string specifying the category to use.
locale	<a href="#">character vector</a> specifying the locale to used. The first successfully set one will be used.
...	Not used.
substitute	If <a href="#">TRUE</a> , argument expr is <a href="#">substitute()</a> :ed, otherwise not.
envir	The <a href="#">environment</a> in which the expression should be evaluated.

**Value**

Returns the results of the expression evaluated.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally, [eval\(\)](#) is used to evaluate the expression. and [Sys.setlocale\(\)](#) to set locale.

**Examples**

```
# Vector
cat("Original vector:\n")
x <- c(letters[1:8], LETTERS[1:8])
print(x)

cat("Sorting with 'C' locale:\n")
y1 <- withLocale(sort(x), "LC_COLLATE", "C")
print(y1)

cat("Sorting with an 'English' locale:\n")
y2 <- withLocale(sort(x), "LC_COLLATE", c("en_US", "en_US.UTF8", "English_United States.1252"))
print(y2)
```

---

withOptions

*Evaluate an R expression with options set temporarily*


---

**Description**

Evaluate an R expression with options set temporarily.

**Usage**

```
withOptions(expr, ..., args=list(), substitute=TRUE, envir=parent.frame())
```

## Arguments

expr	The R expression to be evaluated.
...	Named options to be used.
args	(optional) Additional named options specified as a named <a href="#">list</a> .
substitute	If <code>TRUE</code> , argument <code>expr</code> is <code>substitute()</code> :ed, otherwise not.
envir	The <a href="#">environment</a> in which the expression should be evaluated.

## Details

Upon exit (also on errors), this function will reset *all* options to the state of options available upon entry. This means any options *modified* but also those *added* when evaluating `expr` will also be undone upon exit.

## Value

Returns the results of the expression evaluated.

## Author(s)

Henrik Bengtsson

## See Also

Internally, `eval()` is used to evaluate the expression. and `options()` to set options.

## Examples

```
print(pi)

# Same, i.e. using default
withOptions({
  print(pi)
})

# Printing with two digits
withOptions({
  print(pi)
}, digits=2)

# Printing with two digits then with three more
withOptions({
  print(pi)
  withOptions({
    print(pi)
  }, digits=getOption("digits")+3)
}, digits=2)

# Still printing with the default
print(pi)
```

---

`withRepos`*Evaluate an R expression with repositories set temporarily*

---

**Description**

Evaluate an R expression with repositories set temporarily.

**Usage**

```
withRepos(expr, repos="[[mainstream]]", ..., substitute=TRUE, envir=parent.frame())
```

**Arguments**

<code>expr</code>	The R expression to be evaluated.
<code>repos</code>	A <a href="#">character vector</a> of repositories to use.
<code>...</code>	Additional arguments passed to <a href="#">useRepos()</a> .
<code>substitute</code>	If <code>TRUE</code> , argument <code>expr</code> is <a href="#">substitute()</a> :ed, otherwise not.
<code>envir</code>	The <a href="#">environment</a> in which the expression should be evaluated.

**Value**

Returns the results of the expression evaluated.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally, [eval\(\)](#) is used to evaluate the expression. See also [options\(\)](#) and [install.packages](#).

**Examples**

```
## Not run:
# Install from BioC related repositories only
withRepos(install.packages("edgeR"), repos="[[BioC]]")

# Install from CRAN or BioC related repositories only
withRepos(install.packages("edgeR"), repos=c("CRAN", "[[BioC]]"))

# Install from mainstream repositories only (same as previous)
withRepos(install.packages("edgeR"), repos="[[mainstream]]")

# Install from R-Forge and mainstream repositories only
withRepos(install.packages("R.utils"), repos="[[R-Forge]]")

# Update only CRAN packages
withRepos(update.packages(ask=FALSE), repos="[[CRAN]]")
```

```
# Update only Bioconductor packages
withRepos(update.packages(ask=FALSE), repos="[[BioC]]")

## End(Not run)
```

---

withSeed

*Evaluate an R expression with a temporarily set random set*

---

### Description

Evaluate an R expression with a temporarily set random set.

### Usage

```
withSeed(expr, seed, ..., substitute=TRUE, envir=parent.frame())
```

### Arguments

expr	The R expression to be evaluated.
seed, ...	Arguments passed to <a href="#">set.seed()</a> .
substitute	If <code>TRUE</code> , argument expr is <a href="#">substitute()</a> :ed, otherwise not.
envir	The <a href="#">environment</a> in which the expression should be evaluated.

### Details

Upon exit (also on errors), this function will restore `.Random.seed` in the global environment to the value it had upon entry. If it did not exist, it will be removed.

### Value

Returns the results of the expression evaluated.

### Author(s)

Henrik Bengtsson

### See Also

Internally, [set.seed\(\)](#) is used to set the random seed.

**Examples**

```

# Generate a random number
y0 <- runif(1)
print(y0)

# Generate a random number using the same seed over and over
yp <- NULL
for (ii in 1:10) {
  y <- withSeed({
    runif(1)
  }, seed=0x42)
  print(y)
  # Assert identical
  if (!is.null(yp)) stopifnot(identical(y, yp))
  yp <- y
}

# Generate a random number
y <- runif(1)
print(y)

```

withSink

*Evaluate an R expression while temporarily diverting output***Description**

Evaluate an R expression while temporarily diverting output.

**Usage**

```
withSink(expr, file, append=FALSE, type=c("output", "message"), substitute=TRUE,
  envir=parent.frame())
```

**Arguments**

expr	The R expression to be evaluated.
file	A writable <a href="#">connection</a> or a <a href="#">character</a> string naming the file to write to.
append	If <a href="#">TRUE</a> , the diverted output is appended to the file, otherwise not.
type	A <a href="#">character</a> string specifying whether to divert output sent to the standard output or the standard error. See <a href="#">sink()</a> for details.
substitute	If <a href="#">TRUE</a> , argument expr is <a href="#">substitute()</a> :ed, otherwise not.
envir	The <a href="#">environment</a> in which the expression should be evaluated.

**Details**

Upon exit (also on errors), this function will close the requested "sink". If additional sinks (of any type) were also opened during the evaluation, those will also be closed with a warning.



**Value**

Returns the results of the expression evaluated.

**Author(s)**

Henrik Bengtsson

**See Also**

Internally, `sink()` is used to divert any output.

**Examples**

```
# Divert standard output
pathname <- tempfile(fileext=".output.txt")
res <- withSink(file=pathname, {
  print(letters)
})
mcat(readLines(pathname), sep="\n")

# Divert standard error/messages
pathname <- tempfile(fileext=".message.txt")
res <- withSink(file=pathname, type="message", {
  mprint(LETTERS)
})
mcat(readLines(pathname), sep="\n")
```

---

withTimeout

*Evaluate an R expression and interrupts it if it takes too long*

---

**Description**

Evaluate an R expression and interrupts it if it takes too long.

**Usage**

```
withTimeout(expr, substitute=TRUE, envir=parent.frame(), timeout, cpu=timeout,
  elapsed=timeout, onTimeout=c("error", "warning", "silent"), ...)
```

**Arguments**

<code>expr</code>	The R expression to be evaluated.
<code>substitute</code>	If <code>TRUE</code> , argument <code>expr</code> is <code>substitute()</code> :ed, otherwise not.
<code>envir</code>	The <code>environment</code> in which the expression should be evaluated.

timeout, cpu, elapsed	A <a href="#">numeric</a> specifying the maximum number of seconds the expression is allowed to run before being interrupted by the timeout. The <code>cpu</code> and <code>elapsed</code> arguments can be used to specify whether time should be measured in CPU time or in wall time.
onTimeout	A <a href="#">character</a> specifying what action to take if a timeout event occurs.
...	Not used.

### Details

This method utilizes `setTimeLimit()` by first setting the timeout limits, then evaluating the expression that may or may not timeout. The method is guaranteed to reset the timeout limits to be infinitely long upon exiting, regardless whether it returns normally or preemptively due to a timeout or an error.

### Value

Returns the results of the expression evaluated. If timed out, `NULL` is returned if `onTimeout` was "warning" or "silent". If "error" a `TimeoutException` is thrown.

### Non-supported cases

In order to understand when this function works and when it does not, it is useful to know that it utilizes R's built-in time-out mechanism, which sets the limits on what is possible and not. From `setTimeLimit()`, we learn that:

*"Time limits are checked whenever a user interrupt could occur. This will happen frequently in R code and during `Sys.sleep(*)`, but only at points in compiled C and Fortran code identified by the code author."*

More precisely, if a function is implemented in native code (e.g. C) and the developer of that function does not check for user interrupts, then you cannot interrupt that function neither via a user interrupt (e.g. Ctrl-C) *nor via the built-in time out mechanism*. To change this, you need to contact the developer of that piece of code and ask them to check for R user interrupts in their native code.

Furthermore, it is not possible to interrupt/break out of a "readline" prompt (e.g. `readline()` and `readLines()`) using timeouts; the timeout exception will not be thrown until after the user completes the prompt (i.e. after pressing ENTER).

System calls via `system()` and `system2()` cannot be timed out via the above mechanisms. However, in R ( $\geq 3.5.0$ ) these functions have argument `timeout` providing their own independent timeout mechanism.

Other examples of calls that do *not* support timeout are "atomic" calls that may take very long such as large object allocation and `rnorm(n)` where `n` is very large.

(\*) Note that on Unix and macOS, `Sys.sleep(time)` will signal a timeout error only *after* `time` seconds passed, regardless of `timeout limit (< time)`.

### Author(s)

Henrik Bengtsson

## References

[1] R help thread 'Time out for a R Function' on 2010-12-06. <http://www.mail-archive.com/r-help@r-project.org/msg119344.html>

## See Also

Internally, `eval()` is used to evaluate the expression and `setTimeLimit()` is used to control for timeout events.

## Examples

```
# -----
# Function that takes "a long" time to run
# -----
foo <- function() {
  print("Tic")
  for (kk in 1:100) {
    print(kk)
    Sys.sleep(0.1)
  }
  print("Tac")
}

# -----
# Evaluate code, if it takes too long, generate
# a timeout by throwing a TimeoutException.
# -----
res <- NULL
tryCatch({
  res <- withTimeout({
    foo()
  }, timeout = 1.08)
}, TimeoutException = function(ex) {
  message("Timeout. Skipping.")
})

# -----
# Evaluate code, if it takes too long, generate
# a timeout returning NULL and generate a warning.
# -----
res <- withTimeout({
  foo()
}, timeout = 1.08, onTimeout = "warning")

# The same using an expression object
expr <- quote(foo())
res <- withTimeout(expr, substitute = FALSE,
  timeout = 1.08, onTimeout = "warning")
```

```

# -----
# Evaluate code, if it takes too long, generate
# a timeout, and return silently NULL.
# -----
res <- withTimeout({
  foo()
}, timeout = 1.08, onTimeout = "silent")

```

---

wrap.array

*Reshape an array or a matrix by permuting and/or joining dimensions*


---

### Description

Reshape an array or a matrix by permuting and/or joining dimensions.

A useful application of this is to reshape a multidimensional [array](#) to a [matrix](#), which then can be saved to file using for instance `write.table()`.

### Usage

```

## S3 method for class 'array'
wrap(x, map=list(NA), sep=".", ...)

```

### Arguments

x	An <a href="#">array</a> or a <a href="#">matrix</a> .
map	A <a href="#">list</a> of length equal to the number of dimensions in the reshaped array. Each element should be an <a href="#">integer vector</a> s specifying the dimensions to be joined in corresponding new dimension. One element may equal <code>NA</code> to indicate that that dimension should be a join of all non-specified (remaining) dimensions. Default is to wrap everything into a <a href="#">vector</a> .
sep	A <a href="#">character</a> pasting joined dimension names.
...	Not used.

### Details

If the indices in `unlist(map)` is in a non-increasing order, [aperm\(\)](#) will be called, which requires reshuffling of array elements in memory. In all other cases, the reshaping of the array does not require this, but only fast modifications of attributes `dim` and `dimnames`.

### Value

Returns an [array](#) of `length(map)` dimensions, where the first dimension is of size `prod(map[[1]])`, the second `prod(map[[2]])`, and so on.

**Author(s)**

Henrik Bengtsson

**See Also**[\\*unwrap\(\)](#). See [aperm\(\)](#).**Examples**

```

# Create a 3x2x3 array
dim <- c(3,2,3)
ndim <- length(dim)
dimnames <- list()
for (kk in 1:ndim)
  dimnames[[kk]] <- sprintf("%s%d", letters[kk], 1:dim[kk])
x <- 1:prod(dim)
x <- array(x, dim=dim, dimnames=dimnames)

cat("Array 'x':\n")
print(x)

cat("\nReshape 'x' to its identity:\n")
y <- wrap(x, map=list(1, 2, 3))
print(y)
# Assert correctness of reshaping
stopifnot(identical(y, x))

cat("\nReshape 'x' by swapping dimensions 2 and 3, i.e. aperm(x, perm=c(1,3,2)):\n")
y <- wrap(x, map=list(1, 3, 2))
print(y)
# Assert correctness of reshaping
stopifnot(identical(y, aperm(x, perm=c(1,3,2))))

cat("\nWrap 'x' to a matrix 'y' by keeping dimension 1 and joining the others:\n")
y <- wrap(x, map=list(1, NA))
print(y)
# Assert correctness of reshaping
for (aa in dimnames(x)[[1]]) {
  for (bb in dimnames(x)[[2]]) {
    for (cc in dimnames(x)[[3]]) {
      tt <- paste(bb, cc, sep=".")
      stopifnot(identical(y[aa,tt], x[aa,bb,cc]))
    }
  }
}

```

```

cat("\nUnwrap matrix 'y' back to array 'x':\n")
z <- unwrap(y)
print(z)
stopifnot(identical(z,x))

cat("\nWrap a matrix 'y' to a vector and back again:\n")
x <- matrix(1:8, nrow=2, dimnames=list(letters[1:2], 1:4))
y <- wrap(x)
z <- unwrap(y)
print(z)
stopifnot(identical(z,x))

cat("\nWrap and unwrap a randomly sized and shaped array 'x2':\n")
maxdim <- 5
dim <- sample(1:maxdim, size=sample(2:maxdim))
ndim <- length(dim)
dimnames <- list()
for (kk in 1:ndim)
  dimnames[[kk]] <- sprintf("%s%d", letters[kk], 1:dim[kk])
x2 <- 1:prod(dim)
x2 <- array(x2, dim=dim, dimnames=dimnames)

cat("\nArray 'x2':\n")
print(x2)

# Number of dimensions of wrapped array
ndim2 <- sample(1:(ndim-1), size=1)

# Create a random map for joining dimensions
splits <- NULL;
if (ndim > 2)
  splits <- sort(sample(2:(ndim-1), size=ndim2-1))
splits <- c(0, splits, ndim);
map <- list();
for (kk in 1:ndim2)
  map[[kk]] <- (splits[kk]+1):splits[kk+1];

cat("\nRandom 'map':\n")
print(map)

cat("\nArray 'y2':\n")
y2 <- wrap(x2, map=map)
print(y2)

cat("\nArray 'x2':\n")
z2 <- unwrap(y2)
print(z2)

stopifnot(identical(z2,x2))

```

---

writeBinFragments	<i>Writes binary data to disjoint sections of a connection or a file</i>
-------------------	--

---

### Description

Writes binary data to disjoint sections of a connection or a file.

### Usage

```
## Default S3 method:  
writeBinFragments(con, object, idxs, size=NA, ...)
```

### Arguments

con	A <a href="#">connection</a> or the pathname of an existing file.
object	A <a href="#">vector</a> of objects to be written.
idxs	A <a href="#">vector</a> of (non-duplicated) indices or a Nx2 <a href="#">matrix</a> of N from-to index intervals specifying the elements to be read. Positions are always relative to the start of the file/connection.
size	The size of the data type to be read. If <a href="#">NA</a> , the natural size of the data type is used.
...	Additional arguments passed to <a href="#">writeBin()</a> .

### Value

Returns nothing.

### Author(s)

Henrik Bengtsson

### See Also

[readBinFragments\(\)](#).

### Examples

```
## Not run: # See example(readBinFragments.connection)
```

---

```
writeDataFrame.data.frame
```

*Writes a data.frame to tabular text file*

---

### Description

Writes a data.frame to tabular text file with an optional header.

### Usage

```
## S3 method for class 'data.frame'
writeDataFrame(data, file, path=NULL, sep="\t", quote=FALSE, row.names=FALSE,
  col.names=!append, ..., header=list(), createdBy=NULL,
  createdOn=format(Sys.time(), format = "%Y-%m-%d %H:%M:%S %Z"),
  nbrOfRows=nrow(data), headerPrefix="# ", headerSep=": ", append=FALSE, overwrite=FALSE)
```

### Arguments

data	A <a href="#">data.frame</a> .
file	A <a href="#">connection</a> or a filename to write to.
path	The directory where the file will be written.
sep, quote, row.names, col.names, ...	Additional arguments passed to <a href="#">write.table</a> .
header	An optional named <a href="#">list</a> of header rows to be written at the beginning of the file. If <code>NULL</code> , no header will be written.
createdBy, createdOn, nbrOfRows	If non- <code>NULL</code> , common header rows to be added to the header.
headerPrefix	A <a href="#">character</a> string specifying the prefix of each header row.
headerSep	A <a href="#">character</a> string specifying the character separating the header name and header values.
append	If <code>TRUE</code> , the output is appended to an existing file.
overwrite	If <code>TRUE</code> , an existing file is overwritten.

### Value

Returns (invisibly) the pathname to the file written (or the [connection](#) written to).

### Author(s)

Henrik Bengtsson

### See Also

[write.table](#), [readTable\(\)](#).



# Index

## \*Topic **IO**

- captureOutput, 14
- compressPDF, 20
- createFileAtomically, 23
- createLink, 25
- createWindowsShortcut, 26
- dimNA< -, 29
- displayCode, 30
- fileAccess, 38
- filePath, 40
- findSourceTraceback, 44
- getAbsolutePath, 47
- getParent, 48
- getRelativePath, 49
- hasUrlProtocol, 54
- isAbsolutePath, 62
- isDirectory, 63
- isFile, 64
- isOpen.character, 65
- isUrl, 69
- lastModified, 73
- LComments, 74
- listDirectory, 75
- loadObject, 76
- makedirs, 79
- mpager, 81
- NullVerbose, 82
- popBackupFile, 89
- popTemporaryFile, 90
- pushBackupFile, 93
- pushTemporaryFile, 95
- readBinFragments, 98
- readTable, 101
- readTableIndex, 103
- readWindowsShortcut, 104
- removeDirectory, 105
- resample, 106
- saveObject, 107
- Settings, 110

- shell.exec2, 113
- SmartComments, 114
- sourceDirectory, 116
- sourceTo, 117
- systemR, 123
- TextStatusBar, 124
- touchFile, 127
- toUrl, 129
- useRepos, 130
- VComments, 131
- Verbose, 133
- withLocale, 139
- withOptions, 140
- withRepos, 142
- withSeed, 143
- withSink, 144
- withTimeout, 145
- writeBinFragments, 151
- writeDataFrame.data.frame, 152

## \*Topic **attribute**

- intervalsToSeq.matrix, 61
- seqToHumanReadable, 108
- seqToIntervals, 109

## \*Topic **character**

- as.character.binmode, 8
- intToBin, 61

## \*Topic **classes**

- Arguments, 7
- Assert, 9
- FileProgressBar, 42
- GString, 50
- Java, 71
- LComments, 74
- NullVerbose, 82
- Options, 85
- ProgressBar, 92
- Settings, 110
- SmartComments, 114
- System, 122

- TextStatusBar, 124
  - TimeoutException, 126
  - VComments, 131
  - Verbose, 133
- \*Topic **device**
  - env, 35
- \*Topic **error**
  - TimeoutException, 126
- \*Topic **file**
  - compressFile, 18
  - compressPDF, 20
  - copyDirectory, 21
  - createLink, 25
  - createWindowsShortcut, 26
  - dimNA< -, 29
  - displayCode, 30
  - downloadFile.character, 32
  - installPackages, 59
  - mpager, 81
  - nullfile, 81
  - readWindowsShortcut, 104
  - shell.exec2, 113
  - touchFile, 127
- \*Topic **logic**
  - isZero, 70
- \*Topic **manip**
  - as.character.binmode, 8
  - dataFrame, 28
  - insert, 58
  - intToBin, 61
- \*Topic **methods**
  - attachLocally.list, 10
  - callHooks.function, 13
  - downloadFile.character, 32
  - extract.array, 36
  - inAnyInterval.numeric, 57
  - intervalsToSeq.matrix, 61
  - isEof.connection, 63
  - isOpen.character, 65
  - mapToIntervals.numeric, 77
  - mergeIntervals.numeric, 78
  - TimeoutException, 126
  - unwrap.array, 129
  - wrap.array, 148
  - writeDataFrame.data.frame, 152
- \*Topic **package**
  - isPackageInstalled, 66
  - isPackageLoaded, 66
- R.utils-package, 4
- \*Topic **programming**
  - addFinalizerToLast, 6
  - Arguments, 7
  - as.character.binmode, 8
  - attachLocally.list, 10
  - callHooks, 11
  - callHooks.function, 13
  - capitalize, 13
  - captureOutput, 14
  - cmdArgs, 16
  - colClasses, 17
  - compressFile, 18
  - countLines, 22
  - createFileAtomically, 23
  - detachPackage, 29
  - doCall, 31
  - downloadFile.character, 32
  - egsub, 34
  - extract.array, 36
  - fileAccess, 38
  - finalizeSession, 43
  - findSourceTraceback, 44
  - getAbsolutePath, 47
  - getParent, 48
  - getRelativePath, 49
  - hasUrlProtocol, 54
  - hpaste, 55
  - inAnyInterval.numeric, 57
  - intToBin, 61
  - isAbsolutePath, 62
  - isDirectory, 63
  - isFile, 64
  - isUrl, 69
  - lastModified, 73
  - LComments, 74
  - listDirectory, 75
  - loadObject, 76
  - mapToIntervals.numeric, 77
  - mergeIntervals.numeric, 78
  - mkdirs, 79
  - mpager, 81
  - nullfile, 81
  - NullVerbose, 82
  - onGarbageCollect, 83
  - onSessionExit, 84
  - Options, 85
  - patchCode, 87

- popBackupFile, 89
- popTemporaryFile, 90
- pushBackupFile, 93
- pushTemporaryFile, 95
- readRdHelp, 100
- removeDirectory, 105
- resample, 106
- saveObject, 107
- setOption, 110
- Settings, 110
- SmartComments, 114
- sourceDirectory, 116
- sourceTo, 117
- splitByPattern, 119
- systemR, 123
- TextStatusBar, 124
- TimeoutException, 126
- touchFile, 127
- toUrl, 129
- unwrap.array, 129
- useRepos, 130
- VComments, 131
- Verbose, 133
- withLocale, 139
- withOptions, 140
- withRepos, 142
- withSeed, 143
- withSink, 144
- withTimeout, 145
- wrap.array, 148
- \*Topic utilities**
  - attachLocally.list, 10
  - createFileAtomically, 23
  - dataFrame, 28
  - egsub, 34
  - env, 35
  - inAnyInterval.numeric, 57
  - isOpen.character, 65
  - isPackageInstalled, 66
  - isPackageLoaded, 66
  - mapToIntervals.numeric, 77
  - mergeIntervals.numeric, 78
  - mout, 80
  - patchCode, 87
  - popBackupFile, 89
  - popTemporaryFile, 90
  - printf, 91
  - pushBackupFile, 93
  - pushTemporaryFile, 95
    - withCapture, 137
  - \*intervalsToSeq, 109
  - \*mapToIntervals, 57
  - \*unwrap, 149
  - \*wrap, 130
  - \*writeRaw, 136
  - .Last, 85
  - .Machine, 71
  - .Random.seed, 143
  - .Rprofile, 114
- addFinalizerToLast, 6
- all.equal, 71
- aperm(), 148, 149
- append, 58
- Arguments, 4, 7
- array, 37, 130, 148
- as.character, 50, 86, 92, 134
- as.character.binmode, 8
- as.double, 134
- as.list, 86
- as.logical, 134
- as.numeric, 16
- asByte, 72
- asInt, 72
- asLong, 72
- Assert, 4, 9
- asShort, 72
- attach, 10
- attachLocally, 10
- attachLocally(attachLocally.list), 10
- attachLocally.list, 10
- browseURL, 114
- bunzip2(compressFile), 18
- bzip2(compressFile), 18
- callHooks, 11, 13
- callHooks.function, 13
- callHooks.list, 11, 12
- callHooks.list(callHooks.function), 13
- capitalize, 13
- capture, 134
- capture.output, 15
- captureOutput, 14
- cat, 80, 91, 134
- ccat(mout), 80

- character, [8](#), [10](#), [11](#), [14–17](#), [19](#), [21](#), [22](#), [25](#),  
[28–34](#), [38](#), [41](#), [44–50](#), [53–55](#), [60](#),  
[62–67](#), [70](#), [74](#), [76](#), [79](#), [81](#), [82](#), [84](#), [88](#),  
[89](#), [91](#), [94](#), [97](#), [98](#), [101–105](#), [108](#),  
[110](#), [111](#), [113](#), [114](#), [117](#), [119](#), [123](#),  
[124](#), [128–131](#), [134](#), [138](#), [140](#), [142](#),  
[144](#), [146](#), [148](#), [152](#)
- check, [9](#)
- cleanup, [43](#)
- cmdArg (cmdArgs), [16](#)
- cmdArgs, [16](#)
- ormsg (mout), [80](#)
- colClasses, [17](#), [102](#)
- commandArgs, [16](#), [17](#)
- compactPDF, [21](#)
- Comparison, [71](#)
- compile, [115](#)
- compressFile, [18](#)
- compressPDF, [20](#)
- connection, [15](#), [22](#), [30](#), [45](#), [53](#), [64](#), [65](#), [76](#), [82](#),  
[91](#), [98](#), [102](#), [104](#), [107](#), [117](#), [134](#), [144](#),  
[151](#), [152](#)
- connections, [19](#), [65](#)
- convertComment, [115](#), [133](#)
- copyDirectory, [21](#)
- copyFile, [22](#)
- countLines, [22](#)
- cout (mout), [80](#)
- cprint (mout), [80](#)
- cprintf (mout), [80](#)
- createFileAtomically, [23](#)
- createLink, [25](#)
- createWindowsShortcut, [26](#), [26](#), [105](#)
- cshow (mout), [80](#)
- cstr (mout), [80](#)
- currentTimeMillis, [122](#)
- cut, [78](#)
- data.frame, [10](#), [28](#), [102](#), [152](#)
- dataFrame, [28](#)
- decapitalize (capitalize), [13](#)
- decompressFile (compressFile), [18](#)
- detach, [29](#)
- detachPackage, [29](#)
- digest, [107](#)
- dim, [30](#)
- dimNA< -, [29](#)
- dimNA<- (dimNA< -), [29](#)
- dir.create, [79](#)
- displayCode, [30](#)
- do.call, [32](#)
- doCall, [31](#)
- download.file, [33](#)
- downloadFile (downloadFile.character),  
[32](#)
- downloadFile.character, [32](#)
- duplicated, [67](#), [68](#)
- egsub, [34](#)
- enter, [135](#)
- env, [35](#)
- environment, [10](#), [15](#), [32](#), [34](#), [35](#), [45](#), [53](#), [116](#),  
[117](#), [138](#), [140–145](#)
- equals, [86](#), [135](#)
- eval, [15](#), [138](#), [140–142](#), [147](#)
- evalCapture (withCapture), [137](#)
- evalq, [35](#), [36](#)
- evaluate, [50](#), [135](#)
- evalWithTimeout (withTimeout), [145](#)
- Exception, [126](#), [127](#)
- exit, [135](#)
- expression, [34](#), [35](#)
- extract.array, [36](#)
- extract.default (extract.array), [36](#)
- extract.matrix (extract.array), [36](#)
- FALSE, [6](#), [10](#), [21](#), [46](#), [58](#), [62–64](#), [70](#), [76](#), [79](#), [89](#),  
[90](#), [94–96](#), [101](#), [102](#), [105](#), [117](#), [136](#)
- file.access, [38](#), [39](#)
- file.copy, [22](#)
- file.info, [63](#), [65](#), [74](#), [128](#)
- file.path, [40](#), [41](#)
- file.remove, [105](#)
- file.show, [31](#), [81](#)
- file.symlink, [26](#)
- file\_test, [63](#), [65](#)
- fileAccess, [38](#)
- filePath, [40](#), [105](#)
- FileProgressBar, [5](#), [42](#), [92](#)
- finalizeSession, [43](#), [85](#)
- findGhostscript, [122](#)
- findGraphicsDevice, [122](#)
- findInterval, [78](#)
- findSettings, [111](#)
- findSourceTraceback, [44](#)
- flush, [124](#)
- function, [13](#), [19](#), [23](#), [32](#), [34](#), [84](#), [130](#)

- gcat, [45](#), [51](#), [54](#)
- gcDLLs, [46](#)
- get, [34](#)
- getAbsolutePath, [47](#), [49](#)
- getBarString, [92](#)
- getBuiltinDate, [51](#)
- getBuiltinDatetime, [51](#)
- getBuiltinHostname, [51](#)
- getBuiltinOs, [51](#)
- getBuiltinPid, [51](#)
- getBuiltinRhome, [51](#)
- getBuiltinRversion, [51](#)
- getBuiltinTime, [51](#)
- getBuiltinUsername, [51](#)
- getCharacters, [7](#)
- getDoubles, [7](#)
- getEnvironment, [7](#)
- getFilename, [7](#)
- getHostname, [122](#)
- getIndices, [7](#)
- getInstanceOf, [7](#)
- getIntegers, [7](#)
- getLabel, [125](#)
- getLeaves, [86](#)
- getLoadedDLLs, [47](#)
- getLoadedPathname, [111](#)
- getLogicals, [7](#)
- getMessage, [127](#)
- getNumerics, [7](#)
- getOption, [86](#), [110](#)
- getParent, [48](#)
- getRaw, [51](#)
- getReadablePathname, [7](#)
- getReadablePathnames, [7](#)
- getRegularExpression, [7](#)
- getRelativePath, [49](#)
- getThreshold, [135](#)
- getTimestampFormat, [135](#)
- getUsername, [122](#)
- getVariableValue, [51](#)
- getVector, [7](#)
- getVerbose, [7](#)
- getWritablePathname, [8](#)
- GString, [4](#), [45](#), [50](#), [53](#), [134](#)
- gstring, [45](#), [46](#), [51](#), [53](#)
- gsub, [34](#)
- gunzip (compressFile), [18](#)
- gzip (compressFile), [18](#)
- hasOption, [86](#)
- hasUrlProtocol, [54](#)
- header, [135](#)
- help, [101](#)
- hpaste, [55](#)
- inAnyInterval, [78](#), [79](#)
- inAnyInterval.numeric, [57](#)
- increase, [92](#)
- Inf, [55](#)
- inheritsFrom, [9](#)
- insert, [58](#)
- install.packages, [60](#), [142](#)
- installPackages, [59](#)
- integer, [23](#), [28](#), [29](#), [33](#), [37](#), [38](#), [48](#), [55](#), [61](#), [77](#),  
[79](#), [103](#), [108](#), [109](#), [120](#), [138](#), [148](#)
- intervalsToSeq.matrix, [61](#)
- intToBin, [8](#), [61](#)
- intToHex (intToBin), [61](#)
- intToOct (intToBin), [61](#)
- isAbsolutePath, [48](#), [49](#), [62](#)
- isBzipped (compressFile), [18](#)
- isCompressedFile (compressFile), [18](#)
- isDirectory, [63](#), [65](#)
- isDone, [92](#)
- isEof.connection, [63](#)
- isFile, [63](#), [64](#)
- isGzipped (compressFile), [18](#)
- isMatrix, [9](#)
- isModified, [111](#)
- isOn, [83](#), [135](#)
- isOpen.character, [65](#)
- isPackageInstalled, [66](#), [67](#)
- isPackageLoaded, [66](#), [66](#)
- isReplicated, [67](#), [69](#)
- isScalar, [9](#)
- isSingle, [68](#), [69](#)
- isUrl, [69](#)
- isVector, [9](#)
- isVisible, [83](#), [135](#)
- isZero, [70](#)
- Java, [4](#), [71](#)
- lastModified, [73](#)
- layout, [121](#)
- LComments, [5](#), [74](#), [114](#), [131](#)
- less, [135](#)
- library, [88](#)

- list, [10](#), [13](#), [16](#), [32](#), [37](#), [58](#), [86](#), [101](#), [104](#), [130](#), [131](#), [138](#), [141](#), [148](#), [152](#)
- list.files, [76](#)
- listDirectory, [75](#)
- load, [77](#)
- loadAnywhere, [111](#)
- loadObject, [76](#), [107](#)
- loadToEnv, [77](#)
- logical, [19](#), [23](#), [29](#), [33](#), [54](#), [57](#), [64–67](#), [69](#), [70](#), [80](#), [89](#), [90](#), [94](#), [95](#), [98](#), [102](#), [103](#), [116](#), [123](#), [134](#)
- mapToIntervals, [57](#)
- mapToIntervals.numeric, [77](#)
- match, [78](#), [79](#)
- matrix, [37](#), [61](#), [77](#), [78](#), [98](#), [109](#), [121](#), [130](#), [148](#), [151](#)
- mcat (mout), [80](#)
- mergeIntervals.numeric, [78](#)
- message, [80](#)
- mkdirs, [79](#)
- mode, [98](#)
- more, [135](#)
- mout, [80](#)
- mpager, [81](#)
- mprint (mout), [80](#)
- mprintf (mout), [80](#)
- mshow (mout), [80](#)
- mstr (mout), [80](#)
- mtext, [120](#)
- MultiVerbose, [134](#)
- NA, [16](#), [29](#), [30](#), [40](#), [77](#), [98](#), [136](#), [148](#), [151](#)
- names, [86](#)
- nbrOfOptions, [86](#)
- new.env, [35](#), [36](#)
- newline, [125](#), [135](#)
- NULL, [10](#), [13](#), [15](#), [16](#), [25](#), [30](#), [33](#), [37](#), [48](#), [49](#), [55](#), [67](#), [88](#), [107](#), [117](#), [131](#), [138](#), [146](#), [152](#)
- nullcon (nullfile), [81](#)
- nullfile, [81](#)
- NullVerbose, [82](#), [134](#), [136](#)
- numeric, [16](#), [30](#), [62](#), [76–78](#), [120](#), [124](#), [134](#), [146](#)
- Object, [7](#), [9](#), [42](#), [71](#), [74](#), [82](#), [85](#), [86](#), [92](#), [110](#), [114](#), [122](#), [124](#), [126](#), [131](#), [134](#)
- octmode, [8](#)
- off, [135](#)
- on, [135](#)
- onGarbageCollect, [83](#)
- onSessionExit, [6](#), [43](#), [44](#), [84](#), [111](#)
- openBrowser, [122](#)
- Options, [4](#), [85](#), [110](#), [111](#)
- options, [86](#), [88](#), [141](#), [142](#)
- parse, [51](#), [115](#), [117](#)
- parseDebian, [122](#)
- parseRepos (useRepos), [130](#)
- paste, [55](#)
- patchCode, [87](#)
- popBackupFile, [24](#), [89](#), [93](#), [94](#)
- popMessage, [125](#)
- popTemporaryFile, [23](#), [24](#), [90](#), [95](#), [96](#)
- print, [51](#), [80](#), [135](#)
- printf, [80](#), [91](#), [135](#)
- ProgressBar, [5](#), [42](#), [92](#)
- promptAndSave, [111](#)
- pushBackupFile, [24](#), [90](#), [93](#)
- pushState, [135](#)
- pushTemporaryFile, [23](#), [24](#), [90](#), [95](#)
- queryRCmdCheck, [97](#)
- R.home, [123](#)
- R.utils (R.utils-package), [4](#)
- R.utils-package, [4](#)
- raw, [15](#)
- read.table, [18](#), [28](#), [101](#), [102](#)
- readBin, [98](#)
- readBinFragments, [98](#), [151](#)
- readByte, [72](#)
- readInt, [72](#)
- readline, [146](#)
- readLines, [146](#)
- readRdHelp, [100](#)
- readShort, [72](#)
- readTable, [101](#), [103](#), [152](#)
- readTableIndex, [102](#), [103](#)
- readUTF, [72](#)
- readWindowsShellLink, [41](#)
- readWindowsShortcut, [27](#), [41](#), [104](#)
- removeDirectory, [105](#)
- replicates (isReplicated), [67](#)
- require, [66](#)
- resample, [106](#)
- reset, [92](#), [115](#), [133](#)
- rle, [109](#)
- ruler, [135](#)

- sample, [106](#)
- sample.int, [106](#)
- saveAnywhere, [111](#)
- saveObject, [76](#), [77](#), [107](#)
- saveRDS, [77](#), [107](#)
- seek, [64](#)
- seqToHumanReadable, [108](#)
- seqToIntervals, [61](#), [108](#), [109](#)
- set.seed, [143](#)
- setDefaultLevel, [135](#)
- setLabel, [125](#)
- setLabels, [125](#)
- setMaxValue, [92](#)
- setOption, [86](#), [110](#)
- setProgress, [92](#)
- setStepLength, [92](#)
- setThreshold, [135](#)
- setTicks, [92](#)
- setTimeLimit, [146](#), [147](#)
- setTimestampFormat, [135](#)
- Settings, [5](#), [85](#), [110](#)
- setValue, [92](#)
- shell.exec2, [113](#)
- show, [80](#)
- showConnections, [65](#)
- singles (isSingle), [69](#)
- sink, [144](#), [145](#)
- slice.index, [37](#)
- SmartComments, [5](#), [74](#), [114](#), [131](#)
- source, [44](#), [88](#), [116–118](#), [138](#)
- sourceDirectory, [116](#), [118](#)
- sourceTo, [116](#), [117](#), [138](#)
- sourceutils, [44](#)
- splitByPattern, [119](#)
- sprintf, [17](#), [91](#), [124](#)
- srcfile, [44](#)
- stext, [120](#)
- str, [80](#), [86](#), [135](#)
- strayDLLs (gcDLLs), [46](#)
- strsplit, [119](#)
- subplots, [121](#)
- substitute, [140–145](#)
- summary, [135](#)
- Sys.setenv, [88](#)
- Sys.setFileTime, [128](#)
- Sys.setlocale, [140](#)
- sys.source, [118](#)
- System, [5](#), [122](#)
- system, [123](#), [146](#)
- systemR, [123](#)
- TextStatusBar, [124](#)
- TimeoutException, [126](#), [146](#)
- timestamp, [135](#)
- timestampOn, [135](#)
- toCamelCase, [14](#)
- touchFile, [127](#)
- toUrl, [129](#)
- TRUE, [6](#), [11](#), [15](#), [16](#), [19](#), [21–23](#), [25](#), [26](#), [30](#), [32–34](#), [37](#), [38](#), [41](#), [45–47](#), [49](#), [60–64](#), [67](#), [70](#), [76](#), [77](#), [79](#), [81](#), [88–90](#), [92](#), [94–96](#), [102](#), [104](#), [105](#), [107](#), [116](#), [117](#), [121](#), [129–131](#), [134](#), [136](#), [138](#), [140–145](#), [152](#)
- unlink, [105](#), [106](#)
- unwrap.array, [129](#)
- unwrap.data.frame (unwrap.array), [129](#)
- unwrap.default (unwrap.array), [129](#)
- unwrap.matrix (unwrap.array), [129](#)
- update, [43](#), [93](#), [125](#)
- updateLabels, [125](#)
- URLEncode, [129](#)
- useRepos, [130](#), [142](#)
- validate, [115](#), [133](#)
- VComments, [5](#), [74](#), [114](#), [115](#), [117](#), [131](#)
- vector, [10](#), [14–17](#), [21](#), [22](#), [25](#), [28–30](#), [37](#), [54](#), [57](#), [58](#), [60](#), [65–67](#), [69](#), [70](#), [76–78](#), [81](#), [88](#), [91](#), [98](#), [102](#), [103](#), [106](#), [108](#), [109](#), [116](#), [117](#), [119](#), [120](#), [123](#), [128–131](#), [140](#), [142](#), [148](#), [151](#)
- Verbose, [5](#), [23](#), [33](#), [82](#), [89](#), [90](#), [94](#), [95](#), [98](#), [102](#), [103](#), [116](#), [123](#), [132](#), [133](#)
- warning, [88](#)
- warnings, [135](#)
- withCapture, [137](#)
- withLocale, [139](#)
- withOptions, [140](#)
- withRepos, [131](#), [142](#)
- withSeed, [143](#)
- withSink, [144](#)
- withTimeout, [145](#)
- wrap.array, [148](#)
- wrap.data.frame (wrap.array), [148](#)
- wrap.matrix (wrap.array), [148](#)

`write.table`, [152](#)  
`writeBin`, [151](#)  
`writeBinFragments`, [98](#), [151](#)  
`writeByte`, [72](#)  
`writeDataFrame`  
    (`writeDataFrame.data.frame`),  
    [152](#)  
`writeDataFrame.data.frame`, [152](#)  
`writeInt`, [72](#)  
`writeRaw`, [83](#), [135](#)  
`writeShort`, [72](#)  
`writeUTF`, [72](#)