

# Package ‘reproducible’

December 12, 2024

**Type** Package

**Title** Enhance Reproducibility of R Code

**Description** A collection of high-level, machine- and OS-independent tools for making reproducible and reusable content in R.

The two workhorse functions are `Cache()` and `prepInputs()`. `Cache()` allows for nested caching, is robust to environments and objects with environments (like functions), and deals with some classes of file-backed R objects e.g., from `terra` and `raster` packages.

Both functions have been developed to be foundational components of data retrieval and processing in continuous workflow situations. In both functions, efforts are made to make the first and subsequent calls of functions have the same result, but faster at subsequent times by way of checksums and digesting. Several features are still under development, including cloud storage of cached objects allowing for sharing between users. Several advanced options are available, see `?reproducibleOptions()`.

**SystemRequirements** 'unrar' (Linux/macOS) or '7-Zip' (Windows) to work with '.rar' files.

**URL** <https://reproducible.predictiveecology.org>,  
<https://github.com/PredictiveEcology/reproducible>

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**Depends** R (>= 4.1)

**Imports** cli, data.table (>= 1.10.4), digest, filelock, fpCompare, fs, lobstr, methods, stats, utils

**Suggests** archive, covr, curl, DBI, future, geodata, glue, googledrive, httr, httr2, knitr, parallel, qs, raster (>= 3.5-15), RCurl (>= 1.95-4.8), rlang, rmarkdown, RSQLite, R.utils, sf, sp (>= 1.4-2), terra (>= 1.7-20), testthat, withr

**Encoding** UTF-8

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**License** GPL-3

**VignetteBuilder** knitr, rmarkdown

**BugReports** <https://github.com/PredictiveEcology/reproducible/issues>

**ByteCompile** yes

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**Collate** 'DBI.R' 'cache-helpers.R' 'cache-internals.R' 'robustDigest.R'  
'cache.R' 'cacheGeo.R' 'checksums.R' 'cloud.R' 'convertPaths.R'  
'copy.R' 'download.R' 'messages.R' 'exportedMethods.R' 'gis.R'  
'helpers.R' 'listNamed.R' 'objectSize.R' 'options.R'  
'packages.R' 'paths.R' 'pipe.R' 'postProcess.R'  
'postProcessTo.R' 'preProcess.R' 'prepInputs.R'  
'reproducible-deprecated.R' 'reproducible-package.R' 'search.R'  
'showCacheEtc.R' 'spatialObjects-class.R' 'terra-migration.R'  
'zzz.R'

**NeedsCompilation** no

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---

reproducible-package *The reproducible package*

---

## Description

This package aims at making high-level, robust, machine and OS independent tools for making deeply reproducible and reusable content in R. The core user functions are `Cache` and `prepInputs`. Each of these is built around many core and edge cases required to have reproducible code of arbitrary complexity.

## Main Tools

There are many elements within the reproducible package. However, there are currently two main ones that are critical for reproducible research. The key element for reproducible research is that the code must always return the same content every time it is run, but it must be vastly faster the 2nd, 3rd, 4th etc, time it is run. That way, the entire code sequence for a project of arbitrary size can be run *from the start* every time.

**`Cache()`**: A robust wrapper for any function, including those with environments, disk-backed storage (currently on `Raster` class), operating-system independent, whose first time called will execute the function, second time will compare the inputs to a database of entries, and recover the first result if inputs are identical. If `options("reproducible.useMemoise" = TRUE)`, the second time will be very fast as it will recover the answer from RAM.

**`prepInputs()` for other specifics for other classes.**: Download, or load objects, and possibly post-process them. The main advantage to using this over more direct routes is that it will automatically build checksums tables, use `Cache` internally where helpful, and possibly run a variety of post-processing actions. This means this function can also itself be cached for even more speed. This allows all project data to be stored in custom cloud locations or in their original online data repositories, without altering code between the first, second, third, etc., times the code is run.

## Package options

See `reproducibleOptions()` for a complete description of package `options()` to configure behaviour.

## Author(s)

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### See Also

Useful links:

- <https://reproducible.predictiveecology.org>
- <https://github.com/PredictiveEcology/reproducible>
- Report bugs at <https://github.com/PredictiveEcology/reproducible/issues>

---

.debugCache

*Attach debug info to return for Cache*

---

### Description

Internal use only. Attaches an attribute to the output, usable for debugging the Cache.

### Usage

```
.debugCache(obj, preDigest, ...)
```

### Arguments

obj	An arbitrary R object.
preDigest	A list of hashes.
...	Dots passed from Cache

### Value

The same object as obj, but with 2 attributes set.

### Author(s)

Eliot McIntire

---

<code>.file.move</code>	<i>Move a file to a new location – Defunct – use <code>hardLinkOrCopy</code></i>
-------------------------	--

---

**Description**

This will first try to `file.rename`, and if that fails, then it will `file.copy` then `file.remove`.

**Usage**

```
.file.move(from, to, overwrite = FALSE)
```

**Arguments**

<code>from, to</code>	character vectors, containing file names or paths.
<code>overwrite</code>	logical indicating whether to overwrite destination file if it exists.

**Value**

Logical indicating whether operation succeeded.

---

<code>.isMemoised</code>	<i>Evaluate whether a cacheId is memoised</i>
--------------------------	---

---

**Description**

Intended for internal use. Exported so other packages can use this function.

**Usage**

```
.isMemoised(cacheId, cachePath = getOption("reproducible.cachePath"))
```

**Arguments**

<code>cacheId</code>	Character string. If passed, this will override the calculated hash of the inputs, and return the result from this <code>cacheId</code> in the <code>cachePath</code> . Setting this is equivalent to manually saving the output of this function, i.e., the object will be on disk, and will be recovered in subsequent This may help in some particularly finicky situations where <code>Cache</code> is not correctly detecting unchanged inputs. This will guarantee the object will be identical each time; this may be useful in operational code.
<code>cachePath</code>	A repository used for storing cached objects. This is optional if <code>Cache</code> is used inside a <code>SpaDES</code> module.

**Value**

A logical, length 1 indicating whether the `cacheId` is memoised.

---

.prefix *Add a prefix or suffix to the basename part of a file path*

---

### Description

Prepend (or postpend) a filename with a prefix (or suffix). If the directory name of the file cannot be ascertained from its path, it is assumed to be in the current working directory.

### Usage

```
.prefix(f, prefix = "")
```

```
.suffix(f, suffix = "")
```

### Arguments

f	A character string giving the name/path of a file.
prefix	A character string to prepend to the filename.
suffix	A character string to postpend to the filename.

### Value

A character string or vector with the prefix pre-pended or suffix post-pended on the basename of the f, before the file extension.

### Author(s)

Jean Marchal and Alex Chubaty

### Examples

```
# file's full path is specified (i.e., dirname is known)
myFile <- file.path("~/data", "file.tif")
.prefix(myFile, "small_") ## "/home/username/data/small_file.tif"
.suffix(myFile, "_cropped") ## "/home/username/data/myFile_cropped.shp"

# file's full path is not specified
.prefix("myFile.shp", "small") ## "./small_myFile.shp"
.suffix("myFile.shp", "_cropped") ## "./myFile_cropped.shp"
```

---

`.prepareFileBackedRaster`*Copy the file-backing of a file-backed Raster\* object*

---

## Description

Rasters are sometimes file-based, so the normal save and copy and assign mechanisms in R don't work for saving, copying and assigning. This function creates an explicit file copy of the file that is backing the raster, and changes the pointer (i.e., `filename(object)`) so that it is pointing to the new file.

## Usage

```
.prepareFileBackedRaster(  
  obj,  
  repoDir = NULL,  
  overwrite = FALSE,  
  drv = getDrv(getOption("reproducible.drv", NULL)),  
  conn = getOption("reproducible.conn", NULL),  
  ...  
)
```

## Arguments

<code>obj</code>	The raster object to save to the repository.
<code>repoDir</code>	Character denoting an existing directory in which an artifact will be saved.
<code>overwrite</code>	Logical. Should the raster be saved to disk, overwriting existing file.
<code>drv</code>	if using a database backend, <code>drv</code> must be an object that inherits from <code>DBIDriver</code> e.g., from package <code>RSQLite</code> , e.g., <code>SQLite</code>
<code>conn</code>	an optional <code>DBIConnection</code> object, as returned by <code>dbConnect()</code> .
<code>...</code>	Not used

## Value

A raster object and its newly located file backing. Note that if this is a legitimate Cache repository, the new location will be a subdirectory called `'rasters/'` of `'repoDir/'`. If this is not a repository, the new location will be within `repoDir`.

## Author(s)

Eliot McIntire



---

.removeCacheAtts      *Remove attributes that are highly varying*

---

**Description**

Remove attributes that are highly varying

**Usage**

```
.removeCacheAtts(x)
```

**Arguments**

x                      Any arbitrary R object that could have attributes

---

.requireNamespace      *Provide standard messaging for missing package dependencies*

---

**Description**

This provides a standard message format for missing packages, e.g., detected via requireNamespace.

**Usage**

```
.requireNamespace(  
  pkg = "methods",  
  minVersion = NULL,  
  stopOnFALSE = FALSE,  
  messageStart = NULL  
)
```

**Arguments**

pkg                      Character string indicating name of package required  
minVersion              Character string indicating minimum version of package that is needed  
stopOnFALSE            Logical. If TRUE, this function will create an error (i.e., stop) if the function returns FALSE; otherwise it simply returns FALSE  
messageStart            A character string with a prefix of message to provide

**Value**

A logical or stop if the namespace is not available to be loaded.

---

`.setSubAttrInList`      *Set subattributes within a list by reference*

---

### Description

Sets only a single element within a list attribute.

### Usage

```
.setSubAttrInList(object, attr, subAttr, value)
```

### Arguments

<code>object</code>	An arbitrary object
<code>attr</code>	The attribute name (that is a list object) to change
<code>subAttr</code>	The list element name to change
<code>value</code>	The new value

### Value

This sets or updates the `subAttr` element of a list that is located at `attr(object, attr)`, with the `value`. This, therefore, updates a sub-element of a list attribute and returns that same object with the updated attribute.

---

`.wrap`      *Deal with class for saving to and loading from Cache or Disk*

---

### Description

This generic and some methods will do whatever is required to prepare an object for saving to disk (or RAM) via e.g., `saveRDS`. Some objects (e.g., terra's `Spat*`) cannot be saved without first wrapping them. Also, file-backed objects are similar.

### Usage

```
.wrap(
  obj,
  cachePath,
  preDigest,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  outputObjects = NULL,
  ...
)
```

```
## S3 method for class 'list'
.wrap(
  obj,
  cachePath,
  preDigest,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  outputObjects = NULL,
  ...
)

## S3 method for class 'environment'
.wrap(
  obj,
  cachePath,
  preDigest,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  outputObjects = NULL,
  ...
)

## Default S3 method:
.wrap(
  obj,
  cachePath,
  preDigest,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  ...
)

## Default S3 method:
.unwrap(
  obj,
  cachePath,
  cacheId,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  ...
)

.unwrap(
  obj,
```

```

    cachePath,
    cacheId,
    drv = getDrv(getOption("reproducible.drv", NULL)),
    conn = getOption("reproducible.conn", NULL),
    ...
)

## S3 method for class 'environment'
.unwrap(
  obj,
  cachePath,
  cacheId,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  ...
)

## S3 method for class 'list'
.unwrap(
  obj,
  cachePath,
  cacheId,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  ...
)

```

### Arguments

<code>obj</code>	Any arbitrary R object.
<code>cachePath</code>	A repository used for storing cached objects. This is optional if Cache is used inside a SpaDES module.
<code>preDigest</code>	The list of <code>preDigest</code> that comes from <code>CacheDigest</code> of an object
<code>drv</code>	if using a database backend, <code>drv</code> must be an object that inherits from <code>DBIDriver</code> e.g., from package <code>RSQLite</code> , e.g., <code>SQLite</code>
<code>conn</code>	an optional <code>DBIConnection</code> object, as returned by <code>dbConnect()</code> .
<code>verbose</code>	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t
<code>outputObjects</code>	Optional character vector indicating which objects to return. This is only relevant for <code>list</code> , <code>environment</code> (or similar) objects
<code>...</code>	Arguments passed to methods; default does not use anything in <code>...</code>
<code>cacheId</code>	Used strictly for messaging. This should be the <code>cacheId</code> of the object being recovered.

**Value**

Returns an object that can be saved to disk e.g., via saveRDS.

**Examples**

```
# For SpatExtent
if (requireNamespace("terra")) {
  ex <- terra::ext(c(0, 2, 0, 3))
  exWrapped <- .wrap(ex)
  ex1 <- .unwrap(exWrapped)
}
```

---

assessDataType	<i>Assess the appropriate raster layer data type</i>
----------------	--

---

**Description**

When writing raster-type objects to disk, a datatype can be specified. These functions help identify what smallest datatype can be used.

**Usage**

```
assessDataType(ras, type = "writeRaster")

## Default S3 method:
assessDataType(ras, type = "writeRaster")
```

**Arguments**

ras                    The RasterLayer or RasterStack for which data type will be assessed.  
 type                    Character. "writeRaster" (default) or "GDAL" (defunct) to return the recommended data type for writing from the raster packages, respectively, or "projectRaster" to return recommended resampling type.

**Value**

A character string indicating the data type of the spatial layer (e.g., "INT2U"). See terra::datatype()

**Examples**

```
if (requireNamespace("terra", quietly = TRUE)) {
  ## LOG1S
  rasOrig <- terra::rast(ncols = 10, nrows = 10)
  ras <- rasOrig
  ras[] <- rep(c(0,1),50)
  assessDataType(ras)
```

```
ras <- rasOrig
ras[] <- rep(c(0,1),50)
assessDataType(ras)

ras[] <- rep(c(TRUE,FALSE),50)
assessDataType(ras)

ras[] <- c(NA, NA, rep(c(0,1),49))
assessDataType(ras)

ras <- rasOrig
ras[] <- c(0, NaN, rep(c(0,1),49))
assessDataType(ras)

## INT1S
ras[] <- -1:98
assessDataType(ras)

ras[] <- c(NA, -1:97)
assessDataType(ras)

## INT1U
ras <- rasOrig
ras[] <- 1:100
assessDataType(ras)

ras[] <- c(NA, 2:100)
assessDataType(ras)

## INT2U
ras <- rasOrig
ras[] <- round(runif(100, min = 64000, max = 65000))
assessDataType(ras)

## INT2S
ras <- rasOrig
ras[] <- round(runif(100, min = -32767, max = 32767))
assessDataType(ras)

ras[54] <- NA
assessDataType(ras)

## INT4U
ras <- rasOrig
ras[] <- round(runif(100, min = 0, max = 500000000))
assessDataType(ras)

ras[14] <- NA
assessDataType(ras)

## INT4S
ras <- rasOrig
```

```

ras[] <- round(runif(100, min = -200000000, max = 200000000))
assessDataType(ras)

ras[14] <- NA
assessDataType(ras)

## FLT4S
ras <- rasOrig
ras[] <- runif(100, min = -10, max = 87)
assessDataType(ras)

ras <- rasOrig
ras[] <- round(runif(100, min = -3.4e+26, max = 3.4e+28))
assessDataType(ras)

ras <- rasOrig
ras[] <- round(runif(100, min = 3.4e+26, max = 3.4e+28))
assessDataType(ras)

ras <- rasOrig
ras[] <- round(runif(100, min = -3.4e+26, max = -1))
assessDataType(ras)

## FLT8S
ras <- rasOrig
ras[] <- c(-Inf, 1, rep(c(0,1),49))
assessDataType(ras)

ras <- rasOrig
ras[] <- c(Inf, 1, rep(c(0,1),49))
assessDataType(ras)

ras <- rasOrig
ras[] <- round(runif(100, min = -1.7e+30, max = 1.7e+308))
assessDataType(ras)

ras <- rasOrig
ras[] <- round(runif(100, min = 1.7e+30, max = 1.7e+308))
assessDataType(ras)

ras <- rasOrig
ras[] <- round(runif(100, min = -1.7e+308, max = -1))
assessDataType(ras)

# 2 layer with different types LOG1S and FLT8S
ras <- rasOrig
ras[] <- rep(c(0,1),50)
ras1 <- rasOrig
ras1[] <- round(runif(100, min = -1.7e+308, max = -1))
sta <- c(ras, ras1)
assessDataType(sta)
}

```

---

basename2	<i>A version of base::basename that is NULL resistant</i>
-----------	---

---

**Description**

A version of base::basename that is NULL resistant

**Usage**

```
basename2(x)
```

**Arguments**

x	A character vector of paths
---	-----------------------------

**Value**

NULL if x is NULL, otherwise, as basename.

Same as [base::basename\(\)](#)

---

Cache	<i>Saves a wide variety function call outputs to disk and optionally RAM, for recovery later</i>
-------	--

---

**Description**

A function that can be used to wrap around other functions to cache function calls for later use. This is normally most effective when the function to cache is slow to run, yet the inputs and outputs are small. The benefit of caching, therefore, will decline when the computational time of the "first" function call is fast and/or the argument values and return objects are large. The default setting (and first call to Cache) will always save to disk. The 2nd call to the same function will return from disk, unless `options("reproducible.useMemoise" = TRUE)`, then the 2nd time will recover the object from RAM and is normally much faster (at the expense of RAM use).

**Usage**

```
Cache(
  FUN,
  ...,
  notOlderThan = NULL,
  .objects = NULL,
  .cacheExtra = NULL,
  .functionName = NULL,
  outputObjects = NULL,
  algo = "xxhash64",
```



```

cacheRepo = NULL,
cachePath = NULL,
length = getOption("reproducible.length", Inf),
compareRasterFileLength,
userTags = c(),
omitArgs = NULL,
classOptions = list(),
debugCache = character(),
sideEffect = FALSE,
makeCopy = FALSE,
quick = getOption("reproducible.quick", FALSE),
verbose = getOption("reproducible.verbose", 1),
cacheId = NULL,
useCache = getOption("reproducible.useCache", TRUE),
useCloud = FALSE,
cloudFolderID = NULL,
showSimilar = getOption("reproducible.showSimilar", FALSE),
drv = getDrv(getOption("reproducible.drv", NULL)),
conn = getOption("reproducible.conn", NULL)
)

```

### Arguments

<code>FUN</code>	Either a function (e.g., <code>rnorm</code> ), a function call (e.g., <code>rnorm(1)</code> ), or an unevaluated function call (e.g., using quote).
<code>...</code>	Arguments passed to <code>FUN</code> , if <code>FUN</code> is not an expression.
<code>notOlderThan</code>	A time. Load an object from the Cache if it was created after this.
<code>.objects</code>	Character vector of objects to be digested. This is only applicable if there is a list, environment (or similar) with named objects within it. Only this/these objects will be considered for caching, i.e., only use a subset of the list, environment or similar objects. In the case of nested list-type objects, this will only be applied outermost first.
<code>.cacheExtra</code>	A an arbitrary R object that will be included in the <code>CacheDigest</code> , but otherwise not passed into the <code>FUN</code> . If the user supplies a named list, then <code>Cache</code> will report which individual elements of <code>.cacheExtra</code> have changed when <code>options("reproducible.showSimilar" = TRUE)</code> . This can allow a user more control and understanding for debugging.
<code>.functionName</code>	A an arbitrary character string that provides a name that is different than the actual function name (e.g., "rnorm") which will be used for messaging. This can be useful when the actual function is not helpful for a user, such as <code>do.call</code> .
<code>outputObjects</code>	Optional character vector indicating which objects to return. This is only relevant for list, environment (or similar) objects
<code>algo</code>	The algorithms to be used; currently available choices are <code>md5</code> , which is also the default, <code>sha1</code> , <code>crc32</code> , <code>sha256</code> , <code>sha512</code> , <code>xxhash32</code> , <code>xxhash64</code> , <code>murmur32</code> , <code>spookyhash</code> , <code>blake3</code> , <code>crc32c</code> , <code>xxh3_64</code> , and <code>xxh3_128</code> .
<code>cacheRepo</code>	Same as <code>cachePath</code> , but kept for backwards compatibility.
<code>cachePath</code>	A repository used for storing cached objects. This is optional if <code>Cache</code> is used inside a <code>SpaDES</code> module.

length	Numeric. If the element passed to Cache is a Path class object (from e.g., <code>asPath(filename)</code> ) or it is a Raster with file-backing, then this will be passed to <code>digest::digest</code> , essentially limiting the number of bytes to digest (for speed). This will only be used if <code>quick = FALSE</code> . Default is <code>getOption("reproducible.length")</code> , which is set to <code>Inf</code> .
compareRasterFileLength	Being deprecated; use <code>length</code> .
userTags	A character vector with descriptions of the Cache function call. These will be added to the Cache so that this entry in the Cache can be found using <code>userTags</code> e.g., via <code>showCache()</code> .
omitArgs	Optional character string of arguments in the FUN to omit from the digest.
classOptions	Optional list. This will pass into <code>.robustDigest</code> for specific classes. Should be options that the <code>.robustDigest</code> knows what to do with.
debugCache	Character or Logical. Either "complete" or "quick" (uses partial matching, so "c" or "q" work). TRUE is equivalent to "complete". If "complete", then the returned object from the Cache function will have two attributes, <code>debugCache1</code> and <code>debugCache2</code> , which are the entire <code>list(...)</code> and that same object, but after all <code>.robustDigest</code> calls, at the moment that it is digested using <code>digest</code> , respectively. This <code>attr(mySimOut, "debugCache2")</code> can then be compared to a subsequent call and individual items within the object <code>attr(mySimOut, "debugCache1")</code> can be compared. If "quick", then it will return the same two objects directly, without evaluating the <code>FUN(...)</code> .
sideEffect	Now deprecated. Logical or path. Determines where the function will look for new files following function completion. See Details. <i>NOTE: this argument is experimental and may change in future releases.</i>
makeCopy	Now deprecated. Ignored if used.
quick	Logical or character. If TRUE, no disk-based information will be assessed, i.e., only memory content. See Details section about <code>quick</code> in <code>Cache()</code> .
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t
cacheId	Character string. If passed, this will override the calculated hash of the inputs, and return the result from this <code>cacheId</code> in the <code>cachePath</code> . Setting this is equivalent to manually saving the output of this function, i.e., the object will be on disk, and will be recovered in subsequent This may help in some particularly finicky situations where Cache is not correctly detecting unchanged inputs. This will guarantee the object will be identical each time; this may be useful in operational code.
useCache	Logical, numeric or "overwrite" or "devMode". See details.
useCloud	Logical. See Details.
cloudFolderID	A googledrive dribble of a folder, e.g., using <code>drive_mkdir()</code> . If left as NULL, the function will create a cloud folder with name from last two folder levels of the <code>cachePath</code> path, <code>: paste0(basename(dirname(cachePath)), "_", basename(cachePath))</code> . This <code>cloudFolderID</code> will be added to <code>options("reproducible.cloudFolderID")</code> .

	but this will not persist across sessions. If this is a character string, it will treat this as a folder name to create or use on GoogleDrive.
showSimilar	A logical or numeric. Useful for debugging. If TRUE or 1, then if the Cache does not find an identical archive in the cachePath, it will report (via message) the next most recent similar archive, and indicate which argument(s) is/are different. If a number larger than 1, then it will report the N most recent similar archived objects.
drv	if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite
conn	an optional DBIConnection object, as returned by dbConnect().

### Details

There are other similar functions in the R universe. This version of Cache has been used as part of a robust continuous workflow approach. As a result, we have tested it with many "non-standard" R objects (e.g., RasterLayer, Spat\* objects) and environments (which are always unique, so do not cache readily).

This version of the Cache function accommodates those four special, though quite common, cases by:

1. converting any environments into list equivalents;
2. identifying the dispatched S4 method (including those made through inheritance) before hashing so the correct method is being cached;
3. by hashing the linked file, rather than the raster object. Currently, only file-backed Raster\* or Spat\* objects are digested (e.g., not ff objects, or any other R object where the data are on disk instead of in RAM);
4. Uses `digest::digest()` This is used for file-backed objects as well.
5. Cache will save arguments passed by user in a hidden environment. Any nested Cache functions will use arguments in this order: 1) actual arguments passed at each Cache call; 2) any inherited arguments from an outer Cache call; 3) the default values of the Cache function. See section on *Nested Caching*.

Cache will add a tag to the entry in the cache database called `accessed`, which will assign the time that it was accessed, either read or write. That way, cached items can be shown (using `showCache`) or removed (using `clearCache`) selectively, based on their access dates, rather than only by their creation dates. See example in `clearCache()`.

### Value

Returns the value of the function call or the cached version (i.e., the result from a previous call to this same cached function with identical arguments).

### Nested Caching

Commonly, Caching is nested, i.e., an outer function is wrapped in a Cache function call, and one or more inner functions are also wrapped in a Cache function call. A user *can* always specify arguments in every Cache function call, but this can get tedious and can be prone to errors. The

normal way that R handles arguments is it takes the user passed arguments if any, and default arguments for all those that have no user passed arguments. We have inserted a middle step. The order or precedence for any given Cache function call is

1. user arguments, 2. inherited arguments, 3. default arguments. At this time, the top level Cache arguments will propagate to all inner functions unless each individual Cache call has other arguments specified, i.e., "middle" nested Cache function calls don't propagate their arguments to further "inner" Cache function calls. See example.

`userTags` is unique of all arguments: its values will be appended to the inherited `userTags`.

### quick

The `quick` argument is attempting to sort out an ambiguity with character strings: are they file paths or are they simply character strings. When `quick = TRUE`, Cache will treat these as character strings; when `quick = FALSE`, they will be attempted to be treated as file paths first; if there is no file, then it will revert to treating them as character strings. If user passes a character vector to this, then this will behave like `omitArgs`: `quick = "file"` will treat the argument "file" as character string.

The most often encountered situation where this ambiguity matters is in arguments about filenames: is the filename an input pointing to an object whose content we want to assess (e.g., a file-backed raster), or an output (as in `saveRDS`) and it should not be assessed. If only run once, the output file won't exist, so it will be treated as a character string. However, once the function has been run once, the output file will exist, and `Cache(...)` will assess it, which is incorrect. In these cases, the user is advised to use `quick = "TheOutputFilenameArgument"` to specify the argument whose content on disk should not be assessed, but whose character string should be assessed (distinguishing it from `omitArgs = "TheOutputFilenameArgument"`, which will not assess the file content nor the character string).

This is relevant for objects of class `character`, `Path` and `Raster` currently. For class `character`, it is ambiguous whether this represents a character string or a vector of file paths. If it is known that character strings should not be treated as paths, then `quick = TRUE` is appropriate, with no loss of information. If it is file or directory, then it will digest the file content, or `basename(object)`. For class `Path` objects, the file's metadata (i.e., filename and file size) will be hashed instead of the file contents if `quick = TRUE`. If set to `FALSE` (default), the contents of the file(s) are hashed. If `quick = TRUE`, `length` is ignored. `Raster` objects are treated as paths, if they are file-backed.

### Caching Speed

Caching speed may become a critical aspect of a final product. For example, if the final product is a shiny app, rerunning the entire project may need to take less than a few seconds at most. There are 3 arguments that affect Cache speed: `quick`, `length`, and `algo`. `quick` is passed to `.robustDigest`, which currently only affects `Path` and `Raster*` class objects. In both cases, `quick` means that little or no disk-based information will be assessed.

### Filepaths

If a function has a path argument, there is some ambiguity about what should be done. Possibilities include:

1. hash the string as is (this will be very system specific, meaning a Cache call will not work if copied between systems or directories);
2. hash the basename(path);
3. hash the contents of the file.

If paths are passed in as is (i.e., character string), the result will not be predictable. Instead, one should use the wrapper function `asPath(path)`, which sets the class of the string to a `Path`, and one should decide whether one wants to digest the content of the file (using `quick = FALSE`), or just the filename (`quick = TRUE`). See examples.

### Stochasticity or randomness

In general, it is expected that caching will only be used when randomness is not desired, e.g., `Cache(rnorm(1))` is unlikely to be useful in many cases. However, `Cache` captures the call that is passed to it, leaving all functions unevaluated. As a result `Cache(glm, x ~ y, rnorm(1))` will not work as a means of forcing a new evaluation each time, as the `rnorm(1)` is not evaluated before the call is assessed against the cache database. To force a new call each time, evaluate the randomness prior to the `Cache` call, e.g., `ran = rnorm(1)` then pass this to `.cacheExtra`, e.g., `Cache(glm, x ~ y, .cacheExtra = ran)`

### drv and conn

By default, `drv` uses an SQLite database. This can be sufficient for most cases. However, if a user has dozens or more cores making requests to the `Cache` database, it may be insufficient. A user can set up a different database backend, e.g., PostgreSQL that can handle multiple simultaneous read-write situations. See <https://github.com/PredictiveEcology/SpaDES/wiki/Using-alternate-database-backends>

### useCache

Logical or numeric. If `FALSE` or `0`, then the entire Caching mechanism is bypassed and the function is evaluated as if it was not being Cached. Default is `getOption("reproducible.useCache")`, which is `TRUE` by default, meaning use the `Cache` mechanism. This may be useful to turn all Caching on or off in very complex scripts and nested functions. Increasing levels of numeric values will cause deeper levels of Caching to occur (though this may not work as expected in all cases). The following is no longer supported: Currently, only implemented in `postProcess`: to do both caching of inner `cropInputs`, `projectInputs` and `maskInputs`, and caching of outer `postProcess`, use `useCache = 2`; to skip the inner sequence of 3 functions, use `useCache = 1`. For large objects, this may prevent many duplicated save to disk events.

If `useCache = "overwrite"` (which can be set with `options("reproducible.useCache" = "overwrite")`), then the function invoke the caching mechanism but will purge any entry that is matched, and it will be replaced with the results of the current call.

If `useCache = "devMode"`: The point of this mode is to facilitate using the `Cache` when functions and datasets are continually in flux, and old `Cache` entries are likely stale very often. In `devMode`, the cache mechanism will work as normal if the `Cache` call is the first time for a function OR if it successfully finds a copy in the cache based on the normal `Cache` mechanism. It *differs* from the normal `Cache` if the `Cache` call does *not* find a copy in the `cachePath`, but it does find an entry that matches based on `userTags`. In this case, it will delete the old entry in the `cachePath` (identified based on matching `userTags`), then continue with normal `Cache`. For this to work correctly,

`userTags` must be unique for each function call. This should be used with caution as it is still experimental. Currently, if `userTags` are not unique to a single entry in the `cachePath`, it will default to the behaviour of `useCache = TRUE` with a message. This means that "devMode" is most useful if used from the start of a project.

### `useCloud`

This is experimental and there are many conditions under which this is known to not work correctly. This is a way to store all or some of the local Cache in the cloud. Currently, the only cloud option is Google Drive, via **googledrive**. For this to work, the user must be or be able to be authenticated with `googledrive::drive_auth`. The principle behind this `useCloud` is that it will be a full or partial mirror of a local Cache. It is not intended to be used independently from a local Cache. To share objects that are in the Cloud with another person, it requires 2 steps. 1) share the `cloudFolderID`, which can be retrieved by `getOption("reproducible.cloudFolderID")$id` after at least one Cache call has been made. 2) The other user must then set their `cacheFolderID` in a `Cache(..., reproducible.cloudFolderID = "the ID here")` call or set their option manually `options(\("reproducible.cloudFolderID" = "the ID here")`.

If `TRUE`, then this Cache call will download (if local copy doesn't exist, but cloud copy does exist), upload (local copy does or doesn't exist and cloud copy doesn't exist), or will not download nor upload if object exists in both. If `TRUE` will be at least 1 second slower than setting this to `FALSE`, and likely even slower as the cloud folder gets large. If a user wishes to keep "high-level" control, set this to `getOption("reproducible.useCloud", FALSE)` or `getOption("reproducible.useCloud", TRUE)` (if the default behaviour should be `FALSE` or `TRUE`, respectively) so it can be turned on and off with this option. NOTE: *This argument will not be passed into inner/nested Cache calls.*

### Object attributes

Users should be cautioned that object attributes may not be preserved, especially in the case of objects that are file-backed, such as `Raster` or `SpatRaster` objects. If a user needs to keep attributes, they may need to manually re-attach them to the object after recovery. With the example of `SpatRaster` objects, saving to disk requires `terra::wrap` if it is a memory-backed object. When running `terra::unwrap` on this object, any attributes that a user had added are lost.

### `sideEffect`

This feature is now deprecated. Do not use as it is ignored.

### Note

As indicated above, several objects require pre-treatment before caching will work as expected. The function `.robustDigest` accommodates this. It is an S4 generic, meaning that developers can produce their own methods for different classes of objects. Currently, there are methods for several types of classes. See `.robustDigest()`.

### Author(s)

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**See Also**

`showCache()`, `clearCache()`, `keepCache()`, `CacheDigest()` to determine the digest of a given function or expression, as used internally within `Cache`, `movedCache()`, `.robustDigest()`, and for more advanced uses there are several helper functions, e.g., `rmFromCache()`, `CacheStorageDir()`

**Examples**

```
data.table::setDTthreads(2)
tmpDir <- file.path(tempdir())
opts <- options(reproducible.cachePath = tmpDir)

# Usage -- All below are equivalent; even where args are missing or provided,
# Cache evaluates using default values, if these are specified in formals(FUN)
a <- list()
b <- list(fun = rnorm)
bbb <- 1
ee <- new.env(parent = emptyenv())
ee$qq <- bbb

a[[1]] <- Cache(rnorm(1)) # no evaluation prior to Cache
a[[2]] <- Cache(rnorm, 1) # no evaluation prior to Cache
a[[3]] <- Cache(do.call, rnorm, list(1))
a[[4]] <- Cache(do.call(rnorm, list(1)))
a[[5]] <- Cache(do.call(b$fun, list(1)))
a[[6]] <- Cache(do.call, b$fun, list(1))
a[[7]] <- Cache(b$fun, 1)
a[[8]] <- Cache(b$fun(1))
a[[10]] <- Cache(quote(rnorm(1)))
a[[11]] <- Cache(stats::rnorm(1))
a[[12]] <- Cache(stats::rnorm, 1)
a[[13]] <- Cache(rnorm(1, 0, get("bbb", inherits = FALSE)))
a[[14]] <- Cache(rnorm(1, 0, get("qq", inherits = FALSE, envir = ee)))
a[[15]] <- Cache(rnorm(1, bbb - bbb, get("bbb", inherits = FALSE)))
a[[16]] <- Cache(rnorm(sd = 1, 0, n = get("bbb", inherits = FALSE))) # change order
a[[17]] <- Cache(rnorm(1, sd = get("ee", inherits = FALSE)$qq, mean = 0)

# with base pipe -- this is put in quotes (') because R version 4.0 can't understand this
# if you are using R >= 4.1 or R >= 4.2 if using the _ placeholder,
# then you can just use pipe normally
usingPipe1 <- "b$fun(1) |> Cache()" # base pipe

# For long pipe, need to wrap sequence in { }, or else only last step is cached
usingPipe2 <-
  '{"bbb" |>
    parse(text = _) |>
    eval() |>
    rnorm()} |>
  Cache()'
if (getRversion() >= "4.1") {
  a[[9]] <- eval(parse(text = usingPipe1)) # recovers cached copy
}
if (getRversion() >= "4.2") { # uses the _ placeholder; only available in R >= 4.2
```

```

  a[[18]] <- eval(parse(text = usingPipe2)) # recovers cached copy
}

length(unique(a)) == 1 # all same

### Pipe -- have to use { } or else only final function is Cached
if (getRversion() >= "4.1") {
  b1a <- 'sample(1e5, 1) |> rnorm() |> Cache()'
  b1b <- 'sample(1e5, 1) |> rnorm() |> Cache()'
  b2a <- '{sample(1e5, 1) |> rnorm()} |> Cache()'
  b2b <- '{sample(1e5, 1) |> rnorm()} |> Cache()'
  b1a <- eval(parse(text = b1a))
  b1b <- eval(parse(text = b1b))
  b2a <- eval(parse(text = b2a))
  b2b <- eval(parse(text = b2b))
  all.equal(b1a, b1b) # Not TRUE because the sample is run first
  all.equal(b2a, b2b) # TRUE because of { }
}

#####
# Advanced examples
#####

# .cacheExtra -- add something to digest
Cache(rnorm(1), .cacheExtra = "sfessee11") # adds something other than fn args
Cache(rnorm(1), .cacheExtra = "nothing") # even though fn is same, the extra is different

# omitArgs -- remove something from digest (kind of the opposite of .cacheExtra)
Cache(rnorm(2, sd = 1), omitArgs = "sd") # removes one or more args from cache digest
Cache(rnorm(2, sd = 2), omitArgs = "sd") # b/c sd is not used, this is same as previous

# cacheId -- force the use of a digest -- can give undesired consequences
Cache(rnorm(3), cacheId = "k323431232") # sets the cacheId for this call
Cache(runif(14), cacheId = "k323431232") # recovers same as above, i.e., rnorm(3)

# Turn off Caching session-wide
opts <- options(reproducible.useCache = FALSE)
Cache(rnorm(3)) # doesn't cache
options(opts)

# showSimilar can help with debugging why a Cache call isn't picking up a cached copy
Cache(rnorm(4), showSimilar = TRUE) # shows that the argument `n` is different

#####
# devMode -- enables cache database to stay
#             small even when developing code
#####
opt <- options("reproducible.useCache" = "devMode")
clearCache(tmpDir, ask = FALSE)
centralTendency <- function(x) {
  mean(x)
}
funnyData <- c(1, 1, 1, 1, 10)

```



```

uniqueUserTags <- c("thisIsUnique", "reallyUnique")
ranNumsB <- Cache(centralTendency, funnyData, cachePath = tmpDir,
                 userTags = uniqueUserTags) # sets new value to Cache
showCache(tmpDir) # 1 unique cacheId -- cacheId is 71cd24ec3b0d0cac

# During development, we often redefine function internals
centralTendency <- function(x) {
  median(x)
}
# When we rerun, we don't want to keep the "old" cache because the function will
# never again be defined that way. Here, because of userTags being the same,
# it will replace the entry in the Cache, effectively overwriting it, even though
# it has a different cacheId
ranNumsD <- Cache(centralTendency, funnyData, cachePath = tmpDir, userTags = uniqueUserTags)
showCache(tmpDir) # 1 unique artifact -- cacheId is 632cd06f30e111be

# If it finds it by cacheID, doesn't matter what the userTags are
ranNumsD <- Cache(centralTendency, funnyData, cachePath = tmpDir, userTags = "thisIsUnique")
options(opt)

#####
# For more in depth uses, see vignette
if (interactive())
  browseVignettes(package = "reproducible")

```

---

CacheDigest

*The exact digest function that Cache uses*


---

### Description

This can be used by a user to pre-test their arguments before running Cache, for example to determine whether there is a cached copy.

### Usage

```

CacheDigest(
  objsToDigest,
  ...,
  algo = "xxhash64",
  calledFrom = "CacheDigest",
  .functionName = NULL,
  quick = FALSE
)

```

### Arguments

`objsToDigest` A list of all the objects (e.g., arguments) to be digested  
`...` passed to `.robustDigest`.

algo	The algorithms to be used; currently available choices are md5, which is also the default, sha1, crc32, sha256, sha512, xxhash32, xxhash64, murmur32, spookyhash, blake3, crc32c, xxh3_64, and xxh3_128.
calledFrom	a Character string, length 1, with the function to compare with. Default is "Cache". All other values may not produce robust CacheDigest results.
.functionName	A an arbitrary character string that provides a name that is different than the actual function name (e.g., "rnorm") which will be used for messaging. This can be useful when the actual function is not helpful for a user, such as do.call.
quick	Logical or character. If TRUE, no disk-based information will be assessed, i.e., only memory content. See Details section about quick in <a href="#">Cache()</a> .

### Value

A list of length 2 with the outputHash, which is the digest that Cache uses for cacheId and also preDigest, which is the digest of each sub-element in objsToDigest.

### Examples

```
data.table::setDTthreads(2)
a <- Cache(rnorm, 1)

# like with Cache, user can pass function and args in a few ways
CacheDigest(rnorm(1)) # shows same cacheId as previous line
CacheDigest(rnorm, 1) # shows same cacheId as previous line
```

---

CacheGeo

*Cache-like function for spatial domains*

---

### Description

### Usage

```
CacheGeo(
  targetFile = NULL,
  url = NULL,
  domain,
  FUN,
  destinationPath = getOption("reproducible.destinationPath", "."),
  useCloud = getOption("reproducible.useCloud", FALSE),
  cloudFolderID = NULL,
  purge = FALSE,
  useCache = getOption("reproducible.useCache"),
  overwrite = getOption("reproducible.overwrite"),
  action = c("nothing", "update", "replace", "append"),
  ...
)
```

**Arguments**

<code>targetFile</code>	The (optional) local file (or path to file)
<code>url</code>	The (optional) url of the object on Google Drive (the only option currently). This is only for downloading and uploading to.
<code>domain</code>	An sf polygon object that is the spatial area of interest. If NULL, then this will return the whole object in <code>targetFile</code> .
<code>FUN</code>	A function call that will be called if there is the domain is not already contained within the sf object at <code>url</code> or <code>targetFile</code> . This function call <b>MUST</b> return either a sf class object or a data.frame class object that has a geometry column (which can then be converted to sf with <code>st_as_sf</code> )
<code>destinationPath</code>	Character string of a directory in which to download and save the file that comes from <code>url</code> and is also where the function will look for <code>archive</code> or <code>targetFile</code> . NOTE (still experimental): To prevent repeated downloads in different locations, the user can also set <code>options("reproducible.inputPaths")</code> to one or more local file paths to search for the file before attempting to download. Default for that option is NULL meaning do not search locally.
<code>useCloud</code>	A logical.
<code>cloudFolderID</code>	If this is specified, then it must be either 1) a googledrive url to a folder where the <code>targetFile</code> will be read from or written to, or 2) a googledrive id or 3) an absolute path to a (possibly non-existent yet) folder on your google drive.
<code>purge</code>	Logical or Integer. 0/FALSE (default) keeps existing CHECKSUMS.txt file and prepInputs will write or append to it. 1/TRUE will deleted the entire CHECKSUMS.txt file. Other options, see details.
<code>useCache</code>	Passed to Cache in various places. Defaults to <code>getOption("reproducible.useCache", 2L)</code> in <code>prepInputs</code> , and <code>getOption("reproducible.useCache", FALSE)</code> if calling any of the inner functions manually. For <code>prepInputs</code> , this mean it will use Cache only up to 2 nested levels, which includes <code>preProcess</code> , <code>postProcess</code> and its nested <code>*Input</code> functions (e.g., <code>cropInputs</code> , <code>projectInputs</code> , <code>maskInputs</code> ) are no longer internally cached, as terra processing speeds mean internal caching is more time consuming. We recommend caching the full <code>prepInputs</code> call instead (e.g. <code>prepInputs(...) &gt;Cache()</code> ).
<code>overwrite</code>	Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.
<code>action</code>	A character string, with one of <code>c("nothing", "update", "replace", "append")</code> . Partial matching is used ("n" is sufficient). <code>nothing</code> will prevent any updating of the <code>targetFile</code> , i.e., "read only". <code>append</code> will add the spatial elements in <code>domain</code> to <code>targetFile</code> (and writing it back to disk). <code>update</code> will do the same as <code>append</code> , but will also remove any identical geometries before appending. <code>replace</code> does nothing currently.
<code>...</code>	Any named objects that are needed for <code>FUN</code>

**Details**

This function is a combination of `Cache` and `prepInputs` but for spatial domains. This differs from `Cache` in that the current function call doesn't have to have an identical function call previously

run. Instead, it needs to have had a previous function call where the domain being passes is *within* the geographic limits of the `targetFile` or file located at the `url`. This is similar to a geospatial operation on a remote GIS server, with 2 differences:

1. This downloads the object first before doing the GIS locally, and 2. it will optionally upload an updated object if the geographic area did not yet exist.

This has a very specific use case: assess whether an existing `sf` polygon or multipolygon object (local or remote) covers the spatial area of a domain of interest. If it does, then return only that part of the `sf` object that completely covers the domain. If it does not, then run FUN. It is expected that FUN will produce an `sf` polygon or multipolygon class object. The result of FUN will then be appended to the `sf` object as a new entry (feature) or it will replace the existing "same extent" entry in the `sf` object.

### Value

Returns an object that results from FUN, which will possibly be a subset of a larger spatial object that is specified with `targetFile` or `url`.

### Examples

```
if (requireNamespace("sf", quietly = TRUE) &&
    requireNamespace("terra", quietly = TRUE)) {
  dPath <- checkPath(file.path(tempdir2()), create = TRUE)
  localFileLux <- system.file("ex/lux.shp", package = "terra")

  # 1 step for each layer
  # 1st step -- get study area
  full <- prepInputs(localFileLux, destinationPath = dPath) # default is sf::st_read
  zoneA <- full[3:6, ]
  zoneB <- full[8, ] # not in A
  zoneC <- full[3, ] # yes in A
  zoneD <- full[7:8, ] # not in A, B or C
  zoneE <- full[3:5, ] # yes in A
  # 2nd step: re-write to disk as read/write is lossy; want all "from disk" for this ex.
  writeTo(zoneA, writeTo = "zoneA.shp", destinationPath = dPath)
  writeTo(zoneB, writeTo = "zoneB.shp", destinationPath = dPath)
  writeTo(zoneC, writeTo = "zoneC.shp", destinationPath = dPath)
  writeTo(zoneD, writeTo = "zoneD.shp", destinationPath = dPath)
  writeTo(zoneE, writeTo = "zoneE.shp", destinationPath = dPath)
  # Must re-read to get identical columns
  zoneA <- sf::st_read(file.path(dPath, "zoneA.shp"))
  zoneB <- sf::st_read(file.path(dPath, "zoneB.shp"))
  zoneC <- sf::st_read(file.path(dPath, "zoneC.shp"))
  zoneD <- sf::st_read(file.path(dPath, "zoneD.shp"))
  zoneE <- sf::st_read(file.path(dPath, "zoneE.shp"))

  # The function that is to be run. This example returns a data.frame because
  #   saving `sf` class objects with list-like columns does not work with
  #   many st_driver()
  fun <- function(domain, newField) {
```

```

domain |>
  as.data.frame() |>
  cbind(params = I(lapply(seq_len(NROW(domain)), function(x) newField)))
}

# Run sequence -- A, B will add new entries in targetFile, C will not,
#                               D will, E will not
for (z in list(zoneA, zoneB, zoneC, zoneD, zoneE)) {
  out <- CacheGeo(
    targetFile = "fireSenseParams.rds",
    domain = z,
    FUN = fun(domain, newField = I(list(list(a = 1, b = 1:2, c = "D")))),
    fun = fun, # pass whatever is needed into the function
    destinationPath = dPath,
    action = "update"
    # , cloudFolderID = "cachedObjects" # to upload/download from cloud
  )
}
}

```

---

```
checkAndMakeCloudFolderID
```

*Check for presence of checkFolderID (for Cache(useCloud))*

---

### Description

Will check for presence of a cloudFolderID and make a new one if one not present on Google Drive, with a warning.

### Usage

```

checkAndMakeCloudFolderID(
  cloudFolderID = getOption("reproducible.cloudFolderID", NULL),
  cachePath = NULL,
  create = FALSE,
  overwrite = FALSE,
  verbose = getOption("reproducible.verbose", 1),
  team_drive = NULL
)

```

### Arguments

cloudFolderID	The google folder ID where cloud caching will occur.
cachePath	A repository used for storing cached objects. This is optional if Cache is used inside a SpaDES module.
create	Logical. If TRUE, then the cloudFolderID will be created. This should be used with caution as there are no checks for overwriting. See <code>googledrive::drive_mkdir</code> . Default FALSE.

overwrite	Logical. Passed to googledrive::drive_mkdir.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
team_drive	Logical indicating whether to check team drives.

**Value**

Returns the character string of the cloud folder ID created or reported

---

checkPath	<i>Check directory path</i>
-----------	-----------------------------

---

**Description**

Checks the specified path to a directory for formatting consistencies, such as trailing slashes, etc.

**Usage**

```
checkPath(path, create)

## S4 method for signature 'character,logical'
checkPath(path, create)

## S4 method for signature 'character,missing'
checkPath(path)

## S4 method for signature 'NULL,ANY'
checkPath(path)

## S4 method for signature 'missing,ANY'
checkPath()
```

**Arguments**

path	A character string corresponding to a directory path.
create	A logical indicating whether the path should be created if it does not exist. Default is FALSE.

**Value**

Character string denoting the cleaned up filepath.

**Note**

This will not work for paths to files. To check for existence of files, use `file.exists()`. To normalize a path to a file, use `normPath()` or `normalizePath()`.

**See Also**

`file.exists()`, `dir.create()`, `normPath()`

**Examples**

```
## normalize file paths
paths <- list("./aaa/zzz",
             "./aaa/zzz/",
             "../aaa/zzz",
             "../aaa/zzz/",
             ".\\\\"aaa\\\\"zzz",
             ".\\\\"aaa\\\\"zzz\\\\"",
             file.path(".", "aaa", "zzz"))

checked <- normPath(paths)
length(unique(checked)) ## 1; all of the above are equivalent

## check to see if a path exists
tmpdir <- file.path(tempdir(), "example_checkPath")

dir.exists(tmpdir) ## FALSE
tryCatch(checkPath(tmpdir, create = FALSE), error = function(e) FALSE) ## FALSE

checkPath(tmpdir, create = TRUE)
dir.exists(tmpdir) ## TRUE

unlink(tmpdir, recursive = TRUE)
```

---

checkRelative

*An alternative to basename and dirname when there are sub-folders*

---

**Description**

This confirms that the files which may be absolute actually exist when compared `makeRelative(knownRelativeFiles, absolutePrefix)`. This is different than just using `basename` because it will include any sub-folder structure within the `knownRelativePaths`

**Usage**

```
checkRelative(
  files,
  absolutePrefix,
  knownRelativeFiles,
  verbose = getOption("reproducible.verbose")
)
```

**Arguments**

files	A character vector of files to check to see if they are the same as knownRelativeFiles, once the absolutePrefix is removed
absolutePrefix	A directory to "remove" from files to compare to knownRelativeFiles
knownRelativeFiles	A character vector of relative filenames, that could have sub-folder structure.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t

Checksums

*Calculate checksum***Description**

Verify (and optionally write) checksums. Checksums are computed using `.digest()`, which is simply a wrapper around `digest::digest`.

**Usage**

```
Checksums(
  path,
  write,
  quickCheck = getOption("reproducible.quickCheck", FALSE),
  checksumFile = identifyCHECKSUMStxtFile(path),
  files = NULL,
  verbose = getOption("reproducible.verbose", 1),
  ...
)

## S4 method for signature 'character,logical'
Checksums(
  path,
  write,
  quickCheck = getOption("reproducible.quickCheck", FALSE),
  checksumFile = identifyCHECKSUMStxtFile(path),
  files = NULL,
  verbose = getOption("reproducible.verbose", 1),
  ...
)

## S4 method for signature 'character,missing'
Checksums(
  path,
  write,
```



```

    quickCheck = getOption("reproducible.quickCheck", FALSE),
    checksumFile = identifyCHECKSUMStxtFile(path),
    files = NULL,
    verbose = getOption("reproducible.verbose", 1),
    ...
  )

```

### Arguments

path	Character string giving the directory path containing CHECKSUMS.txt file, or where it will be written if checksumFile = TRUE.
write	Logical indicating whether to overwrite CHECKSUMS.txt. Default is FALSE, as users should not change this file. Module developers should write this file prior to distributing their module code, and update accordingly when the data change.
quickCheck	Logical. If TRUE, then this will only use file sizes, rather than a digest::digest hash. This is generally faster, but will be <i>much</i> less robust.
checksumFile	The filename of the checksums file to read or write to. The default is 'CHECKSUMS.txt' located at file.path(path, module, "data", checksumFile). It is likely not a good idea to change this, and should only be used in cases such as Cache, which can evaluate if the checksumFile has changed.
files	An optional character string or vector of specific files to checksum. This may be very important if there are many files listed in a CHECKSUMS.txt file, but only a few are to be checksummed.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
...	Passed to <code>digest::digest()</code> and <code>utils::write.table()</code> . For <code>digest</code> , the notable argument is <code>algo</code> . For <code>write.table</code> , the notable argument is <code>append</code> .

### Value

A `data.table` with columns: `result`, `expectedFile`, `actualFile`, `checksum.x`, `checksum.y`, `algorithm.x`, `algorithm.y`, `filesize.x`, `filesize.y` indicating the result of comparison between local file (x) and expectation based on the CHECKSUMS.txt file.

### Note

In version 1.2.0 and earlier, two checksums per file were required because of differences in the checksum hash values on Windows and Unix-like platforms. Recent versions use a different (faster) algorithm and only require one checksum value per file. To update your 'CHECKSUMS.txt' files using the new algorithm, see <https://github.com/PredictiveEcology/SpaDES/issues/295#issuecomment-246513405>.

### Author(s)

Alex Chubaty

**Examples**

```

## Not run:
modulePath <- file.path(tempdir(), "myModulePath")
dir.create(modulePath, recursive = TRUE, showWarnings = FALSE)
moduleName <- "myModule"
cat("hi", file = file.path(modulePath, moduleName)) # put something there for this example

## verify checksums of all data files
Checksums(modulePath, files = moduleName)

## write new CHECKSUMS.txt file
Checksums(files = moduleName, modulePath, write = TRUE)

## End(Not run)

```

---

cloudDownload

*Download from cloud, if necessary*


---

**Description**

Meant for internal use, as there are internal objects as arguments.

**Usage**

```

cloudDownload(
  outputHash,
  newFileName,
  gdriveLs,
  cachePath,
  cloudFolderID,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose")
)

```

**Arguments**

outputHash	The cacheId of the object to upload
newFileName	The character string of the local filename that the downloaded object will have
gdriveLs	The result of googledrive::drive_ls(googledrive::as_id(cloudFolderID), pattern = "outputHash")
cachePath	A repository used for storing cached objects. This is optional if Cache is used inside a SpaDES module.

cloudFolderID	A googledrive dribble of a folder, e.g., using <code>drive_mkdir()</code> . If left as <code>NULL</code> , the function will create a cloud folder with name from last two folder levels of the <code>cachePath</code> path, : <code>paste0(basename(dirname(cachePath)), "_", basename(cachePath))</code> . This <code>cloudFolderID</code> will be added to <code>options("reproducible.cloudFolderID")</code> but this will not persist across sessions. If this is a character string, it will treat this as a folder name to create or use on GoogleDrive.
drv	if using a database backend, <code>drv</code> must be an object that inherits from <code>DBIDriver</code> e.g., from package <code>RSQLite</code> , e.g., <code>SQLite</code>
conn	an optional <code>DBIConnection</code> object, as returned by <code>dbConnect()</code> .
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t

---

compareNA	<i>NA-aware comparison of two vectors</i>
-----------	---

---

## Description

Copied from [http://www.cookbook-r.com/Manipulating\\_data/Comparing\\_vectors\\_or\\_factors\\_with\\_NA/](http://www.cookbook-r.com/Manipulating_data/Comparing_vectors_or_factors_with_NA/). This function returns `TRUE` wherever elements are the same, including `NA`'s, and `FALSE` everywhere else.

## Usage

```
compareNA(v1, v2)
```

## Arguments

<code>v1</code>	A vector
<code>v2</code>	A vector

## Value

A logical vector, indicating positions where two vectors are same or differ.

## Examples

```
a <- c(NA, 1, 2, NA)
b <- c(1, NA, 2, NA)
compareNA(a, b)
```

---

convertPaths	<i>Change the absolute path of a file</i>
--------------	---

---

### Description

convertPaths is simply a wrapper around gsub for changing the first part of a path. convertRasterPaths is useful for changing the path to a file-backed raster (e.g., after copying the file to a new location).

### Usage

```
convertPaths(x, patterns, replacements)
```

```
convertRasterPaths(x, patterns, replacements)
```

### Arguments

x	For convertPaths, a character vector of file paths. For convertRasterPaths, a disk-backed RasterLayer object, or a list of such rasters.
patterns	Character vector containing a pattern to match (see ?gsub).
replacements	Character vector of the same length of patterns containing replacement text (see ?gsub).

### Value

A normalized path with the patterns replaced by replacements. Or a list of such objects if x was a list.

### Author(s)

Eliot McIntire and Alex Chubaty

### Examples

```
filenames <- c("/home/user1/Documents/file.txt", "/Users/user1/Documents/file.txt")
oldPaths <- dirname(filenames)
newPaths <- c("/home/user2/Desktop", "/Users/user2/Desktop")
convertPaths(filenames, oldPaths, newPaths)
```

---

Copy	<i>Recursive copying of nested environments, and other "hard to copy" objects</i>
------	---

---

### Description

When copying environments and all the objects contained within them, there are no copies made: it is a pass-by-reference operation. Sometimes, a deep copy is needed, and sometimes, this must be recursive (i.e., environments inside environments).

### Usage

```
Copy(object, ...)

## S4 method for signature 'ANY'
Copy(
  object,
  filebackedDir,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  ...
)

## S4 method for signature 'data.table'
Copy(object, ...)

## S4 method for signature 'list'
Copy(object, ...)

## S4 method for signature 'refClass'
Copy(object, ...)

## S4 method for signature 'data.frame'
Copy(object, ...)
```

### Arguments

object	An R object (likely containing environments) or an environment.
...	Only used for custom Methods
filebackedDir	A directory to copy any files that are backing R objects, currently only valid for Raster classes. Defaults to <code>.reproducibleTempPath()</code> , which is unlikely to be very useful. Can be <code>NULL</code> , which means that the file will not be copied and could therefore cause a collision as the pre-copied object and post-copied object would have the same file backing them.

drv	if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite
conn	an optional DBIConnection object, as returned by dbConnect().
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t

### Details

To create a new Copy method for a class that needs its own method, try something like shown in example and put it in your package (or other R structure).

### Value

The same object as object, but with pass-by-reference class elements "deep" copied. reproducible has methods for several classes.

### Author(s)

Eliot McIntire

### See Also

[.robustDigest\(\)](#), [FileNames\(\)](#)

### Examples

```
e <- new.env()
e$abc <- letters
e$one <- 1L
e$lst <- list(W = 1:10, X = runif(10), Y = rnorm(10), Z = LETTERS[1:10])
ls(e)

# 'normal' copy
f <- e
ls(f)
f$one
f$one <- 2L
f$one
e$one ## uh oh, e has changed!

# deep copy
e$one <- 1L
g <- Copy(e)
ls(g)
g$one
g$one <- 3L
g$one
f$one
e$one
```

```
## To create a new deep copy method, use the following template
## setMethod("Copy", signature = "the class", # where = specify here if not in a package,
##         definition = function(object, filebackendDir, ...) {
##         # write deep copy code here
##         })
```

---

copySingleFile

*Copy a file using robocopy on Windows and rsync on Linux/macOS*


---

### Description

This is replacement for `file.copy`, but for one file at a time. The additional feature is that it will use `robocopy` (on Windows) or `rsync` on Linux or Mac, if they exist. It will default back to `file.copy` if none of these exists. If there is a possibility that the file already exists, then this function should be very fast as it will do "update only", i.e., nothing.

### Usage

```
copySingleFile(
  from = NULL,
  to = NULL,
  useRobocopy = TRUE,
  overwrite = TRUE,
  delDestination = FALSE,
  create = TRUE,
  silent = FALSE
)
```

```
copyFile(
  from = NULL,
  to = NULL,
  useRobocopy = TRUE,
  overwrite = TRUE,
  delDestination = FALSE,
  create = TRUE,
  silent = FALSE
)
```

### Arguments

<code>from</code>	The source file.
<code>to</code>	The new file.
<code>useRobocopy</code>	For Windows, this will use a system call to <code>robocopy</code> which appears to be much faster than the internal <code>file.copy</code> function. Uses <code>/MIR</code> flag. Default <code>TRUE</code> .
<code>overwrite</code>	Passed to <code>file.copy</code>

delDestination Logical, whether the destination should have any files deleted, if they don't exist in the source. This is /purge for robocopy and -delete for rsync.

create Passed to checkPath.

silent Should a progress be printed.

### Value

This function is called for its side effect, i.e., a file is copied from to to.

### Author(s)

Eliot McIntire and Alex Chubaty

### Examples

```
tmpDirFrom <- file.path(tempdir(), "example_fileCopy_from")
tmpDirTo <- file.path(tempdir(), "example_fileCopy_to")
tmpFile1 <- tempfile("file1", tmpDirFrom, ".csv")
tmpFile2 <- tempfile("file2", tmpDirFrom, ".csv")
dir.create(tmpDirFrom, recursive = TRUE, showWarnings = FALSE)
dir.create(tmpDirTo, recursive = TRUE, showWarnings = FALSE)
f1 <- normalizePath(tmpFile1, mustWork = FALSE)
f2 <- normalizePath(tmpFile2, mustWork = FALSE)
t1 <- normalizePath(file.path(tmpDirTo, basename(tmpFile1)), mustWork = FALSE)
t2 <- normalizePath(file.path(tmpDirTo, basename(tmpFile2)), mustWork = FALSE)

write.csv(data.frame(a = 1:10, b = runif(10), c = letters[1:10]), f1)
write.csv(data.frame(c = 11:20, d = runif(10), e = letters[11:20]), f2)
copyFile(c(f1, f2), c(t1, t2))
file.exists(t1) ## TRUE
file.exists(t2) ## TRUE
identical(read.csv(f1), read.csv(f2)) ## FALSE
identical(read.csv(f1), read.csv(t1)) ## TRUE
identical(read.csv(f2), read.csv(t2)) ## TRUE
```

---

createCache

*Low-level functions to create and work with a cache*

---

### Description

**These are intended for advanced use only.**

### Usage

```
createCache(
  cachePath = getOption("reproducible.cachePath"),
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
```



```
    force = FALSE,
    verbose = getOption("reproducible.verbose")
)

loadFromCache(
  cachePath = getOption("reproducible.cachePath"),
  cacheId,
  preDigest,
  fullCacheTableForObj = NULL,
  format = getOption("reproducible.cacheSaveFormat", "rds"),
  .functionName = NULL,
  .dotsFromCache = NULL,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose")
)

extractFromCache(sc, elem, ifNot = NULL)

rmFromCache(
  cachePath = getOption("reproducible.cachePath"),
  cacheId,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  format = getOption("reproducible.cacheSaveFormat", "rds")
)

CacheDBFile(
  cachePath = getOption("reproducible.cachePath"),
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL)
)

CacheStorageDir(cachePath = getOption("reproducible.cachePath"))

CacheStoredFile(
  cachePath = getOption("reproducible.cachePath"),
  cacheId,
  format = NULL,
  obj = NULL
)

CacheDBTableName(
  cachePath = getOption("reproducible.cachePath"),
  drv = getDrv(getOption("reproducible.drv", NULL))
)

CacheIsACache(
```

```

cachePath = getOption("reproducible.cachePath"),
create = FALSE,
drv = getDrv(getOption("reproducible.drv", NULL)),
conn = getOption("reproducible.conn", NULL)
)

```

### Arguments

cachePath	A path describing the directory in which to create the database file(s)
drv	A driver, passed to dbConnect
conn	an optional DBIConnection object, as returned by dbConnect().
force	Logical. Should it create a cache in the cachePath, even if it already exists, overwriting.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
cacheId	The cacheId or otherwise digested hash value, as character string.
preDigest	The list of preDigest that comes from CacheDigest of an object
fullCacheTableForObj	The result of showCache, but subsetted for only the cacheId being loaded or selected
format	The text string representing the file extension used normally by different save formats; currently only "rds" or "qs". Defaults to getOption("reproducible.cacheSaveFormat", "rds")
.functionName	Optional. Used for messaging when this function is called from Cache
.dotsFromCache	Optional. Used internally.
sc	a cache tags data.table object
elem	character string specifying a tagKey value to match
ifNot	character (or NULL) specifying the return value to use if elem not matched
obj	The optional object that is of interest; it may have an attribute "saveRawFile" that would be important.
create	Logical. Currently only affects non <b>RSQLite</b> default drivers. If TRUE and there is no Cache database, the function will create one.

### Details

- createCache() will create a Cache folder structure and necessary files, based on the particular drv or conn provided;
- loadFromCache() retrieves a single object from the cache, given its cacheId;
- extractFromCache() retrieves a single tagValue from the cache based on the tagKey of elem;
- rmFromCache() removes one or more items from the cache, and updates the cache database files.

**Value**

- createCache() returns NULL (invisibly) and intended to be called for side effects;
- loadFromCache() returns the object from the cache that has the particular cacheId;
- extractFromCache() returns the tagValue from the cache corresponding to elem if found, otherwise the value of ifNot;
- rmFromCache() returns NULL (invisibly) and is intended to be called for side effects;
- CacheDBFile() returns the name of the database file for a given Cache, when useDBI() == FALSE, or NULL if TRUE;
- CacheDBFiles() (i.e., plural) returns the name of all the database files for a given Cache when useDBI() == TRUE, or NULL if FALSE;
- CacheStoredFile() returns the file path to the file with the specified hash value, This can be loaded to memory with e.g., [loadFile\(\)](#);
- CacheStorageDir() returns the name of the directory where cached objects are stored;
- CacheStoredFile returns the file path to the file with the specified hash value;
- CacheDBTableName() returns the name of the table inside the SQL database, if that is being used;
- CacheIsACache() returns a logical indicating whether the cachePath is currently a reproducible cache database;

**Examples**

```

data.table::setDTthreads(2)
newCache <- tempdir2()
createCache(newCache)

out <- Cache(rnorm(1), cachePath = newCache)
cacheId <- gsub("cacheId:", "", attr(out, "tags"))
loadFromCache(newCache, cacheId = cacheId)

rmFromCache(newCache, cacheId = cacheId)

# clean up
unlink(newCache, recursive = TRUE)

data.table::setDTthreads(2)
newCache <- tempdir2()

# Given the drv and conn, creates the minimum infrastructure for a cache
createCache(newCache)

CacheDBFile(newCache) # identifies the database file
CacheStorageDir(newCache) # identifies the directory where cached objects are stored

```

```

out <- Cache(rnorm(1), cachePath = newCache)
cacheId <- gsub("cacheId:", "", attr(out, "tags"))
CacheStoredFile(newCache, cacheId = cacheId)

# The name of the table inside the SQL database
CacheDBTableName(newCache)

CacheIsACache(newCache) # returns TRUE

# clean up
unlink(newCache, recursive = TRUE)

```

---

determineFilename      *Determine filename, either automatically or manually*

---

## Description

Determine the filename, given various combinations of inputs.

## Usage

```

determineFilename(
  filename2 = NULL,
  filename1 = NULL,
  destinationPath = getOption("reproducible.destinationPath", "."),
  verbose = getOption("reproducible.verbose", 1),
  prefix = "Small",
  ...
)

```

## Arguments

filename2	filename2 is optional, and is either NULL (no writing of outputs to disk), or several options for writing the object to disk. If TRUE (the default), it will give it a file name determined by <code>.prefix(basename(filename1), prefix)</code> . If a character string, it will use this as its file name. See <code>determineFilename()</code> .
filename1	Character strings giving the file paths of the <i>input</i> object (filename1) filename1 is only used for messaging (i.e., the object itself is passed in as x) and possibly naming of output (see details and filename2).
destinationPath	Optional. If filename2 is a relative file path, then this will be the directory of the resulting absolute file path.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t
prefix	The character string to prepend to filename1, if filename2 not provided.
...	Passed into <code>writeTo()</code>

## Details

The post processing workflow, which includes this function, addresses several scenarios, and depending on which scenario, there are several file names at play. For example, Raster objects may have file-backed data, and so *possess a file name*, whereas Spatial objects do not. Also, if post processing is part of a `prepInputs()` workflow, there will always be a file downloaded. From the perspective of `postProcess`, these are the "inputs" or `filename1`. Similarly, there may or may not be a desire to write an object to disk after all post processing, `filename2`.

This subtlety means that there are two file names that may be at play: the "input" file name (`filename1`), and the "output" filename (`filename2`). When this is used within `postProcess`, it is straight forward.

However, when `postProcess` is used within a `prepInputs` call, the `filename1` file is the file name of the downloaded file (usually automatically known following the downloading, and referred to as `targetFile`) and the `filename2` is the file name of the of post-processed file.

If `filename2` is `TRUE`, i.e., not an actual file name, then the cropped/masked raster will be written to disk with the original `filename1/targetFile` name, with `prefix` prefixed to the `basename(targetFile)`.

If `filename2` is a character string, it will be the path of the saved/written object e.g., passed to `writeOutput`. It will be tested whether it is an absolute or relative path and used as is if absolute or prepended with `destinationPath` if relative.

If `filename2` is `logical`, then the output filename will be `prefix` prefixed to the `basename(filename1)`. If a character string, it will be the path returned. It will be tested whether it is an absolute or relative path and used as is if absolute or prepended with `destinationPath` if provided, and if `filename2` is relative.

---

downloadFile

*A wrapper around a set of downloading functions*

---

## Description

Currently, this only deals with `googledrive::drive_download`, and `utils::download.file()`. In general, this is not intended for use by a user.

## Usage

```
downloadFile(
  archive,
  targetFile,
  neededFiles,
  destinationPath = getOption("reproducible.destinationPath", "."),
  quick,
  checksumFile,
  dlFun = NULL,
  checkSums,
  url,
  needChecksums,
```

```

preDigest,
overwrite = getOption("reproducible.overwrite", TRUE),
alsoExtract = "similar",
verbose = getOption("reproducible.verbose", 1),
purge = FALSE,
.tempPath,
...
)

```

## Arguments

archive	Optional character string giving the path of an archive containing <code>targetFile</code> , or a vector giving a set of nested archives (e.g., <code>c("xxx.tar", "inner.zip", "inner.rar")</code> ). If there is/are (an) inner archive(s), but they are unknown, the function will try all until it finds the <code>targetFile</code> . See table in <code>preProcess()</code> . If it is NA, then it will <i>not</i> attempt to see it as an archive, even if it has archive-like file extension (e.g., <code>.zip</code> ). This may be useful when an R function is expecting an archive directly.
targetFile	Character string giving the filename (without relative or absolute path) to the eventual file (raster, shapefile, csv, etc.) after downloading and extracting from a zip or tar archive. This is the file <i>before</i> it is passed to <code>postProcess</code> . The internal checksumming does not checksum the file after it is <code>postProcessed</code> (e.g., <code>cropped/reprojected/masked</code> ). Using <code>Cache</code> around <code>prepInputs</code> will do a sufficient job in these cases. See table in <code>preProcess()</code> .
neededFiles	Character string giving the name of the file(s) to be extracted.
destinationPath	Character string of a directory in which to download and save the file that comes from <code>url</code> and is also where the function will look for <code>archive</code> or <code>targetFile</code> . NOTE (still experimental): To prevent repeated downloads in different locations, the user can also set <code>options("reproducible.inputPaths")</code> to one or more local file paths to search for the file before attempting to download. Default for that option is NULL meaning do not search locally.
quick	Logical. This is passed internally to <code>Checksums()</code> (the <code>quickCheck</code> argument), and to <code>Cache()</code> (the <code>quick</code> argument). This results in faster, though less robust checking of inputs. See the respective functions.
checksumFile	A character string indicating the absolute path to the CHECKSUMS.txt file.
dlFun	Optional "download function" name, such as <code>"raster::getData"</code> , which does custom downloading, in addition to loading into R. Still experimental.
checkSums	A checksums file, e.g., created by <code>Checksums(..., write = TRUE)</code>
url	Optional character string indicating the URL to download from. If not specified, then no download will be attempted. If not entry exists in the CHECKSUMS.txt (in <code>destinationPath</code> ), an entry will be created or appended to. This CHECKSUMS.txt entry will be used in subsequent calls to <code>prepInputs</code> or <code>preProcess</code> , comparing the file on hand with the ad hoc CHECKSUMS.txt. See table in <code>preProcess()</code> .
needChecksums	A numeric, with 0 indicating do not write a new checksums, 1 write a new one, 2 append new information to existing one.

preDigest	The list of preDigest that comes from CacheDigest of an object
overwrite	Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.
alsoExtract	Optional character string naming files other than targetFile that must be extracted from the archive. If NULL, the default, then it will extract all files. Other options: "similar" will extract all files with the same filename without file extension as targetFile. NA will extract nothing other than targetFile. A character string of specific file names will cause only those to be extracted. See table in <a href="#">preProcess()</a> .
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t
purge	Logical or Integer. 0/FALSE (default) keeps existing CHECKSUMS.txt file and prepInputs will write or append to it. 1/TRUE will deleted the entire CHECKSUMS.txt file. Other options, see details.
.tempPath	Optional temporary path for internal file intermediate steps. Will be cleared on.exit from this function.
...	Passed to dlFun. Still experimental. Can be e.g., type for google docs.

**Value**

This function is called for its side effects, which will be a downloaded file (targetFile), placed in destinationPath. This file will be checksummed, and that checksum will be appended to the checksumFile.

**Author(s)**

Eliot McIntire

---

downloadRemote	<i>Download a remote file</i>
----------------	-------------------------------

---

**Description**

Download a remote file

**Usage**

```
downloadRemote(
  url,
  archive,
  targetFile,
  checkSums,
  dlFun = NULL,
```

```

fileToDownload,
messSkipDownload,
destinationPath,
overwrite,
needChecksums,
.tempPath,
preDigest,
alsoExtract = "similar",
verbose = getOption("reproducible.verbose", 1),
...
)

```

### Arguments

url	Optional character string indicating the URL to download from. If not specified, then no download will be attempted. If not entry exists in the CHECKSUMS.txt (in destinationPath), an entry will be created or appended to. This CHECKSUMS.txt entry will be used in subsequent calls to prepInputs or preProcess, comparing the file on hand with the ad hoc CHECKSUMS.txt. See table in <a href="#">preProcess()</a> .
archive	Optional character string giving the path of an archive containing targetFile, or a vector giving a set of nested archives (e.g., c("xxx.tar", "inner.zip", "inner.rar")). If there is/are (an) inner archive(s), but they are unknown, the function will try all until it finds the targetFile. See table in <a href="#">preProcess()</a> . If it is NA, then it will <i>not</i> attempt to see it as an archive, even if it has archive-like file extension (e.g., .zip). This may be useful when an R function is expecting an archive directly.
targetFile	Character string giving the filename (without relative or absolute path) to the eventual file (raster, shapefile, csv, etc.) after downloading and extracting from a zip or tar archive. This is the file <i>before</i> it is passed to postProcess. The internal checksumming does not checksum the file after it is postProcessed (e.g., cropped/reprojected/masked). Using Cache around prepInputs will do a sufficient job in these cases. See table in <a href="#">preProcess()</a> .
checkSums	TODO
dIFun	Optional "download function" name, such as "raster::getData", which does custom downloading, in addition to loading into R. Still experimental.
fileToDownload	TODO
messSkipDownload	The character string text to pass to messaging if download skipped
destinationPath	Character string of a directory in which to download and save the file that comes from url and is also where the function will look for archive or targetFile. NOTE (still experimental): To prevent repeated downloads in different locations, the user can also set options("reproducible.inputPaths") to one or more local file paths to search for the file before attempting to download. Default for that option is NULL meaning do not search locally.
overwrite	Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.



needChecksums	Logical indicating whether to generate checksums. <b>## TODO:</b> add overwrite arg to the function?
.tempPath	Optional temporary path for internal file intermediate steps. Will be cleared on.exit from this function.
preDigest	The list of preDigest that comes from CacheDigest of an object
alsoExtract	Optional character string naming files other than targetFile that must be extracted from the archive. If NULL, the default, then it will extract all files. Other options: "similar" will extract all files with the same filename without file extension as targetFile. NA will extract nothing other than targetFile. A character string of specific file names will cause only those to be extracted. See table in <a href="#">preProcess()</a> .
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
...	Additional arguments passed to <a href="#">postProcess()</a> and <a href="#">Cache()</a> . Since ... is passed to <a href="#">postProcess()</a> , these will ... will also be passed into the inner functions, e.g., <a href="#">cropInputs()</a> . Possibly useful other arguments include dlFun which is passed to preProcess. See details and examples.

---

extractFromArchive      *Extract files from archive*

---

## Description

Extract zip or tar archive files, possibly nested in other zip or tar archives.

## Usage

```
extractFromArchive(
  archive,
  destinationPath = getOption("reproducible.destinationPath", dirname(archive)),
  neededFiles = NULL,
  extractedArchives = NULL,
  checkSums = NULL,
  needChecksums = 0,
  filesExtracted = character(),
  checksumFilePath = character(),
  quick = FALSE,
  verbose = getOption("reproducible.verbose", 1),
  .tempPath,
  ...
)
```

**Arguments**

archive	Character string giving the path of the archive containing the file to be extracted. This path must exist or be NULL
destinationPath	Character string giving the path where neededFiles will be extracted. Defaults to the archive directory.
neededFiles	Character string giving the name of the file(s) to be extracted.
extractedArchives	Used internally to track archives that have been extracted from.
checkSums	A checksums file, e.g., created by Checksums(..., write = TRUE)
needChecksums	A numeric, with 0 indicating do not write a new checksums, 1 write a new one, 2 append new information to existing one.
filesExtracted	Used internally to track files that have been extracted.
checksumFilePath	The full path to the checksum.txt file
quick	Passed to Checksums
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
.tempPath	Optional temporary path for internal file intermediate steps. Will be cleared on.exit from this function.
...	Passed to unzip or untar, e.g., overwrite

**Value**

A character vector listing the paths of the extracted archives.

**Author(s)**

Jean Marchal and Eliot McIntire

---

fastMask	<i>Faster operations on rasters (DEPRECATED because terra::mask is fast)</i>
----------	--

---

**Description**

Deprecated. Use `maskTo()`.

**Usage**

```
fastMask(
  x,
  y,
  cores = NULL,
  useGDAL = FALSE,
  verbose = getOption("reproducible.verbose", 1),
  ...,
  messageSkipDeprecated = FALSE
)
```

**Arguments**

x	A Raster* object.
y	A SpatialPolygons object. If it is not in the same projection as x, it will be reprojected on the fly to that of x
cores	An integer* or 'AUTO'. This will be used if gdalwarp is triggered. 'AUTO' will calculate 90% of the total number of cores in the system, while an integer or rounded float will be passed as the exact number of cores to be used.
useGDAL	Deprecated. Logical or "force". This is defunct; internals now can use terra if options("reproducible.useTerra" = TRUE), which is not (yet) the default.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
...	Currently unused.
messageSkipDeprecated	Logical. If TRUE, then the message about this function being deprecated will be suppressed.

**Value**

A Raster\* object, masked (i.e., smaller extent and/or several pixels converted to NA)

**Author(s)**

Eliot McIntire

---

FileNames *Return the filename(s) from a Raster\* object*

---

**Description**

This is mostly just a wrapper around filename from the raster package, except that instead of returning an empty string for a RasterStack object, it will return a vector of length >1 for RasterStack.

**Usage**

```
FileNames(obj, allowMultiple = TRUE, returnList = FALSE)

## S4 method for signature 'ANY'
FileNames(obj, allowMultiple = TRUE, returnList = FALSE)

## S4 method for signature 'environment'
FileNames(obj, allowMultiple = TRUE, returnList = FALSE)

## S4 method for signature 'list'
FileNames(obj, allowMultiple = TRUE, returnList = FALSE)

## S4 method for signature 'data.table'
FileNames(obj, allowMultiple = TRUE, returnList = FALSE)

## S4 method for signature 'Path'
FileNames(obj, allowMultiple = TRUE, returnList = FALSE)
```

**Arguments**

<code>obj</code>	A Raster* object (i.e., RasterLayer, RasterStack, RasterBrick)
<code>allowMultiple</code>	Logical. If TRUE, the default, then all relevant filenames will be returned, i.e., in cases such as .grd where multiple files are required. If FALSE, then only the first file will be returned, e.g., filename.grd, in the case of default Raster format in R.
<code>returnList</code>	Default FALSE. If FALSE, then return format will be a character vector. When TRUE, list or environment objects will return a list of character strings or vectors. When returned as a character vector, then the names of objects with >1 filename associated with them will be given a numeric suffix, which means the name in the returned vector does not match the object in the list or environment. When returned as a list, their names are preserved.

**Details**

New methods can be made for this generic.

**Value**

A character vector of filenames that are part of the objects passed to `obj`. This returns NULL is the object is not file-backed or does not have a method to recover the file-backed filename.

**Author(s)**

Eliot McIntire

---

fixErrorsIn	<i>Fix common errors in GIS layers, using terra</i>
-------------	---

---

### Description

Currently, this only tests for validity of a SpatVect file, then if there is a problem, it will run `terra::makeValid`

### Usage

```
fixErrorsIn(
  x,
  error = NULL,
  verbose = getOption("reproducible.verbose"),
  fromFnName = ""
)
```

### Arguments

<code>x</code>	The SpatStat or SpatVect object to try to fix.
<code>error</code>	The error message, e.g., coming from <code>try(...)</code>
<code>verbose</code>	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t
<code>fromFnName</code>	The function name that produced the error, e.g., <code>maskTo</code>

### Value

An object of the same class as `x`, but with some errors fixed via `terra::makeValid()`

---

gdalProject	<i>3-Step postProcess sequence for SpatRasters using gdalwarp</i>
-------------	---

---

### Description

`gdalProject` is a thin wrapper around `sf::gdal_utils('gdalwarp', ...)` with specific options set, notably, `-r` to method (in the ...), `-t_srs` to the crs of the `toRas`, `-te` to the extent of the `toRas`, `-te_srs` to the crs of the `toRas`, `-dstnodata = NA`, and `-overwrite`.

`gdalResample` is a thin wrapper around `sf::gdal_utils('gdalwarp', ...)` with specific options set, notably, `"-r"`, `"near"`, `-te`, `-te_srs`, `tr`, `-dstnodata = NA`, `-overwrite`.

`gdalMask` is a thin wrapper around `sf::gdal_utils('gdalwarp', ...)` with specific options set, notably, `-cutline`, `-dstnodata = NA`, and `-overwrite`.

**Usage**

```

gdalProject(
    fromRas,
    toRas,
    filenameDest,
    verbose = getOption("reproducible.verbose"),
    ...
)

gdalResample(
    fromRas,
    toRas,
    filenameDest,
    verbose = getOption("reproducible.verbose"),
    ...
)

gdalMask(
    fromRas,
    maskToVect,
    writeTo = NULL,
    verbose = getOption("reproducible.verbose"),
    ...
)

```

**Arguments**

fromRas	see from argument from <a href="#">postProcessTo()</a> , but can only be a SpatRaster.
toRas	see to argument from <a href="#">postProcessTo()</a> , but can only be a SpatRaster.
filenameDest	A filename with an appropriate extension (e.g., .tif) for gdal to write the output to. Since this function is conceived to be part of a chain, and not the final step, this function does not use writeTo, which is reserved for the final step in the chain.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
...	For gdalProject, this can be method. For gdalMask can be destinationPath and touches. For all gdal*, this can also be and datatype.
maskToVect	see maskTo argument from <a href="#">maskTo()</a> , but can only be a SpatVector
writeTo	Optional character string of a filename to use writeRaster to save the final object. Default is NULL, which means there is no writeRaster

**Details**

These three functions are used within `postProcessTo`, in the sequence: `gdalProject`, `gdalResample` and `gdalMask`, when `from` and `projectTo` are `SpatRaster` and `maskTo` is a `SpatVector`, but only

if `options(reproducible.gdalwarp = TRUE)` is set.

This sequence is a slightly different order than the sequence when `gdalwarp = FALSE` or the arguments do not match the above. This sequence was determined to be faster and more accurate than any other sequence, including running all three steps in one `gdalwarp` call (which `gdalwarp` can do). Using one-step `gdalwarp` resulted in very coarse pixelation when converting from a coarse resolution to fine resolution, which visually was inappropriate in test cases.

### See Also

[gdalResample\(\)](#), and [gdalMask\(\)](#) and the overarching [postProcessTo\(\)](#)

### Examples

```
# prepare dummy data -- 3 SpatRasters, 2 SpatVectors
# need 2 SpatRaster
rf <- system.file("ex/elev.tif", package = "terra")
elev1 <- terra::rast(rf)
#'
ras2 <- terra::deepcopy(elev1)
ras2[ras2 > 200 & ras2 < 300] <- NA_integer_
terra::values(elev1) <- rep(1L, terra::ncell(ras2))
#'
# a polygon vector
f <- system.file("ex/lux.shp", package = "terra")
vOrig <- terra::vect(f)
v <- vOrig[1:2, ]
#'
utm <- terra::crs("epsg:23028") # $wkt
vInUTM <- terra::project(vOrig, utm)
vAsRasInLongLat <- terra::rast(vOrig, resolution = 0.008333333)
res100 <- 100
rInUTM <- terra::rast(vInUTM, resolution = res100, vals = 1)
# crop, reproject, mask, crop a raster with a vector in a different projection
# --> gives message about not enough information
t1 <- postProcessTo(elev1, to = vInUTM)
# crop, reproject, mask a raster to a different projection, then mask
t2a <- postProcessTo(elev1, to = vAsRasInLongLat, maskTo = vInUTM)

# using gdal directly --> slightly different mask
opts <- options(reproducible.gdalwarp = TRUE)
t2b <- postProcessTo(elev1, to = vAsRasInLongLat, maskTo = vInUTM)
t3b <- postProcessTo(elev1, to = rInUTM, maskTo = vInUTM)
options(opts)
```

**Description**

Extracting relative file paths.

**Usage**

```
getRelative(path, relativeToPath)
```

```
makeRelative(files, absoluteBase)
```

**Arguments**

path                    character vector or list specifying file paths  
relativeToPath        directory against which path will be relativized.  
files                   character vector or list specifying file paths  
absoluteBase         base directory (as absolute path) to prepend to files

**Details**

- `getRelative()` searches path "from the right" (instead of "from the left") and tries to reconstruct it relative to directory specified by `relativeToPath`. This is useful when dealing with symlinked paths.
- `makeRelative()` checks to see if files and `normPath(absoluteBase)` share a common path (i.e., "from the left"), otherwise it returns files.

**Examples**

```
## create a project directory (e.g., on a hard drive)
(tmp1 <- tempdir2("myProject", create = TRUE))

## create a cache directory elsewhere (e.g., on an SSD)
(tmp2 <- tempdir2("my_cache", create = TRUE))

## symlink the project cache directory to tmp2
## files created here are actually stored in tmp2
prjCache <- file.path(tmp1, "cache")
file.symlink(tmp2, prjCache)

## create a dummy cache object file in the project cache dir
(tmpf <- tempfile("cache_", prjCache))
cat(rnorm(100), file = tmpf)
file.exists(tmpf)
normPath(tmpf) ## note the 'real' location (i.e., symlink resolved)

getRelative(tmpf, prjCache) ## relative path
getRelative(tmpf, tmp2) ## relative path

makeRelative(tmpf, tmp2) ## abs path; tmpf and normPath(tmp2) don't share common path
makeRelative(tmpf, prjCache) ## abs path; tmpf and normPath(tmp2) don't share common path
makeRelative(normPath(tmpf), prjCache) ## rel path; share common path when both normPath-ed
```



```
unlink(tmp1, recursive = TRUE)
unlink(tmp2, recursive = TRUE)
```

---

internetExists	<i>Checks for existed of a url or the internet using <a href="https://CRAN.R-project.org">https://CRAN.R-project.org</a></i>
----------------	--

---

### Description

A lightweight function that may be less reliable than more purpose built solutions such as checking a specific web page using `RCurl::url.exists`. However, this is slightly faster and is sufficient for many uses.

### Usage

```
internetExists()
urlExists(url)
```

### Arguments

`url` A url of the form `https://...` to test for existence.

### Value

Logical, TRUE if internet site exists, FALSE otherwise  
Logical, TRUE if internet site exists, FALSE otherwise.

---

isUpdated	<i>Has a cached object has been updated?</i>
-----------	--

---

### Description

Has a cached object has been updated?

### Usage

```
isUpdated(x)
```

### Arguments

`x` cached object

### Value

logical

---

keepOrigGeom	<i>Keep original geometries of sf objects</i>
--------------	---

---

### Description

When intersections occur, what was originally 2 polygons features can become LINESTRING and/or POINT and any COLLECTIONS or MULTI- versions of these. This function evaluates what the original geometry was and drops any newly created *different* geometries. For example, if a POLYGON becomes a COLLECTION of MULTIPOLYGON, POLYGON and POINT geometries, the POINT geometries will be dropped. This function is used internally in [postProcessTo\(\)](#).

### Usage

```
keepOrigGeom(newObj, origObj)
```

### Arguments

newObj	The new, derived sf object
origObj	The previous, object whose geometries should be used.

### Value

The original newObj, but with only the type of geometry that entered into the function.

---

linkOrCopy	<i>Hardlink, symlink, or copy a file</i>
------------	--

---

### Description

Attempt first to make a hardlink. If that fails, try to make a symlink (on non-windows systems and `symlink = TRUE`). If that fails, copy the file.

### Usage

```
linkOrCopy(
  from,
  to,
  symlink = TRUE,
  overwrite = TRUE,
  verbose = getOption("reproducible.verbose", 1)
)
```

**Arguments**

from, to	Character vectors, containing file names or paths. to can alternatively be the path to a single existing directory.
symlink	Logical indicating whether to use symlink (instead of hardlink). Default FALSE.
overwrite	Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t

**Value**

This function is called for its side effects, which will be a `file.link` is that is available or `file.copy` if not (e.g., the two directories are not on the same physical disk).

**Note**

Use caution with files-backed objects (e.g., rasters). See examples.

**Author(s)**

Alex Chubaty and Eliot McIntire

**See Also**

[file.link\(\)](#), [file.symlink\(\)](#), [file.copy\(\)](#).

**Examples**

```
tmpDir <- file.path(tempdir(), "symlink-test")
tmpDir <- normalizePath(tmpDir, winslash = "/", mustWork = FALSE)
dir.create(tmpDir)

f0 <- file.path(tmpDir, "file0.csv")
write.csv(iris, f0)

d1 <- file.path(tmpDir, "dir1")
dir.create(d1)
write.csv(iris, file.path(d1, "file1.csv"))

d2 <- file.path(tmpDir, "dir2")
dir.create(d2)
f2 <- file.path(tmpDir, "file2.csv")

## create link to a file
linkOrCopy(f0, f2)
file.exists(f2) ## TRUE
identical(read.table(f0), read.table(f2)) ## TRUE
```

```

## deleting the link shouldn't delete the original file
unlink(f0)
file.exists(f0) ## FALSE
file.exists(f2) ## TRUE

if (requireNamespace("terra", quietly = TRUE)) {
  ## using spatRasters and other file-backed objects
  f3a <- system.file("ex/test.grd", package = "terra")
  f3b <- system.file("ex/test.gri", package = "terra")
  r3a <- terra::rast(f3a)
  f4a <- file.path(tmpDir, "raster4.grd")
  f4b <- file.path(tmpDir, "raster4.gri")
  linkOrCopy(f3a, f4a) ## hardlink
  linkOrCopy(f3b, f4b) ## hardlink
  r4a <- terra::rast(f4a)

  isTRUE(all.equal(r3a, r4a)) # TRUE

  ## cleanup
  unlink(tmpDir, recursive = TRUE)
}

```

---

listNamed

---

*Create a list with names from object names*


---

## Description

This is a convenience wrapper around `newList <- list(a = 1); names(newList) <- "a"`.

## Usage

```
listNamed(...)
```

## Arguments

... Any elements to add to a list, as in `base::list`

## Details

This will return a named list, where names are the object names, captured internally in the function and assigned to the list. If a user manually supplies names, these will be kept (i.e., not overwritten by the object name).

## Examples

```

a <- 1
b <- 2
d <- 3
(newList <- listNamed(a, b, dManual = d)) # "dManual" name kept

```

---

loadFile	<i>Load a file from the cache</i>
----------	-----------------------------------

---

**Description**

Load a file from the cache

**Usage**

```
loadFile(file, format = NULL)
```

**Arguments**

file	character specifying the path to the file
format	(optional) character string specifying file extension ("qs" or "rds") of file; if not specified (i.e., NULL), will be deduced from the file extension of file.

**Value**

the object loaded from file

---

mergeCache	<i>Merge two cache repositories together</i>
------------	--

---

**Description****Usage**

```
mergeCache(
  cacheTo,
  cacheFrom,
  drvTo = getDrv(getOption("reproducible.drv", NULL)),
  drvFrom = getDrv(getOption("reproducible.drv", NULL)),
  connTo = NULL,
  connFrom = NULL,
  verbose = getOption("reproducible.verbose")
)
```

```
## S4 method for signature 'ANY'
```

```
mergeCache(
  cacheTo,
  cacheFrom,
  drvTo = getDrv(getOption("reproducible.drv", NULL)),
```

```

    drvFrom = getDrv(getOption("reproducible.drv", NULL)),
    connTo = NULL,
    connFrom = NULL,
    verbose = getOption("reproducible.verbose")
)

```

### Arguments

cacheTo	The cache repository (character string of the file path) that will become larger, i.e., merge into this
cacheFrom	The cache repository (character string of the file path) from which all objects will be taken and copied from
drvTo	The database driver for the cacheTo.
drvFrom	The database driver for the cacheFrom
connTo	The connection for the cacheTo. If not provided, then a new one will be made from drvTo and cacheTo
connFrom	The database for the cacheFrom. If not provided, then a new one will be made from drvFrom and cacheFrom
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t

### Details

All the cacheFrom artifacts will be put into cacheTo repository. All userTags will be copied verbatim, including accessed, with 1 exception: date will be the current `Sys.time()` at the time of merging. The `createdDate` column will be similarly the current time of merging.

### Value

The character string of the path of cacheTo, i.e., not the objects themselves.

---

messageDF	<i>Use message with a consistent use of verbose</i>
-----------	---

---

### Description

This family has a consistent use of verbose allowing messages to be turned on or off or verbosity increased or decreased throughout the family of messaging in reproducible.

**Usage**

```
messageDF(  
  df,  
  round,  
  colour = NULL,  
  colnames = NULL,  
  indent = NULL,  
  verbose = getOption("reproducible.verbose"),  
  verboseLevel = 1,  
  appendLF = TRUE  
)  
  
messagePrepInputs(  
  ...,  
  appendLF = TRUE,  
  verbose = getOption("reproducible.verbose"),  
  verboseLevel = 1  
)  
  
messagePreProcess(  
  ...,  
  appendLF = TRUE,  
  verbose = getOption("reproducible.verbose"),  
  verboseLevel = 1  
)  
  
messageCache(  
  ...,  
  colour = getOption("reproducible.messageColourCache"),  
  verbose = getOption("reproducible.verbose"),  
  verboseLevel = 1,  
  appendLF = TRUE  
)  
  
messageQuestion(..., verboseLevel = 0, appendLF = TRUE)  
  
.messageFunctionFn(  
  ...,  
  appendLF = TRUE,  
  verbose = getOption("reproducible.verbose"),  
  verboseLevel = 1  
)  
  
messageColoured(  
  ...,  
  colour = NULL,  
  indent = NULL,  
  hangingIndent = TRUE,
```

```

    verbose = getOption("reproducible.verbose", 1),
    verboseLevel = 1,
    appendLF = TRUE
  )

```

### Arguments

df	A data.frame, data.table, matrix
round	An optional numeric to pass to round
colour	Any colour that can be understood by cli
colnames	Logical or NULL. If TRUE, then it will print column names even if there aren't any in the df (i.e., they will) be V1 etc., NULL will print them if they exist, and FALSE which will omit them.
indent	An integer, indicating whether to indent each line
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
verboseLevel	The numeric value for this message* call, equal or above which verbose must be. The higher this is set, the more unlikely the call will show a message.
appendLF	logical: should messages given as a character string have a newline appended?
...	Any character vector, passed to paste0(...)
hangingIndent	Logical. If there are \n, should there be a hanging indent of 2 spaces. Default is TRUE

### Details

- messageDF uses message to print a clean square data structure.
- messageColoured allows specific colours to be used.
- messageQuestion sets a high level for verbose so that the message always gets asked.

### Value

Used for side effects. This will produce a message of a structured data.frame.

---

minFn	<i>Get min or maximum value of a (Spat)Raster</i>
-------	---

---

### Description

During the transition from raster to terra, some functions are not drop in replacements, such as minValue and maxValue became terra::minmax. This helper allows one function to be used, which calls the correct max or min function, depending on whether the object is a Raster or SpatRaster.



**Usage**

```

minFn(x)

maxFn(x)

dataType2(x, ...)

nLayers2(x)

values2(x, ...)

```

**Arguments**

```

x          A Raster or SpatRaster object.
...       Passed to the functions in raster or terra, as needed.

```

**Value**

A vector (not matrix as in `terra::minmax`) with the minimum or maximum value on the Raster or SpatRaster, one value per layer.

**Examples**

```

if (requireNamespace("terra", quietly = TRUE)) {
  ras <- terra::rast(terra::ext(0, 10, 0, 10), vals = 1:100)
  maxFn(ras)
  minFn(ras)
}

```

---

movedCache

*Deal with moved cache issues*

---

**Description**

If a user manually copies a complete Cache folder (including the db file and rasters folder), there are issues that must be addressed, depending on the Cache backend used. If using DBI (e.g., RSQLite or Postgres), the db table must be renamed. Run this function after a manual copy of a cache folder. See examples for one way to do that.

**Usage**

```

movedCache(
  new,
  old,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose")
)

```

**Arguments**

new	Either the path of the new cachePath where the cache was moved or copied to, or the new DB Table Name
old	Optional, if there is only one table in the new cache path. Either the path of the previous cachePath where the cache was moved or copied from, or the old DB Table Name
drv	if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite
conn	an optional DBIConnection object, as returned by dbConnect().
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t

**Details**

When the backend database for a reproducinle cache is an SQL database, the files on disk cannot be copied manually to a new location because they contain internal tables. Because reproducible gives the main table a name based on the cachePath path, calls to Cache will attempt to call this internally if it detects a name mismatch.

**Value**

movedCache does not return anything; it is called for its side effects.

**Examples**

```
data.table::setDTthreads(2)
tmpdir <- "tmpdir"
tmpCache <- "tmpCache"
tmpCacheDir <- normalizePath(file.path(tmpdir(), tmpCache), mustWork = FALSE)
tmpdirPath <- normalizePath(file.path(tmpdir(), tmpdir), mustWork = FALSE)
bb <- Cache(rnorm, 1, cachePath = tmpCacheDir)

# Copy all files from tmpCache to tmpdir
froms <- normalizePath(dir(tmpCacheDir, recursive = TRUE, full.names = TRUE),
  mustWork = FALSE
)
dir.create(file.path(tmpdirPath, "rasters"), recursive = TRUE, showWarnings = FALSE)
dir.create(file.path(tmpdirPath, "cacheOutputs"), recursive = TRUE, showWarnings = FALSE)
file.copy(
  from = froms, overwrite = TRUE,
  to = gsub(tmpCache, tmpdir, froms)
)

# Can use 'movedCache' to update the database table, though will generally
# happen automatically, with message indicating so
movedCache(new = tmpdirPath, old = tmpCacheDir)
bb <- Cache(rnorm, 1, cachePath = tmpdirPath) # should recover the previous call
```

---

normPath	<i>Normalize file paths</i>
----------	-----------------------------

---

### Description

Checks the specified path for formatting consistencies:

1. use slash instead of backslash;
2. do tilde etc. expansion;
3. remove trailing slash.

### Usage

```
normPath(path)

## S4 method for signature 'character'
normPath(path)

## S4 method for signature 'list'
normPath(path)

## S4 method for signature 'NULL'
normPath(path)

## S4 method for signature 'missing'
normPath()

## S4 method for signature 'logical'
normPath(path)

normPathRel(path)
```

### Arguments

path            A character vector of filepaths.

### Details

Additionally, normPath() attempts to create a absolute paths, whereas normPathRel() maintains relative paths.

```
d> getwd()
[1] "/home/achubaty/Documents/GitHub/PredictiveEcology/reproducible"
d> normPathRel("potato/chips")
[1] "potato/chips"
d> normPath("potato/chips")
[1] "/home/achubaty/Documents/GitHub/PredictiveEcology/reproducible/potato/chips"
```

**Value**

Character vector of cleaned up filepaths.

**Examples**

```
## normalize file paths
paths <- list("./aaa/zzz",
             "./aaa/zzz/",
             "../aaa/zzz",
             "../aaa/zzz/",
             ".\\\\"aaa\\\\"zzz",
             ".\\\\"aaa\\\\"zzz\\\\"",
             file.path(".", "aaa", "zzz"))

checked <- normPath(paths)
length(unique(checked)) ## 1; all of the above are equivalent

## check to see if a path exists
tmpdir <- file.path(tempdir(), "example_checkPath")

dir.exists(tmpdir) ## FALSE
tryCatch(checkPath(tmpdir, create = FALSE), error = function(e) FALSE) ## FALSE

checkPath(tmpdir, create = TRUE)
dir.exists(tmpdir) ## TRUE

unlink(tmpdir, recursive = TRUE)
```

---

objSize

*Wrapper around* lobstr::obj\_size

---

**Description**

This function attempts to estimate the real object size of an object. If the object has pass-by-reference semantics, it may not estimate the object size well without a specific method developed. For the case of terra class objects, this will be accurate (both RAM and file size), but only if it is not passed inside a list or environment. To get an accurate size of these, they should be passed individually.

**Usage**

```
objSize(x, quick = FALSE, ...)
```

```
objSizeSession(sumLevel = Inf, enclosingEnvs = TRUE, .prevEnvs = list())
```

**Arguments**

x	An object
quick	Logical. If FALSE, then an attribute, "objSize" will be added to the returned value, with each of the elements' object size returned also.
...	Additional arguments (currently unused), enables backwards compatible use.
sumLevel	Numeric, indicating at which depth in the list of objects should the object sizes be summed (summarized). Default is Inf, meaning no sums. Currently, the only option other than Inf is 1: objSizeSession(1), which gives the size of each package.
enclosingEnvs	Logical indicating whether to include enclosing environments. Default TRUE.
.prevEnvs	For internal account keeping to identify and prevent duplicate counting

**Details**

For functions, a user can include the enclosing environment as described <https://www.r-bloggers.com/2015/03/using-closures-as-objects-in-r/> and <http://adv-r.had.co.nz/memory.html>. It is not entirely clear which estimate is better. However, if the enclosing environment is the .GlobalEnv, it will not be included even though enclosingEnvs = TRUE.

objSizeSession will give the size of the whole session, including loaded packages. Because of the difficulties in calculating the object size of base and methods packages and Autoloads, these are omitted.

**Value**

This will return the result from lobstr::obj\_size, i.e., a lobstr\_bytes which is a numeric. If quick = FALSE, it will also have an attribute, "objSize", which will be a list with each element being the objSize of the individual elements of x. This is particularly useful if x is a list or environment. However, because of the potential for shared memory, the sum of the individual elements will generally not equal the value returned from this function.

**Examples**

```
library(utils)

foo <- new.env()
foo$b <- 1:10
foo$d <- 1:10

objSize(foo) # all the elements in the environment
utils::object.size(foo) # different - only measuring the environment as an object

utils::object.size(prepareInputs) # only the function, without its enclosing environment
objSize(prepareInputs) # the function, plus its enclosing environment

os1 <- utils::object.size(as.environment("package:reproducible"))
(os1) # very small -- just the environment container
```

---

paddedFloatToChar      *Convert numeric to character with padding*

---

### Description

This will pad floating point numbers, right or left. For integers, either class integer or functionally integer (e.g., 1.0), it will not pad right of the decimal. For more specific control or to get exact padding right and left of decimal, try the `stringi` package. It will also not do any rounding. See examples.

### Usage

```
paddedFloatToChar(x, padL = ceiling(log10(x + 1)), padR = 3, pad = "0")
```

### Arguments

<code>x</code>	numeric. Number to be converted to character with padding
<code>padL</code>	numeric. Desired number of digits on left side of decimal. If not enough, pad will be used to pad.
<code>padR</code>	numeric. Desired number of digits on right side of decimal. If not enough, pad will be used to pad.
<code>pad</code>	character to use as padding ( <code>nchar(pad) == 1</code> must be TRUE).

### Value

Character string representing the filename.

### Author(s)

Eliot McIntire and Alex Chubaty

### Examples

```
paddedFloatToChar(1.25)
paddedFloatToChar(1.25, padL = 3, padR = 5)
paddedFloatToChar(1.25, padL = 3, padR = 1) # no rounding, so keeps 2 right of decimal
```

---

 Path-class

*Coerce a character string to a class "Path"*


---

**Description**

Allows a user to specify that their character string is indeed a filepath. Thus, methods that require only a filepath can be dispatched correctly.

**Usage**

```
asPath(obj, nParentDirs = 0)

## S3 method for class 'character'
asPath(obj, nParentDirs = 0)

## S3 method for class 'null'
asPath(obj, nParentDirs = 0)
```

**Arguments**

<code>obj</code>	A character string to convert to a Path.
<code>nParentDirs</code>	A numeric indicating the number of parent directories starting from <code>basename(obj)</code> = 0 to keep for the digest

**Details**

It is often difficult or impossible to know algorithmically whether a character string corresponds to a valid filepath. In the case where it is an existing file, `file.exists` can work. But if it does not yet exist, e.g., for a save, it is difficult to know whether it is a valid path before attempting to save to the path.

This function can be used to remove any ambiguity about whether a character string is a path. It is primarily useful for achieving repeatability with Caching. Essentially, when Caching, arguments that are character strings should generally be digested verbatim, i.e., it must be an exact copy for the Cache mechanism to detect a candidate for recovery from the cache. Paths, are different. While they are character strings, there are many ways to write the same path. Examples of identical meaning, but different character strings are: path expanding of `~` vs. `not`, double back slash vs. single forward slash, relative path vs. absolute path. All of these should be assessed for their actual file or directory location, NOT their character string. By converting all character string that are actual file or directory paths with this function, then Cache will correctly assess the location, NOT the character string representation.

**Value**

A vector of class Path, which is similar to a character, but has an attribute indicating how deep the Path should be considered "digestible". In other words, most of the time, only some component of an absolute path is relevant for evaluating its purpose in a Cache situation. In general, this is usually equivalent to just the "relative" path

## Examples

```
tmpf <- tempfile(fileext = ".csv")
file.exists(tmpf) ## FALSE
tmpfPath <- asPath(tmpf)
is(tmpf, "Path") ## FALSE
is(tmpfPath, "Path") ## TRUE
```

---

postProcess

*Generic function to post process objects*

---

## Description

The method for GIS objects (terra `Spat*` & `sf` classes) will crop, reproject, and mask, in that order. This is a wrapper for `cropTo()`, `fixErrorsIn()`, `projectTo()`, `maskTo()` and `writeTo()`, with a required amount of data manipulation between these calls so that the crs match.

## Usage

```
postProcess(x, ...)

## S3 method for class 'list'
postProcess(x, ...)

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

## Arguments

`x` A GIS object of postProcessing, e.g., `Spat*` or `sf*`. This can be provided as a `rlang::quosure` or a normal R object.

`...` Additional arguments passed to methods. For `spatialClasses`, these are: `cropTo()`, `fixErrorsIn()`, `projectTo()`, `maskTo()`, `determineFilename()`, and `writeTo()`. Each of these may also pass `...` into other functions, like `writeTo()`. This might include potentially important arguments like `datatype`, `format`. Also passed to `terra::project`, with likely important arguments such as `method = "bilinear"`. See details.

## Value

A GIS file (e.g., `RasterLayer`, `SpatRaster` etc.) that has been appropriately cropped, reprojected, masked, depending on the inputs.



### Post processing sequence

If the rasterToMatch or studyArea are passed, then the following sequence will occur:

1. Fix errors `fixErrorsIn()`. Currently only errors fixed are for SpatialPolygons using `buffer(..., width = 0)`.
2. Crop using `cropTo()`
3. Project using `projectTo()`
4. Mask using `maskTo()`
5. Determine file name `determineFilename()`
6. Write that file name to disk, optionally `writeTo()`

NOTE: checksumming does not occur during the post-processing stage, as there are no file downloads. To achieve fast results, wrap prepInputs with Cache

### Backwards compatibility with rasterToMatch and/or studyArea arguments

For backwards compatibility, postProcess will continue to allow passing rasterToMatch and/or studyArea arguments. Depending on which of these are passed, different things will happen to the targetFile located at filename1.

See *Use cases* section in `postProcessTo()` for post processing behaviour with the new from and to arguments.

#### If targetFile is a raster (Raster\*, or SpatRaster) object::

	rasterToMatch	studyArea	Both
extent	Yes	Yes	rasterToMatch
resolution	Yes	No	rasterToMatch
projection	Yes	No*	rasterToMatch*
alignment	Yes	No	rasterToMatch
mask	No**	Yes	studyArea**

\*Can be overridden with useSAcrs.

\*\*Will mask with NAs from rasterToMatch if maskWithRTM.

#### If targetFile is a vector (Spatial\*, sf or SpatVector) object::

	rasterToMatch	studyArea	Both
extent	Yes	Yes	rasterToMatch
resolution	NA	NA	NA
projection	Yes	No*	rasterToMatch*
alignment	NA	NA	NA
mask	No	Yes	studyArea

\*Can be overridden with useSAcrs

**See Also**

prepInputs

**Examples**

```

if (requireNamespace("terra", quietly = TRUE) && requireNamespace("sf", quietly = TRUE)) {
  library(reproducible)
  od <- setwd(tempdir2())
  # download a (spatial) file from remote url (which often is an archive) load into R
  # need 3 files for this example; 1 from remote, 2 local
  dPath <- file.path(tempdir2())
  remoteTifUrl <- "https://github.com/rspatial/terra/raw/master/inst/ex/elev.tif"

  localFileLuxSm <- system.file("ex/luxSmall.shp", package = "reproducible")
  localFileLux <- system.file("ex/lux.shp", package = "terra")

  # 1 step for each layer
  # 1st step -- get study area
  studyArea <- prepInputs(localFileLuxSm, fun = "terra::vect") # default is sf::st_read

  # 2nd step: make the input data layer like the studyArea map
  # Test only relevant if connected to internet -- so using try just in case
  elevForStudy <- try(prepareInputs(url = remoteTifUrl, to = studyArea, res = 250,
    destinationPath = dPath, useCache = FALSE))

  # Alternate way, one step at a time. Must know each of these steps, and perform for each layer

  dir.create(dPath, recursive = TRUE, showWarnings = FALSE)
  file.copy(localFileLuxSm, file.path(dPath, basename(localFileLuxSm)))
  studyArea2 <- terra::vect(localFileLuxSm)
  if (!all(terra::is.valid(studyArea2))) studyArea2 <- terra::makeValid(studyArea2)
  tf <- tempfile(fileext = ".tif")
  download.file(url = remoteTifUrl, destfile = tf, mode = "wb", quiet = TRUE)
  Checksums(dPath, write = TRUE, files = tf)
  elevOrig <- terra::rast(tf)
  studyAreaCrs <- terra::crs(studyArea)
  elevForStudy2 <- terra::project(elevOrig, studyAreaCrs, res = 250) |>
    terra::mask(studyArea2) |>
    terra::crop(studyArea2)

  isTRUE(all.equal(elevForStudy, elevForStudy2)) # TRUE!

  # sf class
  studyAreaSmall <- prepInputs(localFileLuxSm)
  studyAreas <- list()
  studyAreas[["orig"]] <- prepInputs(localFileLux)
  studyAreas[["reprojected"]] <- projectTo(studyAreas[["orig"]], studyAreaSmall)
  studyAreas[["cropped"]] <- suppressWarnings(cropTo(studyAreas[["orig"]], studyAreaSmall))
  studyAreas[["masked"]] <- suppressWarnings(maskTo(studyAreas[["orig"]], studyAreaSmall))

  # SpatVector-- note: doesn't matter what class the "to" object is, only the "from"

```

```

studyAreas <- list()
studyAreas[["orig"]] <- prepInputs(localFileLux, fun = "terra::vect")
studyAreas[["reprojected"]] <- projectTo(studyAreas[["orig"]], studyAreaSmall)
studyAreas[["cropped"]] <- suppressWarnings(cropTo(studyAreas[["orig"]], studyAreaSmall))
studyAreas[["masked"]] <- suppressWarnings(maskTo(studyAreas[["orig"]], studyAreaSmall))
if (interactive()) {
  par(mfrow = c(2,2));
  out <- lapply(studyAreas, function(x) terra::plot(x))
}

setwd(od)
}

```

---

postProcessTo	<i>Transform a GIS dataset so it has the properties (extent, projection, mask) of another</i>
---------------	---

---

### Description

This function provides a single step to achieve the GIS operations "pre-crop-with-buffer-to-speed-up-projection", "project", "post-projection-crop", "mask" and possibly "write". It uses primarily the terra package internally (with some minor functions from sf) in an attempt to be as efficient as possible, except if all inputs are sf objects. (in which case sf is used). Currently, this function is tested with sf, SpatVector, SpatRaster, Raster\* and Spatial\* objects passed to from, and the same plus SpatExtent, and crs passed to to or the relevant \*to functions. For this function, Gridded means a Raster\* class object from raster or a SpatRaster class object from terra. Vector means a Spatial\* class object from sp, a sf class object from sf, or a SpatVector class object from terra. This function is also used internally with the deprecated family `postProcess()`, `*Inputs`, such as `cropInputs()`.

### Usage

```

postProcessTo(
  from,
  to,
  cropTo = NULL,
  projectTo = NULL,
  maskTo = NULL,
  writeTo = NULL,
  overwrite = TRUE,
  verbose = getOption("reproducible.verbose"),
  ...
)

postProcessTerra(
  from,
  to,
  cropTo = NULL,

```

```
projectTo = NULL,  
maskTo = NULL,  
writeTo = NULL,  
overwrite = TRUE,  
verbose = getOption("reproducible.verbose"),  
...  
)  
  
maskTo(  
  from,  
  maskTo,  
  overwrite = FALSE,  
  verbose = getOption("reproducible.verbose"),  
  ...  
)  
  
projectTo(  
  from,  
  projectTo,  
  overwrite = FALSE,  
  verbose = getOption("reproducible.verbose"),  
  ...  
)  
  
cropTo(  
  from,  
  cropTo = NULL,  
  needBuffer = FALSE,  
  overwrite = FALSE,  
  verbose = getOption("reproducible.verbose"),  
  ...  
)  
  
writeTo(  
  from,  
  writeTo,  
  overwrite = getOption("reproducible.overwrite"),  
  isStack = NULL,  
  isBrick = NULL,  
  isRaster = NULL,  
  isSpatRaster = NULL,  
  verbose = getOption("reproducible.verbose"),  
  ...  
)
```

**Arguments**

from	A Gridded or Vector dataset on which to do one or more of: crop, project, mask, and write
------	---

to	A Gridded or Vector dataset which is the object whose metadata will be the target for cropping, projecting, and masking of from.
cropTo	Optional Gridded or Vector dataset which, if supplied, will supply the extent with which to crop from. To omit cropping completely, set this to NA. If supplied, this will override to for the cropping step. Defaults to NULL, which means use to
projectTo	Optional Gridded or Vector dataset, or crs object (e.g., sf::st_crs). If Gridded it will supply the crs, extent, res, and origin to project the from to. If Vector, it will provide the crs only. The resolution and extent will be taken from res(from) (i.e. ncol(from)*nrow(from)). If a Vector, the extent of the projectTo is not used (unless it is also passed to cropTo. To omit projecting, set this to NA. If supplied, this will override to for the projecting step. Defaults to NULL, which means use to. <b>Attention.</b> Conflicts may arise with when projectTo is a Vector/CRS object with a distinct CRS from to. Because to is used for masking <i>after</i> from is re-projected using projectTo, the extents of to and from may no longer overlap (as in <i>align</i> ) perfectly leading to failure during the masking step. We recommend passing a raster templates to projectTo whose extent and CRS are both compatible with the object used later for masking (either to or maskTo).
maskTo	Optional Gridded or Vector dataset which, if supplied, will supply the extent with which to mask from. If Gridded, it will mask with the NA values on the maskTo; if Vector, it will mask on the terra::aggregate(maskTo). To omit masking completely, set this to NA. If supplied, this will override to for the masking step. Defaults to NULL, which means use to
writeTo	Optional character string of a filename to use writeRaster to save the final object. Default is NULL, which means there is no writeRaster
overwrite	Logical. Used if writeTo is not NULL; also if terra determines that the object requires writing to disk during a crop, mask or project call e.g., because it is too large.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
...	Arguments passed to terra::mask (for maskTo), terra::project (for projectTo) or terra::writeRaster (for writeTo) and not used for cropTo, as well postProcess's rasterToMatch and studyArea arguments (see below). Commonly used arguments might be method, touches, and datatype. If filename is passed, it will be ignored; use writeTo = . If reproducible.gdalwarp = TRUE, then these will be passed to the gdal* functions. See them for details.
needBuffer	Logical. Defaults to FALSE, meaning nothing is done out of the ordinary. If TRUE, then a buffer around the cropTo, so that if a reprojection has to happen on the cropTo prior to using it as a crop layer, then a buffer of 1.5 * res(cropTo) will occur prior, so that no edges are cut off.
isStack, isBrick, isRaster, isSpatRaster	Logical. Default NULL. Used to convert from back to these classes prior to writing, if provided.

## Details

postProcessTo is a wrapper around (an initial "wide" crop for speed) cropTo(needBuffer = TRUE), projectTo, cropTo (the actual crop for precision), maskTo, writeTo. Users can call each of these individually.

postProcessTerra is the early name of this function that is now postProcessTo.

This function is meant to replace `postProcess()` with the more efficient and faster terra functions.

## Value

An object of the same class as from, but potentially cropped (via `cropTo()`), projected (via `projectTo()`), masked (via `maskTo()`), and written to disk (via `writeTo()`).

## Use Cases

The table below shows what will result from passing different classes to from and to:

from	to	from will have:
Gridded	Gridded	the extent, projection, origin, resolution and masking where there are NA from the to
Gridded	Vector	the projection, origin, and mask from to, and extent will be a round number of pixels that fit within the
Vector	Vector	the projection, origin, extent and mask from to

If one or more of the \*To arguments are supplied, these will override individual components of to. If to is omitted or NULL, then only the \*To arguments that are used will be performed. In all cases, setting a \*To argument to NA will prevent that step from happening.

## projectTo

Since these functions use the gis capabilities of sf and terra, they will only be able to do things that those functions can do. One key caution, which is stated clearly in `?terra::project` is that projection of a raster (i.e., gridded) object should always be with another gridded object. If the user chooses to supply a projectTo that is a vector object for a from that is gridded, there may be unexpected failures due e.g., to extents not overlapping during the maskTo stage.

## Backwards compatibility with postProcess

`rasterToMatch` **and** `studyArea`::

If these are supplied, postProcessTo will use them instead of to. If only rasterToMatch is supplied, it will be assigned to to. If only studyArea is supplied, it will be used for cropTo and maskTo; it will only be used for projectTo if useSAcrs = TRUE. If both rasterToMatch and studyArea are supplied, studyArea will only be applied to maskTo (unless maskWithRTM = TRUE), and, optionally, to projectTo (if useSAcrs = TRUE); everything else will be from rasterToMatch.

`targetCRS`, `filename2`, `useSAcrs`, `maskWithRTM`::

targetCRS if supplied will be assigned to projectTo. filename2 will be assigned to writeTo. If useSAcrs is set, then the studyArea will be assigned to projectTo. If maskWithRTM is used, then the rasterToMath will be assigned to maskTo. All of these will override any existing values for these arguments.

See also [postProcess\(\)](#) documentation section on *Backwards compatibility with rasterToMatch and/or studyArea* for further detail.

## Cropping

If `cropTo` is not NA, `postProcessTo` does cropping twice, both the first and last steps. It does it first for speed, as cropping is a very fast algorithm. This will quickly remove a bunch of pixels that are not necessary. But, to not create bias, this first crop is padded by  $2 * \text{res}(\text{from})[1]$ , so that edge cells still have a complete set of neighbours. The second crop is at the end, after projecting and masking. After the projection step, the crop is no longer tight. Under some conditions, masking will effectively mask and crop in one step, but under some conditions, this is not true, and the mask leaves padded NAs out to the extent of the `from` (as it is after `crop`, `project`, `mask`). Thus the second crop removes all NA cells so they are tight to the mask.

## See Also

[maskTo\(\)](#), [cropTo\(\)](#), [projectTo\(\)](#), [writeTo\(\)](#), and [fixErrorsIn\(\)](#). Also the functions that call `sf::gdal_utils(...)` directly: [gdalProject\(\)](#), [gdalResample\(\)](#), [gdalMask\(\)](#)

## Examples

```
# prepare dummy data -- 3 SpatRasters, 2 SpatVectors
# need 2 SpatRaster
rf <- system.file("ex/elev.tif", package = "terra")
elev1 <- terra::rast(rf)
#'
ras2 <- terra::deecopy(elev1)
ras2[ras2 > 200 & ras2 < 300] <- NA_integer_
terra::values(elev1) <- rep(1L, terra::ncell(ras2))
#'
# a polygon vector
f <- system.file("ex/lux.shp", package = "terra")
vOrig <- terra::vect(f)
v <- vOrig[1:2, ]
#'
utm <- terra::crs("epsg:23028") # $wkt
vInUTM <- terra::project(vOrig, utm)
vAsRasInLongLat <- terra::rast(vOrig, resolution = 0.008333333)
res100 <- 100
rInUTM <- terra::rast(vInUTM, resolution = res100, vals = 1)
# crop, reproject, mask, crop a raster with a vector in a different projection
# --> gives message about not enough information
t1 <- postProcessTo(elev1, to = vInUTM)
# crop, reproject, mask a raster to a different projection, then mask
t2a <- postProcessTo(elev1, to = vAsRasInLongLat, maskTo = vInUTM)

# using gdal directly --> slightly different mask
opts <- options(reproducible.gdalwarp = TRUE)
t2b <- postProcessTo(elev1, to = vAsRasInLongLat, maskTo = vInUTM)
t3b <- postProcessTo(elev1, to = rInUTM, maskTo = vInUTM)
options(opts)
```

---

```
prepInputs
```

---

*Download and optionally post-process files*

---

## Description

## Usage

```
prepInputs(
  targetFile = NULL,
  url = NULL,
  archive = NULL,
  alsoExtract = NULL,
  destinationPath = getOption("reproducible.destinationPath", "."),
  fun = NULL,
  quick = getOption("reproducible.quick"),
  overwrite = getOption("reproducible.overwrite", FALSE),
  purge = FALSE,
  useCache = getOption("reproducible.useCache", 2),
  .tempPath,
  verbose = getOption("reproducible.verbose", 1),
  ...
)
```

## Arguments

targetFile	Character string giving the filename (without relative or absolute path) to the eventual file (raster, shapefile, csv, etc.) after downloading and extracting from a zip or tar archive. This is the file <i>before</i> it is passed to postProcess. The internal checksumming does not checksum the file after it is postProcessed (e.g., cropped/reprojected/masked). Using Cache around prepInputs will do a sufficient job in these cases. See table in <a href="#">preProcess()</a> .
url	Optional character string indicating the URL to download from. If not specified, then no download will be attempted. If not entry exists in the CHECKSUMS.txt (in destinationPath), an entry will be created or appended to. This CHECKSUMS.txt entry will be used in subsequent calls to prepInputs or preProcess, comparing the file on hand with the ad hoc CHECKSUMS.txt. See table in <a href="#">preProcess()</a> .
archive	Optional character string giving the path of an archive containing targetFile, or a vector giving a set of nested archives (e.g., c("xxx.tar", "inner.zip", "inner.rar")). If there is/are (an) inner archive(s), but they are unknown, the function will try all until it finds the targetFile. See table in <a href="#">preProcess()</a> . If it is NA, then it will <i>not</i> attempt to see it as an archive, even if it has archive-like file extension (e.g., .zip). This may be useful when an R function is expecting an archive directly.



alsoExtract	Optional character string naming files other than <code>targetFile</code> that must be extracted from the archive. If <code>NULL</code> , the default, then it will extract all files. Other options: "similar" will extract all files with the same filename without file extension as <code>targetFile</code> . <code>NA</code> will extract nothing other than <code>targetFile</code> . A character string of specific file names will cause only those to be extracted. See table in <code>preProcess()</code> .
destinationPath	Character string of a directory in which to download and save the file that comes from <code>url</code> and is also where the function will look for archive or <code>targetFile</code> . NOTE (still experimental): To prevent repeated downloads in different locations, the user can also set <code>options("reproducible.inputPaths")</code> to one or more local file paths to search for the file before attempting to download. Default for that option is <code>NULL</code> meaning do not search locally.
fun	Optional. If specified, this will attempt to load whatever file was downloaded during <code>preProcess</code> via <code>d1Fun</code> . This can be either a function (e.g., <code>sf::st_read</code> ), character string (e.g., "base::load"), <code>NA</code> (for no loading, useful if <code>d1Fun</code> already loaded the file) or if extra arguments are required in the function call, it must be a call naming <code>targetFile</code> (e.g., <code>sf::st_read(targetFile, quiet = TRUE)</code> ) as the file path to the file to load. See details and examples below.
quick	Logical. This is passed internally to <code>Checksums()</code> (the <code>quickCheck</code> argument), and to <code>Cache()</code> (the <code>quick</code> argument). This results in faster, though less robust checking of inputs. See the respective functions.
overwrite	Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.
purge	Logical or Integer. <code>0/FALSE</code> (default) keeps existing <code>CHECKSUMS.txt</code> file and <code>prepInputs</code> will write or append to it. <code>1/TRUE</code> will delete the entire <code>CHECKSUMS.txt</code> file. Other options, see details.
useCache	Passed to <code>Cache</code> in various places. Defaults to <code>getOption("reproducible.useCache", 2L)</code> in <code>prepInputs</code> , and <code>getOption("reproducible.useCache", FALSE)</code> if calling any of the inner functions manually. For <code>prepInputs</code> , this means it will use <code>Cache</code> only up to 2 nested levels, which includes <code>preProcess</code> , <code>postProcess</code> and its nested <code>*Input</code> functions (e.g., <code>cropInputs</code> , <code>projectInputs</code> , <code>maskInputs</code> ) are no longer internally cached, as <code>terra</code> processing speeds mean internal caching is more time consuming. We recommend caching the full <code>prepInputs</code> call instead (e.g. <code>prepInputs(... )  &gt; Cache()</code> ).
.tempPath	Optional temporary path for internal file intermediate steps. Will be cleared on <code>exit</code> from this function.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t
...	Additional arguments passed to <code>postProcess()</code> and <code>Cache()</code> . Since <code>...</code> is passed to <code>postProcess()</code> , these will <code>...</code> will also be passed into the inner functions, e.g., <code>cropInputs()</code> . Possibly useful other arguments include <code>d1Fun</code> which is passed to <code>preProcess</code> . See details and examples.

## Details

This function can be used to prepare R objects from remote or local data sources. The object of this function is to provide a reproducible version of a series of commonly used steps for getting, loading, and processing data. This function has two stages: Getting data (download, extracting from archives, loading into R) and post-processing (for `Spatial*` and `Raster*` objects, this is crop, reproject, mask/intersect). To trigger the first stage, provide `url` or `archive`. To trigger the second stage, provide `studyArea` or `rasterToMatch`. See examples.

## Value

This is an omnibus function that will return an R object that will have resulted from the running of `preProcess()` and `postProcess()` or `postProcessTo()`. Thus, if it is a GIS object, it may have been cropped, reprojected, "fixed", masked, and written to disk.

### Stage 1 - Getting data

See `preProcess()` for combinations of arguments.

1. Download from the web via either `googledrive::drive_download()`, `utils::download.file()`;
2. Extract from archive using `unzip()` or `untar()`;
3. Load into R using `terra::rast`, `sf::st_read`, or any other function passed in with `fun`;
4. Checksumming of all files during this process. This is put into a 'CHECKSUMS.txt' file in the `destinationPath`, appending if it is already there, overwriting the entries for same files if entries already exist.

### Stage 2 - Post processing

This will be triggered if either `rasterToMatch` or `studyArea` is supplied.

1. Fix errors. Currently only errors fixed are for `SpatialPolygons` using `buffer(..., width = 0)`;
2. Crop using `cropTo()`;
3. Project using `projectTo()`;
4. Mask using `maskTo()`;
5. write the file to disk via `writeTo()`.

NOTE: checksumming does not occur during the post-processing stage, as there are no file downloads. To achieve fast results, wrap `prepInputs` with `Cache`.

NOTE: `sf` objects are still very experimental.

#### **postProcessing of `Spat*`, `sf`, `Raster*` and `Spatial*` objects:**

The following has been DEPRECATED because there are a sufficient number of ambiguities that this has been changed in favour of `from` and the `*to` family. See `postProcessTo()`.

DEPRECATED: If `rasterToMatch` or `studyArea` are used, then this will trigger several subsequent functions, specifically the sequence, *Crop, reproject, mask*, which appears to be a common sequence while preparing spatial data from diverse sources. See `postProcess()` documentation section on *Backwards compatibility with rasterToMatch and/or studyArea arguments* to understand various combinations of `rasterToMatch` and/or `studyArea`.

**fun**

fun offers the ability to pass any custom function with which to load the file obtained by preProcess into the session. There are two cases that are dealt with: when the preProcess downloads a file (including via dlFun), fun must deal with a file; and, when preProcess creates an R object (e.g., raster::getData returns an object), fun must deal with an object.

fun can be supplied in three ways: a function, a character string (i.e., a function name as a string), or an expression. If a character string or function, it should have the package name e.g., "terra::rast" or as an actual function, e.g., base::readRDS. In these cases, it will evaluate this function call while passing targetFile as the first argument. These will only work in the simplest of cases.

When more precision is required, the full call can be written and where the filename can be referred to as targetFile if the function is loading a file. If preProcess returns an object, fun should be set to fun = NA.

If there is a custom function call, is not in a package, prepInputs may not find it. In such cases, simply pass the function as a named argument (with same name as function) to prepInputs. See examples. NOTE: passing fun = NA will skip loading object into R. Note this will essentially replicate the functionality of simply calling preProcess directly.

**purge**

In options for control of purging the CHECKSUMS.txt file are:

- 0 keep file
- 1 delete file in destinationPath, all records of downloads need to be rebuilt
- 2 delete entry with same targetFile
- 4 delete entry with same alsoExtract
- 3 delete entry with same archive
- 5 delete entry with same targetFile & alsoExtract
- 6 delete entry with same targetFile, alsoExtract & archive
- 7 delete entry that same targetFile, alsoExtract & archive & url

will only remove entries in the CHECKSUMS.txt that are associated with targetFile, alsoExtract or archive When prepInputs is called, it will write or append to a (if already exists) CHECKSUMS.txt file. If the CHECKSUMS.txt is not correct, use this argument to remove it.

**Note**

This function is still experimental: use with caution.

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**See Also**

[postProcessTo\(\)](#), [downloadFile\(\)](#), [extractFromArchive\(\)](#), [postProcess\(\)](#).

**Examples**

```

if (requireNamespace("terra", quietly = TRUE) &&
    requireNamespace("sf", quietly = TRUE)) {
  library(reproducible)
  # Make a dummy study area map -- user would supply this normally
  coords <- structure(c(-122.9, -116.1, -99.2, -106, -122.9, 59.9, 65.7, 63.6, 54.8, 59.9),
    .Dim = c(5L, 2L)
  )
  studyArea <- terra::vect(coords, "polygons")
  terra::crs(studyArea) <- "+proj=longlat +datum=WGS84 +no_defs +ellps=WGS84 +towgs84=0,0,0"
  # Make dummy "large" map that must be cropped to the study area
  outerSA <- terra::buffer(studyArea, 50000)
  terra::crs(outerSA) <- "+proj=longlat +datum=WGS84 +no_defs +ellps=WGS84 +towgs84=0,0,0"
  tf <- normPath(file.path(tempdir2("prepInputsEx"), "prepInputs2.shp"))
  terra::writeVector(outerSA, tf, overwrite = TRUE)

  # run prepInputs -- load file, postProcess it to the studyArea

  studyArea2 <- prepInputs(
    targetFile = tf, to = studyArea,
    fun = "terra::vect",
    destinationPath = tempdir2()
  ) |>
    suppressWarnings() # not relevant warning here

  # clean up
  unlink("CHECKSUMS.txt")

#####
# Remote file using `url`
#####
if (internetExists()) {
  data.table::setDTthreads(2)
  origDir <- getwd()
  # download a zip file from internet, unzip all files, load as shapefile, Cache the call
  # First time: don't know all files - prepInputs will guess, if download file is an archive,
  # then extract all files, then if there is a .shp, it will load with sf::st_read
  dPath <- file.path(tempdir(), "ecozones")
  shpUrl <- "http://sis.agr.gc.ca/cansis/nsdb/ecostrat/zone/ecozone_shp.zip"

  # Wrapped in a try because this particular url can be flaky
  shpEcozone <- try(prepInputs(
    destinationPath = dPath,
    url = shpUrl
  ))
  if (!is(shpEcozone, "try-error")) {
    # Robust to partial file deletions:
    unlink(dir(dPath, full.names = TRUE)[1:3])
    shpEcozone <- prepInputs(
      destinationPath = dPath,
      url = shpUrl
    )
  }
}

```

```

unlink(dPath, recursive = TRUE)

# Once this is done, can be more precise in operational code:
# specify targetFile, alsoExtract, and fun, wrap with Cache
ecozoneFilename <- file.path(dPath, "ecozones.shp")
ecozoneFiles <- c(
  "ecozones.dbf", "ecozones.prj",
  "ecozones.sbn", "ecozones.sbx", "ecozones.shp", "ecozones.shx"
)
shpEcozone <- prepInputs(
  targetFile = ecozoneFilename,
  url = shpUrl,
  fun = "terra::vect",
  alsoExtract = ecozoneFiles,
  destinationPath = dPath
)
unlink(dPath, recursive = TRUE)

# Add a study area to Crop and Mask to
# Create a "study area"
coords <- structure(c(-122.98, -116.1, -99.2, -106, -122.98, 59.9, 65.73, 63.58, 54.79, 59.9),
  .Dim = c(5L, 2L)
)
studyArea <- terra::vect(coords, "polygons")
terra::crs(studyArea) <- "+proj=longlat +datum=WGS84 +no_defs +ellps=WGS84 +towgs84=0,0,0"

# specify targetFile, alsoExtract, and fun, wrap with Cache
ecozoneFilename <- file.path(dPath, "ecozones.shp")
# Note, you don't need to "alsoExtract" the archive... if the archive is not there, but the
# targetFile is there, it will not redownload the archive.
ecozoneFiles <- c(
  "ecozones.dbf", "ecozones.prj",
  "ecozones.sbn", "ecozones.sbx", "ecozones.shp", "ecozones.shx"
)
shpEcozoneSm <- Cache(prepInputs,
  url = shpUrl,
  targetFile = reproducible::asPath(ecozoneFilename),
  alsoExtract = reproducible::asPath(ecozoneFiles),
  studyArea = studyArea,
  fun = "terra::vect",
  destinationPath = dPath,
  writeTo = "EcozoneFile.shp"
) # passed to determineFilename

terra::plot(shpEcozone[, 1])
terra::plot(shpEcozoneSm[, 1], add = TRUE, col = "red")
unlink(dPath)
}
}
}

## Using quoted dlFun and fun -- this is not intended to be run but used as a template

```

```
## prepInputs(..., fun = customFun(x = targetFile), customFun = customFun)
## # or more complex
## test5 <- prepInputs(
##   targetFile = targetFileLuxRDS,
##   dlFun =
##     getDataFn(name = "GADM", country = "LUX", level = 0) # preProcess keeps file from this!
##   ,
##   fun = {
##     out <- readRDS(targetFile)
##     sf::st_as_sf(out)}
## )
```

---

```
preProcessParams      Download, checksum, extract files
```

---

## Description

This does downloading (via `downloadFile`), checksumming (`Checksums`), and extracting from archives (`extractFromArchive`), plus cleaning up of input arguments (e.g., paths, function names). This is the first stage of three used in `prepInputs`.

## Usage

```
preProcessParams(n = NULL)

preProcess(
  targetFile = NULL,
  url = NULL,
  archive = NULL,
  alsoExtract = NULL,
  destinationPath = getOption("reproducible.destinationPath", "."),
  fun = NULL,
  dlFun = NULL,
  quick = getOption("reproducible.quick"),
  overwrite = getOption("reproducible.overwrite", FALSE),
  purge = FALSE,
  verbose = getOption("reproducible.verbose", 1),
  .tempPath,
  ...
)
```

## Arguments

`n` Number of non-null arguments passed to `preProcess`. E.g., passing `n = 1` returns combinations with only a single non-NULL parameter. If NULL (default), all parameter combinations are returned.

targetFile	Character string giving the filename (without relative or absolute path) to the eventual file (raster, shapefile, csv, etc.) after downloading and extracting from a zip or tar archive. This is the file <i>before</i> it is passed to postProcess. The internal checksumming does not checksum the file after it is postProcessed (e.g., cropped/reprojected/masked). Using Cache around prepInputs will do a sufficient job in these cases. See table in <a href="#">preProcess()</a> .
url	Optional character string indicating the URL to download from. If not specified, then no download will be attempted. If not entry exists in the CHECKSUMS.txt (in destinationPath), an entry will be created or appended to. This CHECKSUMS.txt entry will be used in subsequent calls to prepInputs or preProcess, comparing the file on hand with the ad hoc CHECKSUMS.txt. See table in <a href="#">preProcess()</a> .
archive	Optional character string giving the path of an archive containing targetFile, or a vector giving a set of nested archives (e.g., c("xxx.tar", "inner.zip", "inner.rar")). If there is/are (an) inner archive(s), but they are unknown, the function will try all until it finds the targetFile. See table in <a href="#">preProcess()</a> . If it is NA, then it will <i>not</i> attempt to see it as an archive, even if it has archive-like file extension (e.g., .zip). This may be useful when an R function is expecting an archive directly.
alsoExtract	Optional character string naming files other than targetFile that must be extracted from the archive. If NULL, the default, then it will extract all files. Other options: "similar" will extract all files with the same filename without file extension as targetFile. NA will extract nothing other than targetFile. A character string of specific file names will cause only those to be extracted. See table in <a href="#">preProcess()</a> .
destinationPath	Character string of a directory in which to download and save the file that comes from url and is also where the function will look for archive or targetFile. NOTE (still experimental): To prevent repeated downloads in different locations, the user can also set options("reproducible.inputPaths") to one or more local file paths to search for the file before attempting to download. Default for that option is NULL meaning do not search locally.
fun	Optional. If specified, this will attempt to load whatever file was downloaded during preProcess via dlFun. This can be either a function (e.g., sf::st_read), character string (e.g., "base::load"), NA (for no loading, useful if dlFun already loaded the file) or if extra arguments are required in the function call, it must be a call naming targetFile (e.g., sf::st_read(targetFile, quiet = TRUE)) as the file path to the file to load. See details and examples below.
dlFun	Optional "download function" name, such as "raster::getData", which does custom downloading, in addition to loading into R. Still experimental.
quick	Logical. This is passed internally to <a href="#">Checksums()</a> (the quickCheck argument), and to <a href="#">Cache()</a> (the quick argument). This results in faster, though less robust checking of inputs. See the respective functions.
overwrite	Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.
purge	Logical or Integer. 0/FALSE (default) keeps existing CHECKSUMS.txt file and prepInputs will write or append to it. 1/TRUE will deleted the entire CHECKSUMS.txt file. Other options, see details.

verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t
.tempPath	Optional temporary path for internal file intermediate steps. Will be cleared on.exit from this function.
...	Additional arguments passed to <code>postProcess()</code> and <code>Cache()</code> . Since ... is passed to <code>postProcess()</code> , these will ... will also be passed into the inner functions, e.g., <code>cropInputs()</code> . Possibly useful other arguments include <code>dIFun</code> which is passed to <code>preProcess</code> . See details and examples.

**Value**

A list with 5 elements: `checkSums` (the result of a Checksums after downloading), `dots` (cleaned up ..., including deprecated argument checks), `fun` (the function to be used to load the preProcessed object from disk), and `targetFilePath` (the fully qualified path to the `targetFile`).

**Combinations of targetFile, url, archive, alsoExtract**

Use `preProcessParams()` for a table describing various parameter combinations and their outcomes.

\* If the `url` is a file on Google Drive, checksumming will work even without a `targetFile` specified because there is an initial attempt to get the remove file information (e.g., file name). With that, the connection between the `url` and the filename used in the 'CHECKSUMS.txt' file can be made.

**Author(s)**

Eliot McIntire

---

rasterRead	<i>A helper to getOption("reproducible.rasterRead")</i>
------------	---

---

**Description**

A helper to `getOption("reproducible.rasterRead")`

**Usage**

```
rasterRead(...)
```

**Arguments**

... Passed to the function parsed and evaluated from `getOption("reproducible.rasterRead")`

**Value**

A function, that will be the evaluated, parsed character string, e.g., `eval(parse(text = "terra::rast"))`



---

remapFileNames	<i>Remap file names</i>
----------------	-------------------------

---

**Description**

Update file path metadata for file-backed objects (e.g., SpatRasters). Useful when moving saved objects between projects or machines.

**Usage**

```
remapFileNames(obj, tags, cachePath, ...)
```

**Arguments**

obj	(optional) object whose file path metadata will be remapped
tags	cache tags data.table object
cachePath	character string specifying the path to the cache directory or NULL
...	Additional path arguments, passed to <code>absoluteBase()</code> and <code>modifyListPaths()</code>

---

reproducibleOptions	<i>reproducible options</i>
---------------------	-----------------------------

---

**Description**

These provide top-level, powerful settings for a comprehensive reproducible workflow. To see defaults, run `reproducibleOptions()`. See Details below.

**Usage**

```
reproducibleOptions()
```

**Details**

Below are options that can be set with `options("reproducible.xxx" = newValue)`, where `xxx` is one of the values below, and `newValue` is a new value to give the option. Sometimes these options can be placed in the user's `.Rprofile` file so they persist between sessions.

The following options are likely of interest to most users:

`ask` Default: TRUE. Used in `clearCache()` and `keepCache()`.

`cachePath` Default: `.reproducibleTempCacheDir`. Used in `Cache()` and many others. The default path for repositories if not passed as an argument.

`cacheSaveFormat` Default: "rds". What save format to use; currently, "qs" or "rds".

- `cacheSpeed` Default "slow". One of "slow" or "fast" (1 or 2). "slow" uses `digest::digest` internally, which is transferable across operating systems, but much slower than `digest::digest(algo = "spooky")`. So, if all caching is happening on a single machine, "fast" would be a good setting.
- `conn` Default: NULL. Sets a specific connection to a database, e.g., `dbConnect(drv = RSQLite::SQLite())` or `dbConnect(drv = RPostgres::Postgres())`. For remote database servers, setting one connection may be far faster than using `drv` which must make a new connection every time.
- `destinationPath` Default: NULL. Used in `prepInputs()` and `preProcess()`. Can be set globally here.
- `drv` Default: `RSQLite::SQLite()`. Sets the default driver for the backend database system. Only tested with `RSQLite::SQLite()` and `RPostgres::Postgres()`.
- `futurePlan` Default: FALSE. On Linux OSes, `Cache` and `cloudCache` have some functionality that uses the `future` package. Default is to not use these, as they are experimental. They may, however, be very effective in speeding up some things, specifically, uploading cached elements via `googledrive` in `cloudCache`.
- `gdalwarp` Default: FALSE. Experimental. During `postProcessTo` the standard approach is to use `terra` functions directly, with several strategic uses of `sf`. However, in the special case when `from` is a `SpatRaster` or `Raster`, `maskTo` is a `SpatVector` or `SFC_POLYGON` and `projectTo` is a `SpatRaster` or `Raster`, setting this option to TRUE will use `sf::gdal_utils("warp")`. In many test cases, this is much faster than the `terra` sequence. The resulting `SpatRaster` is not identical, but it is very similar.
- `gdalwarpThreads` Default: 2. This will set `-wo NUM_THREADS=` to this number. Default is now 2, meaning `gdalwarp` will use 2 threads with `gdalProject`. To turn off threading, set to 0, 1 or NA.
- `inputPaths` Default: NULL. Used in `prepInputs()` and `preProcess()`. If set to a path, this will cause these functions to save their downloaded and preprocessed file to this location, with a `hardlink` (via `file.link`) to the file created in the `destinationPath`. This can be used so that individual projects that use common data sets can maintain modularity (by placing downloaded objects in their `destinationPath`, but also minimize re-downloading the same (perhaps large) file over and over for each project. Because the files are `hardlinks`, there is no extra space taken up by the apparently duplicated files.
- `inputPathsRecursive` Default: FALSE. Used in `prepInputs()` and `preProcess()`. Should the `reproducible.inputPaths` be searched recursively for existence of a file?
- `memoisePersist` Default: FALSE. Used in `Cache()`. Should the memoised copy of the `Cache` objects persist even if `reproducible` reloads e.g., via `devtools::load_all`? This is mostly useful for developers of `reproducible`. If TRUE, a object named `paste0(".reproducibleMemoise_", cachePath)` will be placed in the `.GlobalEnv`, i.e., one for each `cachePath`.
- `nThreads` Default: 1. The number of threads to use for reading/writing cache files.
- `objSize` Default: TRUE. Logical. If TRUE, then object sizes will be included in the cache database. Simplying calculating object size of large objects can be time consuming, so setting this to FALSE will make caching up to 10% faster, depending on the objects.
- `overwrite` Default: FALSE. Used in `prepInputs()`, `preProcess()`, `downloadFile()`, and `postProcess()`.
- `quick` Default: FALSE. Used in `Cache()`. This will cause `Cache` to use `file.size(file)` instead of the `digest::digest(file)`. Less robust to changes, but faster. *NOTE: this will only affect objects on disk.*

- `rasterRead` Used during `prepInputs` when reading `.tif`, `.grd`, and `.asc` files. Default: `terra::rast`. Can be `raster::raster` for backwards compatibility. Can be set using environment variable `R_REPRODUCIBLE_RASTER_READ`.
- `shapefileRead` Default `NULL`. Used during `prepInputs` when reading a `.shp` file. If `NULL`, it will use `sf::st_read` if `sf` package is available; otherwise, it will use `raster::shapefile`
- `showSimilar` Default `FALSE`. Passed to `Cache`.
- `testCharacterAsFile` Default `FALSE`. The behaviour of `.robustDigest` on character vectors prior to `reproducible == 2.1.2` was that the function would test for whether they were filenames by using `file.exists`. If it was a filename, then it would digest the file content. In cases of a character vector or a `data.frame` of "filenames", this could cause long hanging of the R system as it tries to digest the file contents of potentially many files. This behaviour is not transparent to a user. Now the default is to **not** digest the file content of a character vector even if they are filenames. To force file content digesting, then convert to either `asPath` or `fs::as_fs_path`. Or set this option to `TRUE` and the previous behaviour will return, where it tries to guess whether a character vector is filenames or not, and if it is, then digest the file content.
- `timeout` Default `1200`. Used in `preProcess` when downloading occurs. If a user has `R.utils` package installed, `R.utils::withTimeout( , timeout = getOption("reproducible.timeout"))` will be wrapped around the download so that it will timeout (and error) after this many seconds.
- `useCache` Default: `TRUE`. Used in `Cache()`. If `FALSE`, then the entire `Cache` machinery is skipped and the functions are run as if there was no `Cache` occurring. Can also take 2 other values: `'overwrite'` and `'devMode'`. `'overwrite'` will cause no recovery of objects from the cache repository, only new ones will be created. If the hash is identical to a previous one, then this will overwrite the previous one. `'devMode'` will function as normally `Cache` except it will use the `userTags` to determine if a previous function has been run. If the `userTags` are identical, but the digest value is different, the old value will be deleted from the cache repository and this new value will be added. This addresses a common situation during the development stage: functions are changing frequently, so any entry in the cache repository will be stale following changes to functions, i.e., they will likely never be relevant again. This will therefore keep the cache repository clean of stale objects. If there is ambiguity in the `userTags`, i.e., they do not uniquely identify a single entry in the `cachePath`, then this option will default back to the non-dev-mode behaviour to avoid deleting objects. This, therefore, is most useful if the user is using unique values for `userTags`.
- `useCloud` Default `FALSE`. Passed to `Cache`.
- `useDBI` Default: `TRUE` if **DBI** is available. Default value can be overridden by setting environment variable `R_REPRODUCIBLE_USE_DBI`. As of version 0.3, the backend is now **DBI** instead of **archivist**.
- `useGdown` Default: `FALSE`. If a user provides a Google Drive url to `preProcess/prepInputs`, `reproducible` will use the `googledrive` package. This works reliably in most cases. However, for large files on unstable internet connections, it will stall and stop the download with no error. If a user is finding this behaviour, they can install the `gdown` package, making sure it is available on the `PATH`. This call to `gdown` will only work for files that do not need authentication. If authentication is needed, `d1Google` will fall back to `googledrive::drive_download`, even if this option is `TRUE`, with a message. .

- `useMemoise` Default: FALSE. Used in `Cache()`. If TRUE, recovery of cached elements from the `cachePath` will use `memoise::memoise`. This means that the 2nd time running a function will be much faster than the first in a session (which either will create a new cache entry to disk or read a cached entry from disk). *NOTE: memoised values are removed when the R session is restarted. This option will use more RAM* and so may need to be turned off if RAM is limiting. `clearCache` of any sort will cause all memoising to be 'forgotten' (`memoise::forget`).
- `useNewDigestAlgorithm` Default: 1. Option 1 is the version that has existed for sometime. There is now an option 2 which is substantially faster. It will, however, create Caches that are not compatible with previous ones. Options 1 and 2 are not compatible with the earlier 0. 1 and 2 will make Cache less sensitive to minor but irrelevant changes (like changing the order of arguments) and will work successfully across operating systems (especially relevant for the new `cloudCache` function).
- `useTerra` Default: FALSE. The GIS operations in `postProcess`, by default use primarily the Raster package. The newer terra package does similar operations, but usually faster. A user can now set this option to TRUE and `prepInputs` and several components of `postProcess` will use terra internally.
- `verbose` Default: FALSE. If set to TRUE then every Cache call will show a summary of the objects being cached, their `object.size` and the time it took to digest them and also the time it took to run the call and save the call to the cache repository or load the cached copy from the repository. This may help diagnosing some problems that may occur.

### Value

This function returns a list of all the options that the reproducible package sets and uses. See below for details of each.

### Advanced

The following options are likely not needed by a user.

- `cloudChecksumsFilename` Default: `file.path(dirname(.reproducibleTempCacheDir()), "checksums.rds")`. Used as an experimental argument in `Cache()`
- `length` Default: `Inf`. Used in `Cache()`, specifically to the internal calls to `CacheDigest()`. This is passed to `digest::digest`. Mostly this would be changed from default `Inf` if the digesting is taking too long. Use this with caution, as some objects will have *many* NA values in their first *many* elements
- `useragent` Default: `"https://github.com/PredictiveEcology/reproducible"`. User agent for downloads using this package.

---

retry

*A wrapper around try that retries on failure*

---

### Description

This is useful for functions that are "flaky", such as `curl`, which may fail for unknown reasons that do not persist.

**Usage**

```

retry(
  expr,
  envir = parent.frame(),
  retries = 5,
  exponentialDecayBase = 1.3,
  silent = TRUE,
  exprBetween = NULL,
  messageFn = message
)

```

**Arguments**

<code>expr</code>	An expression to run, i.e., <code>rnorm(1)</code> , similar to what is passed to <code>try</code>
<code>envir</code>	The environment in which to evaluate the quoted expression, default to <code>parent.frame(1)</code>
<code>retries</code>	Numeric. The maximum number of retries.
<code>exponentialDecayBase</code>	Numeric > 1.0. The delay between successive retries will be <code>runif(1, min = 0, max = exponentialDecayBase ^ i - 1)</code> where <code>i</code> is the retry number (i.e., follows <code>seq_len(retries)</code> )
<code>silent</code>	Logical indicating whether to try silently.
<code>exprBetween</code>	Another expression that should be run after a failed attempt of the <code>expr</code> . This should return a named list, where the names indicate the object names to update in the main <code>expr</code> , and the return value is the new value. (previous versions allowed a non-list return, but where the final line had to be an assignment operator, specifying what object (that is used in <code>expr</code> ) will be updated prior to running the <code>expr</code> again. For backwards compatibility, this still works).
<code>messageFn</code>	A function for messaging to console. Defaults to <code>message</code>

**Details**

Based on <https://github.com/jennybc/googlesheets/issues/219#issuecomment-195218525>.

**Value**

As with `try`, so the successfully returned `return()` from the `expr` or a `try-error`.

---

<code>saveToCache</code>	<i>Save an object to Cache</i>
--------------------------	--------------------------------

---

**Description**

This is not expected to be used by a user as it requires that the `cacheId` be calculated in exactly the same as it calculated inside `Cache` (which requires `match.call` to match arguments with their names, among other things).

**Usage**

```
saveToCache(
  cachePath = getOption("reproducible.cachePath"),
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  obj,
  userTags,
  cacheId,
  linkToCacheId = NULL,
  verbose = getOption("reproducible.verbose")
)
```

**Arguments**

cachePath	A repository used for storing cached objects. This is optional if Cache is used inside a SpaDES module.
drv	if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite
conn	an optional DBIConnection object, as returned by dbConnect().
obj	The R object to save to the cache
userTags	A character vector with descriptions of the Cache function call. These will be added to the Cache so that this entry in the Cache can be found using userTags e.g., via <a href="#">showCache()</a> .
cacheId	The hash string representing the result of <code>.robustDigest</code>
linkToCacheId	Optional. If a cacheId is provided here, then a file.link will be made to the file with that cacheId name in the cache repo. This is used when identical outputs exist in the cache. This will save disk space.
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce t

**Value**

This is used for its side effects, namely, it will add the object to the cache and cache database.

---

searchFull	<i>Search up the full scope for functions</i>
------------	---

---

**Description**

This is like `base::search` but when used inside a function, it will show the full scope (see figure in the section *Binding environments* on <http://adv-r.had.co.nz/Environments.html>). This full search path will be potentially much longer than just `search()` (which always starts at `.GlobalEnv`). `searchFullEx` shows an example function that is inside this package whose only function is to show the Scope of a package function.

**Usage**

```
searchFull(env = parent.frame(), simplify = TRUE)
```

```
searchFullEx()
```

**Arguments**

`env` The environment to start searching at. Default is calling environment, i.e., `parent.frame()`

`simplify` Logical. Should the output be simplified to character, if possible (usually it is not possible because environments don't always coerce correctly)

**Details**

`searchFullEx` can be used to show an example of the use of `searchFull`.

**Value**

A list of environments that is the actual search path, unlike `search()` which only prints from `.GlobalEnv` up to base through user attached packages.

**See Also**

[base::search\(\)](#)

**Examples**

```
seeScope <- function() {
  searchFull()
}
seeScope()
searchFull()
searchFullEx()
```

---

`set.randomseed`      *Set seed with a random value using Sys.time()*

---

**Description**

This will set a random seed.

**Usage**

```
set.randomseed(set.seed = TRUE)
```

**Arguments**

`set.seed` Logical. If TRUE, the default, then the function will call `set.seed` internally with the new random seed.

**Details**

This function uses 6 decimal places of `Sys.time()`, i.e., microseconds. Due to integer limits, it also truncates at 1000 seconds, so there is a possibility that this will be non-unique after 1000 seconds (at the microsecond level). In tests, this showed no duplicates after  $1e7$  draws in a loop, as expected.

**Value**

This will return the new seed invisibly. However, this is also called for its side effects, which is a new seed set using `set.seed`

**Note**

This function does not appear to be as reliable on `R <= 4.1.3`

---

 showCache

*Examining and modifying the cache*


---

**Description**

These are convenience wrappers around DBI package functions. They allow the user a bit of control over what is being cached.

**Usage**

```
clearCache(
  x,
  userTags = character(),
  after = NULL,
  before = NULL,
  fun = NULL,
  cacheId = NULL,
  ask = getOption("reproducible.ask"),
  useCloud = FALSE,
  cloudFolderID = getOption("reproducible.cloudFolderID", NULL),
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  ...
)
```

```
## S4 method for signature 'ANY'
```

```
clearCache(
  x,
  userTags = character(),
  after = NULL,
  before = NULL,
  fun = NULL,
```



```
    cacheId = NULL,
    ask = getOption("reproducible.ask"),
    useCloud = FALSE,
    cloudFolderID = getOption("reproducible.cloudFolderID", NULL),
    drv = getDrv(getOption("reproducible.drv", NULL)),
    conn = getOption("reproducible.conn", NULL),
    verbose = getOption("reproducible.verbose"),
    ...
)

cc(secs, ..., verbose = getOption("reproducible.verbose"))

showCache(
  x,
  userTags = character(),
  after = NULL,
  before = NULL,
  fun = NULL,
  cacheId = NULL,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  ...
)

## S4 method for signature 'ANY'
showCache(
  x,
  userTags = character(),
  after = NULL,
  before = NULL,
  fun = NULL,
  cacheId = NULL,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  ...
)

keepCache(
  x,
  userTags = character(),
  after = NULL,
  before = NULL,
  ask = getOption("reproducible.ask"),
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
```

```

    ...
  )

## S4 method for signature 'ANY'
keepCache(
  x,
  userTags = character(),
  after = NULL,
  before = NULL,
  ask = getOption("reproducible.ask"),
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  verbose = getOption("reproducible.verbose"),
  ...
)

```

### Arguments

x	A simList or a directory containing a valid Cache repository. Note: For compatibility with Cache argument, cachePath can also be used instead of x, though x will take precedence.
userTags	Character vector. If used, this will be used in place of the after and before. Specifying one or more userTag here will clear all objects that match those tags. Matching is via regular expression, meaning partial matches will work unless strict beginning (^) and end (\$) of string characters are used. Matching will be against any of the 3 columns returned by showCache(), i.e., artifact, tagValue or tagName. Also, if length(userTags) > 1, then matching is by and. For or matching, use   in a single character string. See examples.
after	A time (POSIX, character understandable by data.table). Objects cached after this time will be shown or deleted.
before	A time (POSIX, character understandable by data.table). Objects cached before this time will be shown or deleted.
fun	An optional character vector describing the function name to extract. Only functions with this/these functions will be returned.
cacheId	An optional character vector describing the cacheIds to extract. Only entries with this/these cacheIds will be returned. If useDBI(FALSE), this will also be dramatically faster than using userTags, for a large cache.
ask	Logical. If FALSE, then it will not ask to confirm deletions using clearCache or keepCache. Default is TRUE
useCloud	Logical. If TRUE, then every object that is deleted locally will also be deleted in the cloudFolderID, if it is non-NULL
cloudFolderID	A googledrive dribble of a folder, e.g., using drive_mkdir(). If left as NULL, the function will create a cloud folder with name from last two folder levels of the cachePath path, : paste0(basename(dirname(cachePath)), "_", basename(cachePath)). This cloudFolderID will be added to options("reproducible.cloudFolderID") but this will not persist across sessions. If this is a character string, it will treat this as a folder name to create or use on GoogleDrive.

drv	if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite
conn	an optional DBIConnection object, as returned by dbConnect().
verbose	Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce t
...	Other arguments. Can be in the form of tagKey = tagValue, such as, class = "numeric" to find all entries that are numerics in the cache. Note: the special cases of cacheId and fun have their own named arguments in these functions. Also can be regexp = xx, where xx is TRUE if the user is passing a regular expression. Otherwise, userTags will need to be exact matches. Default is missing, which is the same as TRUE. If there are errors due to regular expression problem, try FALSE. For cc, it is passed to clearCache, e.g., ask, userTags. For showCache, it can also be sorted = FALSE to return the object unsorted.
secs	Currently 3 options: the number of seconds to pass to clearCache(after = secs), a POSIXct time e.g., from Sys.time(), or missing. If missing, the default, then it will delete the most recent entry in the Cache.

## Details

If neither after or before are provided, nor userTags, then all objects will be removed. If both after and before are specified, then all objects between after and before will be deleted. If userTags is used, this will override after or before.

cc(secs) is just a shortcut for clearCache(repo = currentRepo, after = secs), i.e., to remove any cache entries touched in the last secs seconds. Since, secs can be missing, this is also be a shorthand for "remove most recent entry from the cache".

clearCache remove items from the cache based on their userTag or times values.

keepCache remove all cached items *except* those based on certain userTags or times values.

showCache display the contents of the cache.

By default the return of showCache is sorted by cacheId. For convenience, a user can optionally have it unsorted (passing sorted = FALSE), which may be noticeably faster when the cache is large (> 1e4 entries).

## Value

Will clear all objects (or those that match userTags, or those between after or before) from the repository located in cachePath. Invisibly returns a data.table of the removed items.

## Note

If the cache is larger than 10MB, and clearCache is used, there will be a message and a pause, if interactive, to prevent accidentally deleting of a large cache repository.

**See Also**

[mergeCache\(\)](#). Many more examples in [Cache\(\)](#).

**Examples**

```
data.table::setDTthreads(2)

tmpDir <- file.path(tempdir(), "reproducible_examples", "Cache")
try(clearCache(tmpDir, ask = FALSE), silent = TRUE) # just to make sure it is clear

# Basic use
ranNumsA <- Cache(rnorm, 10, 16, cachePath = tmpDir)

# All same
ranNumsB <- Cache(rnorm, 10, 16, cachePath = tmpDir) # recovers cached copy
ranNumsD <- Cache(quote(rnorm(n = 10, 16)), cachePath = tmpDir) # recovers cached copy

# Any minor change makes it different
ranNumsE <- Cache(rnorm, 10, 6, cachePath = tmpDir) # different

## Example 1: basic cache use with tags
ranNumsA <- Cache(rnorm, 4, cachePath = tmpDir, userTags = "objectName:a")
ranNumsB <- Cache(runif, 4, cachePath = tmpDir, userTags = "objectName:b")
ranNumsC <- Cache(runif, 40, cachePath = tmpDir, userTags = "objectName:b")

showCache(tmpDir, userTags = c("objectName"))
showCache(tmpDir, userTags = c("^a$")) # regular expression ... "a" exactly

# Fine control of cache elements -- pick out only the large runif object, and remove it
cache1 <- showCache(tmpDir, userTags = c("runif")) # show only cached objects made during runif
toRemove <- cache1[tagKey == "object.size"][as.numeric(tagValue) > 700]$cacheId
clearCache(tmpDir, userTags = toRemove, ask = FALSE)
cacheAfter <- showCache(tmpDir, userTags = c("runif")) # Only the small one is left

data.table::setDTthreads(2)
tmpDir <- file.path(tempdir(), "reproducible_examples", "Cache")
try(clearCache(tmpDir, ask = FALSE), silent = TRUE) # just to make sure it is clear

Cache(rnorm, 1, cachePath = tmpDir)
thisTime <- Sys.time()
Cache(rnorm, 2, cachePath = tmpDir)
Cache(rnorm, 3, cachePath = tmpDir)
Cache(rnorm, 4, cachePath = tmpDir)
showCache(x = tmpDir) # shows all 4 entries
cc(ask = FALSE, x = tmpDir)
showCache(x = tmpDir) # most recent is gone
cc(thisTime, ask = FALSE, x = tmpDir)
showCache(x = tmpDir) # all those after thisTime gone, i.e., only 1 left
cc(ask = FALSE, x = tmpDir) # Cache is
cc(ask = FALSE, x = tmpDir) # Cache is already empty
```

---

studyAreaName	<i>Get a unique name for a given study area</i>
---------------	---

---

**Description**

Digest a spatial object to get a unique character string (hash) of the study area. Use `.suffix()` to append the hash to a filename, e.g., when using `filename2` in `prepInputs`.

**Usage**

```
studyAreaName(studyArea, ...)

## S4 method for signature 'character'
studyAreaName(studyArea, ...)

## S4 method for signature 'ANY'
studyAreaName(studyArea, ...)
```

**Arguments**

<code>studyArea</code>	Spatial object.
<code>...</code>	Other arguments (not currently used)

**Value**

A character string using the `.robustDigest` of the `studyArea`. This is only intended for use with spatial objects.

**Examples**

```
studyAreaName("Ontario")
```

---

tempdir2	<i>Make a temporary (sub-)directory</i>
----------	---

---

**Description**

Create a temporary subdirectory in `getOption("reproducible.tempPath")`.

**Usage**

```
tempdir2(
  sub = "",
  tempdir = getOption("reproducible.tempPath", .reproducibleTempPath()),
  create = TRUE
)
```

**Arguments**

sub	Character string, length 1. Can be a result of <code>file.path("smth", "smth2")</code> for nested temporary subdirectories. If the zero length character, then a random sub-directory will be created.
tempdir	Optional character string where the temporary directory should be placed. Defaults to <code>getOption("reproducible.tempPath")</code> .
create	Logical. Should the directory be created. Default TRUE.

**Value**

A character string of a path (that will be created if `create = TRUE`) in a sub-directory of the `tempdir()`.

**See Also**

[tempfile2](#)

---

tempfile2	<i>Make a temporary file in a temporary (sub-)directory</i>
-----------	---

---

**Description**

Make a temporary file in a temporary (sub-)directory

**Usage**

```
tempfile2(
  sub = "",
  tempdir = getOption("reproducible.tempPath", .reproducibleTempPath()),
  ...
)
```

**Arguments**

sub	Character string, length 1. Can be a result of <code>file.path("smth", "smth2")</code> for nested temporary subdirectories. If the zero length character, then a random sub-directory will be created.
tempdir	Optional character string where the temporary directory should be placed. Defaults to <code>getOption("reproducible.tempPath")</code> .
...	passed to <code>tempfile</code> , e.g., <code>fileext</code>

**Value**

A character string of a path to a file in a sub-directory of the `tempdir()`. This file will likely not exist yet.

**See Also**

[tempdir2](#)

---

unrarPath	<i>The known path for unrar or 7z</i>
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---

**Description**

The known path for unrar or 7z

**Usage**

```
.unrarPath
```

**Format**

An object of class NULL of length 0.

---

writeFuture	<i>Write to cache repository, using future::future</i>
-------------	--

---

**Description**

This will be used internally if options("reproducible.futurePlan" = TRUE). This is still experimental.

**Usage**

```
writeFuture(
  written,
  outputToSave,
  cachePath,
  userTags,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  cacheId,
  linkToCacheId = NULL
)
```

**Arguments**

written	Integer. If zero or positive then it needs to be written still. Should be 0 to start.
outputToSave	The R object to save to repository
cachePath	The file path of the repository
userTags	Character string of tags to attach to this outputToSave in the CacheRepo
drv	if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite

conn	an optional DBIConnection object, as returned by dbConnect().
cacheId	Character string. If passed, this will override the calculated hash of the inputs, and return the result from this cacheId in the cachePath. Setting this is equivalent to manually saving the output of this function, i.e., the object will be on disk, and will be recovered in subsequent This may help in some particularly finicky situations where Cache is not correctly detecting unchanged inputs. This will guarantee the object will be identical each time; this may be useful in operational code.
linkToCacheId	Optional. If a cacheId is provided here, then a file.link will be made to the file with that cacheId name in the cache repo. This is used when identical outputs exist in the cache. This will save disk space.

**Value**

Run for its side effect. This will add the objectToSave to the cache located at cachePath, using cacheId as its id, while updating the database entry. It will do this using the future package, so it is written in a future.



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