

# Package ‘reproducible’

January 8, 2026

**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 (>= 4.3)

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**Suggests** archive, covr, curl (>= 7.0.0), DBI, future, geodata, glue, googledrive, httr, httr2 (>= 1.2.1), knitr, parallel, parallelly, qs2, raster (>= 3.5-15), RCurl (>= 1.95-4.8), rlang, rmarkdown, RSQLite, R.utils, rvest, sf, sp (>= 1.4-2), terra (>= 1.7-20), testthat, withr

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**Language** en-CA

**License** GPL-3

**VignetteBuilder** knitr, rmarkdown

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

**ByteCompile** yes

**RoxygenNote** 7.3.3

**Collate** 'DBI.R' 'messages.R' 'GPT2.R' 'cache-helpers.R'  
 'cache-internals.R' 'robustDigest.R' 'cache.R' 'cacheGeo.R'  
 'checksums.R' 'cloud.R' 'convertPaths.R' 'copy.R' 'download.R'  
 'downloadTileAndUpload.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>

---

## Description

This hidden function appends a single tag (key-value pair) to the metadata of a cached object identified by its `cacheId`. Tags can be stored either in a database (via DBI) or in a file-based cache system.

Updates the value of an existing tag for a cached object identified by its `cacheId`. If the tag does not exist and `add = TRUE`, the tag will be added. This function supports both database-backed and file-based cache systems.

## Usage

```

.addTagsRepo(
  cacheId,
  cachePath =getOption("reproducible.cachePath"),
  tagKey = character(),
  tagValue = character(),
  cacheSaveFormat =getOption("reproducible.cacheSaveFormat"),
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn =getOption("reproducible.conn", NULL)
)

.updateTagsRepo(
  cacheId,
  cachePath =getOption("reproducible.cachePath"),
  tagKey = character(),
  tagValue = character(),
  add = TRUE,
  cacheSaveFormat =getOption("reproducible.cacheSaveFormat"),
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn =getOption("reproducible.conn", NULL)
)

```

## Arguments

cacheId	character(1) Unique identifier of the cached object. Must be of length 1.
cachePath	character(1) Path to the cache directory. Defaults to <code>getOption("reproducible.cachePath")</code> .
tagKey	character(1) The key for the tag. Must be supplied.
tagValue	character(1) The new value for the tag. Must be supplied.
cacheSaveFormat	character(1) Format used for saving cache files. Defaults to <code>getOption("reproducible.cacheSaveFormat")</code> .
drv	A DBI driver object. Defaults to <code>getDrv(getOption("reproducible.drv", NULL))</code> .
conn	A DBI connection object. If <code>NULL</code> , a new connection is created internally.
add	logical(1) If <code>TRUE</code> , adds the tag if it does not exist. Defaults to <code>TRUE</code> .

## Details

This function is primarily used internally by the `reproducible` package to maintain metadata about cached objects. It supports both database-backed and file-based caching systems.

- If `useDBI()` returns `TRUE`, the tag update is performed in the database table.
- If no rows are affected and `add = TRUE`, the tag is inserted using `.addTagsRepo()`.
- For file-based caches, the function modifies the tag in the corresponding metadata file.

### Value

NULL (invisibly). The function is called for its side effects.  
NULL (invisibly). Called for its side effects.

### See Also

[.addTagsRepo\(\)](#) for adding tags without updating.

### Examples

```
## Not run:  
a <- Cache(rnorm(1))  
.addTagsRepo(cacheId = gsub("cacheId:", "", attr(a, "tags")),  
            tagKey = "status", tagValue = "processed")  
showCache() # last entry is the above line  
  
## End(Not run)  
  
## Not run:  
a <- Cache(rnorm(1))  
# Update an existing tag  
.updateTagsRepo(cacheId = gsub("cacheId:", "", attr(a, "tags")),  
                tagKey = "status", tagValue = "second")  
  
# Add a tag if it doesn't exist  
.updateTagsRepo(cacheId = gsub("cacheId:", "", attr(a, "tags")),  
                tagKey = "status", tagValue = "new", add = TRUE)  
  
## End(Not run)
```

---

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

### Arguments

<code>obj</code>	An arbitrary R object.
<code>preDigest</code>	A list of hashes.
<code>...</code>	Dots passed from Cache
<code>fullCall</code>	The original call to Cache

**Value**

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

**Author(s)**

Eliot McIntire

---

`.file.move`

*Move a file to a new location – Defunct – use `hardLinkOrCopy`*

---

**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.

---

`.isGridded`

*Some spatial helper functions*

---

**Description**

Some spatial helper functions

**Usage**

```
.isGridded(x)

.isVector(x)

.isSF(x)

.isSpat(x)

.isSpatialAny(x)

.isCRSany(x)
```

## Arguments

- × A spatial object.

## Details

- .isGridded returns TRUE if the object is a SpatRaster or Raster
- .isVector returns TRUE if the object is SpatVector, spatial or sf
- .isSF returns TRUE if the object is sf or sfc
- .isSpat returns TRUE if the object is SpatVector or SpatRaster
- .isSpatialAny returns TRUE if the object returns TRUE for .isGridded or .isVector

## Value

Logical.

---

.isMemoised      *Evaluate whether a cacheId is memoised*

---

## Description

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

## Usage

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

## Arguments

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.
cachePath	A repository used for storing cached objects. This is optional if Cache is used inside a SpaDES module.

## Value

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

---

.objSizeWithTry	lobstr::obj_size with a try to address issue #72
-----------------	--

---

### Description

It is not clear why, but it appears that running `lobstr::obj_size` again, after a bad binding error, it will work.

### Usage

```
.objSizeWithTry(x, useTry = TRUE)
```

### Arguments

x	An object
useTry	Logical. If TRUE, the default, then it will use <code>try</code> . Can optionally avoid this if set to FALSE. The <code>try</code> takes sufficient extra compute time that it is worth avoiding it if possible.

### Value

The size of an object, using `lobstr::obj_size` or `object.size` if the first fails

---

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

**Arguments**

obj	The raster object to save to the repository.
repoDir	Character denoting an existing directory in which an artifact will be saved.
overwrite	Logical. Should the raster be saved to disk, overwriting existing file.
drv	If using a database backend, <code>drv</code> must be an object that inherits from <code>DBIDriver</code> (e.g., <code>RSQLite::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
...	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	<i>Provide standard messaging for missing package dependencies</i>
-------------------	--

---

## 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

<code>pkg</code>	Character string indicating name of package required
<code>minVersion</code>	Character string indicating minimum version of package that is needed
<code>stopOnFALSE</code>	Logical. If TRUE, this function will create an error (i.e., <code>stop</code> ) if the function returns FALSE; otherwise it simply returns FALSE
<code>messageStart</code>	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	<i>Set subattributes within a list by reference</i>
-------------------	---

---

## 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.

`.whereInStack`*Search for objects in the call stack***Description**

Normally, this is only used in special, advanced uses. The standard approach to getting an object from an environment in the call stack is to explicitly pass it into the function.

**Usage**

```
.whereInStack(obj, startingEnv = parent.frame())
```

**Arguments**

<code>obj</code>	Character string. The object name to search.
<code>startingEnv</code>	An environment to start searching in.

**Value**

The environment in which the object exists. It will return the first environment it finds, searching outwards from where the function is used.

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

```
cacheId = NULL,
...
)

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

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

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

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

## S3 method for class 'PackedSpatExtent2'
.unwrap(
  obj,
  cachePath = getOption("reproducible.cachePath"),
  cacheId = NULL,
```

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

## S3 method for class 'PackedSpatVector2'
.unwrap(
  obj,
  cachePath = getOption("reproducible.cachePath"),
  cacheId = NULL,
  drv = getDrv(getOption("reproducible.drv", NULL)),
  conn = getOption("reproducible.conn", NULL),
  ...
)

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

## S3 method for class 'PackedSpatVector'
.unwrap(
  obj,
  cachePath = getOption("reproducible.cachePath"),
  cacheId = NULL,
  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., <code>RSSQLite::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

outputObjects	Optional character vector indicating which objects to return. This is only relevant for list, environment (or similar) objects
cacheId	Used strictly for messaging. This should be the cacheId of the object being recovered. Default is NULL.
...	Arguments passed to methods; default does not use anything in ....

### 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*A version of base::basename that is NULL resistant*

---

**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*Saves a wide variety function call outputs to disk and optionally RAM, for recovery later*

---

**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,
  ...,
  dryRun =getOption("reproducible.dryRun", FALSE),
  notOlderThan = NULL,
  .objects = NULL,
  .cacheExtra = NULL,
  .functionName = NULL,
  .cacheChaining =getOption("reproducible.cacheChaining", NULL),
  outputObjects = NULL,
  algo = "xxhash64",
  cachePath = NULL,
  length =getOption("reproducible.length", Inf),
  userTags = c(),
  omitArgs = NULL,
  classOptions = list(),
  debugCache = character(),
  quick =getOption("reproducible.quick", FALSE),
  verbose =getOption("reproducible.verbose", 1),
  cacheId = NULL,
  cacheSaveFormat =getOption("reproducible.cacheSaveFormat"),
  useCache =getOption("reproducible.useCache", TRUE),
  useCloud =getOption("reproducible.useCloud", FALSE),
  cloudFolderID =getOption("reproducible.cloudFolderID", NULL),
  showSimilar =getOption("reproducible.showSimilar", FALSE),
  drv =getOption("reproducible.drv", NULL),
  conn =getOption("reproducible.conn", NULL)
)

cache2(
  FUN,
  ...,
  dryRun =getOption("reproducible.dryRun", FALSE),
  notOlderThan = NULL,
  .objects = NULL,
  .cacheExtra = NULL,
  .functionName = NULL,
  .cacheChaining =getOption("reproducible.cacheChaining", NULL),
  outputObjects = NULL,
  algo = "xxhash64",
  cachePath = NULL,
  length =getOption("reproducible.length", Inf),
  userTags = c(),
  omitArgs = NULL,
  classOptions = list(),
  debugCache = character(),
  quick =getOption("reproducible.quick", FALSE),

```

```

verbose =getOption("reproducible.verbose", 1),
cacheId = NULL,
cacheSaveFormat =getOption("reproducible.cacheSaveFormat"),
useCache =getOption("reproducible.useCache", TRUE),
useCloud =getOption("reproducible.useCloud", FALSE),
cloudFolderID =getOption("reproducible.cloudFolderID", NULL),
showSimilar =getOption("reproducible.showSimilar", FALSE),
drv =getOption("reproducible.drv", NULL),
conn =getOption("reproducible.conn", NULL)
)

CacheV2(
  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(),
  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

FUN	Either a function (e.g., <code>rnorm</code> ), a function call (e.g., <code>rnorm(1)</code> ), or an unevaluated function call (e.g., <code>using quote()</code> ).
...	Arguments passed to FUN, if FUN is not an expression.
dryRun	See <a href="#">reproducibleOptions</a> .
notOlderThan	A time. Load an object from the Cache if it was created after this.

.objects	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.
.cacheExtra	A an arbitrary R object that will be included in the CacheDigest, but otherwise not passed into the FUN. If the user supplies a named list, then Cache will report which individual elements of .cacheExtra have changed when options("reproducible.showSimilar" = TRUE). This can allow a user more control and understanding for debugging.
.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.
.cacheChaining	A logical or a the name of a function. If TRUE, then the current Cache call will evaluate the function "outside" the Cache call (via sys.function(-1)) and attach the digest of that outer function to the entry for this Cache call. This will then be used by any subsequent Cache call within the same function. If the outer function is unchanged, and there is one or more objects that had been returned by a previous Cache call, then those objects will not be digested; rather their cacheId tag will be used in place of a new digest. This <i>should</i> cause no change in Caching outcomes, and it should be faster in cases where there are several Cache calls within the same function. If FALSE (current default), then this feature is not used. If set to NULL (i.e., unset, the current default), then it will not use cache chaining, but it will attach more information to the Cache entries for each cacheId, as well as new entries for "surroundingFunction" digest, so that if a user switches to .cacheChaining = TRUE, then it will be able to begin using cache chaining without needing to rerun the calls again. Can be set by an option.
outputObjects	Optional character vector indicating which objects to return. This is only relevant for list, environment (or similar) objects
algo	The digest algorithm to use. Default xxhash64 (see <a href="#">digest::digest()</a> for others).
cachePath	A repository used for storing cached objects. This is optional if Cache is used inside a SpaDES module.
length	Numeric. If the element passed to Cache is a Path class object (from e.g., asPath(filename)) or it is a Raster with file-backing, then this will be passed to digest::digest, essentially limiting the number of bytes to digest (for speed). This will only be used if quick = FALSE. Default isgetOption("reproducible.length"), which is set to Inf.
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> .
omitArgs	Optional character string of arguments in the FUN to omit from the digest.
classOptions	Optional list. This will pass into .robustDigest for specific classes. Should be options that the .robustDigest 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, debugCache1 and debugCache2, which are the entire list(...) and that same object, but after all .robustDigest calls, at the moment that it is digested using digest, respectively. This attr(mySimOut, "debugCache2") can then be compared to a subsequent call and individual items within the object attr(mySimOut, "debugCache1") can be compared. If "quick", then it will return the same two objects directly, without evaluating the FUN(...).
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> .
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	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.
cacheSaveFormat	Character string: currently either qs or rds. Defaults togetOption("reproducible.cacheSaveFormat") qs may be faster but appears to have narrower range of conditions that work; rds is safer, and may be slower.
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 cachePath path, : paste0(basename(dirname(cachePath)), "_", basename(cachePath)). This <code>cloudFolderID</code> 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.
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, <code>drv</code> must be an object that inherits from <code>DBIDriver</code> (e.g., <code>RSSQLite::SQLite</code> ).
conn	an optional <code>DBIConnection</code> object, as returned by <code>dbConnect()</code> .
cacheRepo	Same as <code>cachePath</code> , but kept for backwards compatibility.
compareRasterFileLength	Being deprecated; use <code>length</code> .
makeCopy	Now deprecated. Ignored if used.

## 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$id`, which can be retrieved by `getOption("reproducible.cloudFolderID")$id` after at least one Cache call has been made. 2) The other user must then set their `cacheFolderID` in a Cache\(..., `reproducible.cloudFolderID` 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)

Eliot McIntire

### 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 <- 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 { }, sample is not run
}

#####
# 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	<i>The exact digest function that Cache uses</i>
-------------	--

---

## 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 <code>.robustDigest</code> .
algo	The digest algorithm to use. Default <code>xxhash64</code> (see <a href="#">digest::digest()</a> for others).
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 <code>do.call</code> .
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 <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,
  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"),
  bufferOK = FALSE,
  verbose = getOption("reproducible.verbose"),
  ...
)
```

### Arguments

targetFile	The (optional) local file (or path to file) name for a <code>sf</code> object or <code>data.frame</code> that can be coerced to a <code>sf</code> object (i.e., has a <code>geometry</code> column). If <code>cloudFolderID</code> is specified, then this will be the name of the file stored and/or accessed in that cloud folder.
domain	An <code>sf</code> polygon object that is the spatial area of interest. If <code>NULL</code> , then this will return the whole object in <code>targetFile</code> .
FUN	A function call that will be called if there is the domain is not already contained within the <code>sf</code> object at <code>targetFile</code> . This function call MUST return either a <code>sf</code> class object or a <code>data.frame</code> class object that has a <code>geometry</code> column (which can then be converted to <code>sf</code> with <code>sf::st_as_sf</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 <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 <code>NULL</code> meaning do not search locally.
useCloud	A logical.
cloudFolderID	If this is specified, then it must be either 1) a Google Drive url to a folder where the <code>targetFile</code> will be read from or written to, or 2) a <code>googledrive</code> id or 3) an absolute path to a (possibly non-existent yet) folder on your Google drive.
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 deleted 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 mean 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(...)</code> <code> &gt; Cache()</code> ).
overwrite	Logical. Passed to <code>writeTo</code> (possibly inside <code>postProcess</code> ) and <code>postProcess</code> .
action	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.
bufferOK	A logical. If <code>TRUE</code> , then after testing whether the <code>domain</code> is within the <code>targetFile</code> spatial object, and if it returns <code>FALSE</code> , then the function will create a larger object, buffered by 2.5% of the extent of the object. If <code>FALSE</code> , then it will be strict about whether the <code>domain</code> is within the <code>targetFile</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 <code>Caching</code> , which may help diagnose <code>Caching</code> challenges. Can set globally with an option, e.g., <code>options('reproducible.verbose' = 0)</code> to reduce the output.
...	All named objects that are needed for <code>FUN</code> , including the function itself, if it is not in a package.

## 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`. 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`.

### 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
  )
}
}

```

---

cacheId	<i>Extract the cache id of an object</i>
---------	--

---

### Description

Any object that was returned from the Cache or was calculated as part of a Cache call will have an attribute, tags and an entry with cacheId: prefix. This is a lightweight helper to extract that cacheId.

### Usage

cacheId(obj)

### Arguments

obj	Any R object
-----	--------------

### Value

The cacheId if this was part of a Cache call. Otherwise NULL

---

checkAndMakeCloudFolderID	<i>Check for presence of checkFolderID (for Cache(useCloud))</i>
---------------------------	--

---

### 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 googledrive::drive_mkdir. 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**

<code>files</code>	A character vector of files to check to see if they are the same as <code>knownRelativeFiles</code> , once the <code>absolutePrefix</code> is removed
<code>absolutePrefix</code>	A directory to "remove" from <code>files</code> to compare to <code>knownRelativeFiles</code>
<code>knownRelativeFiles</code>	A character vector of relative filenames, that could have sub-folder structure.
<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

---

**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 = identifyCHECKSUMTxtFile(path),
  files = NULL,
  verbose = getOption("reproducible.verbose", 1),
  ...
)

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

## S4 method for signature 'character,missing'
Checksums(
  path,
  write,
  quickCheck = getOption("reproducible.quickCheck", FALSE),
  checksumFile = identifyCHECKSUMTxtFile(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 digest, the notable argument is algo. For write.table, the notable argument is append.

## 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	<i>Download from cloud, if necessary</i>
---------------	--

---

## 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 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 <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., <code>RSQlite::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 the amount of output.

---

**compareNA***NA-aware comparison of two vectors*

---

**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**

v1	A vector
v2	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)
```

---

**convertCallToCommonFormat***Convert all ways of calling a function into canonical form, including defaults*

---

**Description**

e.g., stats::rnorm(1) → rnorm(n = 1, mean = 0, sd = 1)

**Usage**

```
convertCallToCommonFormat(call, usesDots, isSquiggly, .callingEnv)
```

**Arguments**

call	The full captured call as it was passed by user.
usesDots	Logical. Whether the original Cache call used ...
isSquiggly	Logical. Whether there are curly braces e.g., as in a pipe sequence.
.callingEnv	Environment. The environment from which Cache was called.

---

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 = getDrvgetOption("reproducible.drv", NULL),
  conn = getgetOption("reproducible.conn", NULL),
  verbose = getgetOption("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 `.reproducibleTempPath()`, which is unlikely to be very useful. Can be `NULL`, 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, <code>drv</code> must be an object that inherits from <code>DBIDriver</code> (e.g., <code>RSQLite::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

## 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 TRUE.
<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
```

### 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,
    cacheSaveFormat =getOption("reproducible.cacheSaveFormat", .rdsFormat),
    .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),
    cacheSaveFormat =getOption("reproducible.cacheSaveFormat", .rdsFormat),
    verbose,
    ...
  )

  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,
    cacheSaveFormat =getOption("reproducible.cacheSaveFormat"),
    obj = NULL,
    readOnly = FALSE
  )

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

```

## 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
cacheSaveFormat	The text string representing the file extension used normally by different save formats; currently only "rds" or "qs" (which now uses qs2 package. 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
...	Arguments passed to FUN, if FUN is not an expression.
obj	The optional object that is of interest; it may have an attribute "saveRawFile" that would be important.
readOnly	Logical. Only relevant during transition from qs to qs2. Essentially, during transition, qs objects can be read, but not saved. If TRUE then the CacheStoredFile can return a .qs file; if FALSE, then this will not be able to return qs; instead it will return qs2 files.
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)

```

---

**detectActiveCores** *Count Active Threads Based on CPU Usage*

---

## Description

This function counts the number of active system processes (threads) that match a given pattern and exceed a specified minimum CPU usage threshold. It works on Unix-like systems (e.g., Linux, macOS) and does not support Windows.

## Usage

```
detectActiveCores(pattern = "", minCPU = 50)
```

## Arguments

pattern	A character string used to filter process lines. Only processes whose command line matches this pattern will be considered. Default is "" (matches all).
minCPU	A numeric value specifying the minimum CPU usage (in percent) for a process to be considered active. Default is 50.

## Value

An integer representing the number of active threads matching the pattern and exceeding the CPU usage threshold. Returns NULL with a message if run on Windows.

**Note**

This function uses the `ps -ef` system command and regular expressions to parse CPU usage. It may not be portable across all Unix variants.

**Examples**

```
## Not run:
detectActiveCores(pattern = "R", minCPU = 30)

## End(Not run)
```

---

determineFilename	<i>Determine filename, either automatically or manually</i>
-------------------	---

---

**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 <a href="#">determineFilename()</a> .
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 <a href="#">writeTo()</a>

## 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 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,
  .callingEnv,
  ...
)

```

## 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 <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., <code>.zip</code> ). This may be useful when an R function is expecting an archive directly.
<code>targetFile</code>	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., cropped/reprojected/masked). Using <code>Cache</code> around <code>prepInputs</code> will do a sufficient job in these cases. See table in <a href="#">preProcess()</a> .
<code>neededFiles</code>	Character string giving the name of the file(s) to be extracted.
<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>quick</code>	Logical. This is passed internally to <a href="#">Checksums()</a> (the <code>quickCheck</code> argument), and to <a href="#">Cache()</a> (the <code>quick</code> argument). This results in faster, though less robust checking of inputs. See the respective functions.
<code>checksumFile</code>	A character string indicating the absolute path to the <code>CHECKSUMS.txt</code> file.
<code>dlFun</code>	Optional "download function" name, such as <code>"raster::getData"</code> , which does custom downloading, in addition to loading into R. Still experimental.
<code>checkSums</code>	A checksums file, e.g., created by <code>Checksums(..., write = TRUE)</code>
<code>url</code>	Optional character string indicating the URL to download from. If not specified, then no download will be attempted. If not entry exists in the <code>CHECKSUMS.txt</code> (in <code>destinationPath</code> ), an entry will be created or appended to. This <code>CHECKSUMS.txt</code> entry will be used in subsequent calls to <code>prepInputs</code> or <code>preProcess</code> , comparing the file on hand with the ad hoc <code>CHECKSUMS.txt</code> . See table in <a href="#">preProcess()</a> .
<code>needChecksums</code>	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. If TRUE then the download will overwrite an existing file if it exists.
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
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.
.callingEnv	The environment where the function was called from. Used to find objects, if necessary.
...	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),
.callingEnv = parent.frame(),
...
)

```

## 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
dlFun	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. Passed to <code>writeTo</code> (possibly inside <code>postProcess</code> ) and <code>postProcess</code> .
needChecksums	Logical indicating whether to generate checksums. ## TODO: add overwrite arg to the function?
.tempPath	Optional temporary path for internal file intermediate steps. Will be cleared on <code>.exit</code> from this function.
preDigest	The list of <code>preDigest</code> that comes from <code>CacheDigest</code> of an object
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> .
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
.callingEnv	The environment where the function was called from. Used to find objects, if necessary.
...	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>dlFun</code> which is passed to <code>preProcess</code> . See details and examples.

---

<code>extractFromArchive</code>	<i>Extract files from archive</i>
---------------------------------	-----------------------------------

---

## 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

---

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**

obj	A Raster* object (i.e., RasterLayer, RasterStack, RasterBrick)
allowMultiple	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.
returnList	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 if the object is not file-backed or does not have a method to recover the file-backed filename.

**Author(s)**

Eliot McIntire

---

fixErrorsIn*Fix common errors in GIS layers, using terra*

---

## 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*3-Step postProcess sequence for SpatRasters using gdalwarp*

---

## Description

DEFUNCT: Please use the `postProcessTo` functions.

`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., <code>options('reproducible.verbose' = 0)</code> 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

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.

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

```
if (require("terra", quietly = TRUE)) {
  # prepare dummy data -- 3 SpatRasters, 2 SpatVectors
  # need 2 SpatRaster
  rf <- system.file("ex/elev.tif", package = "terra")
  elev1 <- terra::rast(rf)

  # a polygon vector
  f <- system.file("ex/lux.shp", package = "terra")
  v0orig <- terra::vect(f)
  v <- v0orig[1:2, ]

  # utm <- terra::crs("epsg:23028") # $wkt
  utm <- "+proj=utm +zone=28 +datum=WGS84 +units=m +no_defs"
  vInUTM <- terra::project(v0orig, utm)
  vAsRasInLongLat <- terra::rast(v0orig, 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)
  t3a <- postProcessTo(elev1, to = rInUTM, maskTo = vInUTM)

}
```

---

## 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)
```

---

harmonizeCall	<i>Harmonize all forms of call</i>
---------------	------------------------------------

---

### Description

This will convert all known (imagined) calls so that they have the same canonical format i.e., `rnorm(n = 1, mean = 0, sd = 1)`

### Usage

```
harmonizeCall(callList, .callingEnv, .functionName = NULL)
```

### Arguments

<code>callList</code>	A named list with elements <code>call</code> , <code>usesDots</code> and <code>FUNorig</code>
<code>.callingEnv</code>	The calling environment where Cache was called from
<code>.functionName</code>	A possible function name. If omitted, then it will be deduced from the <code>callList</code> and may be inaccurate.

### Value

A named list. We illustrate with the example `rnorm(1)`. The named list will have the original `callList` (`call` (the original call, without quote), `FUNorig`, the original value passed by user to `FUN`, and `usesDots` which is a logical indicating whether the `...` are used), and appended with `new_call` (the harmonized call, with the function and arguments evaluated, e.g., `(function (n, mean = 0, sd = 1) .Call(C_rnorm, n, mean, sd))(1)`), `func_call`, the same harmonized call with neither function nor arguments not evaluated (e.g., `rnorm(1)`), `func` which will be function or method definition `(n, mean = 0, sd = 1) .Call(C_rnorm, n, mean, sd)`, and `.functionName`, which will be the function name as a character string (`rnorm`) either directly passed from the user's `.functionName` or deduced from the `func_call`.

---

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

*Has a cached object has been updated?*

**Description**

Has a cached object has been updated?

**Usage**

`isUpdated(x)`

**Arguments**

x                    cached object

**Value**

logical

keepOrigGeom

*Keep original geometries of sf objects*

**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. Passed to writeTo (possibly inside postProcess) and postProcess.
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)
}

```

**Description**

This is a convenience wrapper around `a <- 1; newList <- list(a); 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
```

**Description**

Load a file from the cache

**Usage**

```
loadFile(file, ...)
```

**Arguments**

file character specifying the path to the file  
... Allows format for backward compatibility

**Value**

the object loaded from file

---

`matchCall2`*Remove quote and determine if call uses ...*

---

**Description**

Minor cleaning up of the FUN and ... to be used subsequently. This does only very minor things as it is run even if useCache = FALSE, i.e., even if the Cache is skipped.

**Usage**

```
matchCall2(definition, call, envir, envir2 = parent.frame(), FUN)
```

**Arguments**

<code>definition</code>	a function, by default the function from which <code>match.call</code> is called. See details.
<code>call</code>	an unevaluated call to the function specified by <code>definition</code> , as generated by <code>call</code> .
<code>envir</code>	an environment, from which the ... in <code>call</code> are retrieved, if any.
<code>envir2</code>	Environment. The environment where <code>matchCall2</code> was called.
<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., <code>using quote()</code> ).

**Value**

A named list with `call1` (the original call, without quote), `FUNorig`, the original value passed by user to `FUN`, and `usesDots` which is a logical indicating whether the ... are used.

---

`mergeCache`*Merge two cache repositories together*

---

**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 = getDrvgetOption("reproducible.drv", NULL),  
  drvFrom = getDrvgetOption("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., options('reproducible.verbose' = 0) 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 handing 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***Get min or maximum value of a (Spat)Raster*

---

## 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

<code>x</code>	A <code>Raster</code> or <code>SpatRaster</code> object.
<code>...</code>	Passed to the functions in <code>raster</code> or <code>terra</code> , 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 = getDrvgetOption("reproducible.drv", NULL),  
  conn = getgetOption("reproducible.conn", NULL),  
  verbose = getgetOption("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., RSQLite:::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 reproducible 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(tempdir(), tmpCache), mustWork = FALSE)
tmpdirPath <- normalizePath(file.path(tempdir(), 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

*Normalize file paths*

## 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 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)
```

---

numCoresToUse

*Estimate Number of CPU Cores to Use for Parallel Processing*

---

## Description

This function estimates the number of CPU cores that can be safely used for parallel processing, taking into account a minimum threshold, the total number of physical cores, and currently active threads.

## Usage

```
numCoresToUse(min = 2, max = NULL)
```

## Arguments

min	An integer specifying the minimum number of cores to use. Default is 2.
max	An integer specifying the maximum number of cores available, typically the number of physical cores. Default is max(1L, getOption("Ncpus", 1L), parallel::detectCores() - 1, logical = FALSE, na.rm = TRUE).

## Value

An integer representing the number of cores that can be used for parallel tasks, ensuring at least min cores are used, while subtracting one for the current process and an estimate of actively used threads (via detectActiveCores()).

## Note

This function depends on detectActiveCores() and is not supported on Windows systems.

## See Also

[detectActiveCores\(\)](#)

## Examples

```
if (FALSE) {
  numCoresToUse()
  numCoresToUse(min = 4)
}
```

---

**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, recursive = FALSE, ...)  
objSizeSession(sumLevel = Inf, enclosingEnvs = TRUE, .prevEnvirs = list())
```

## Arguments

<code>x</code>	An object
<code>quick</code>	Logical. If <code>FALSE</code> , then an attribute, "objSize" will be added to the returned value, with each of the elements' object size returned also.
<code>recursive</code>	Logical. If <code>TRUE</code> , then, in addition to evaluating the whole object, it will also return the recursive sizes of the elements of a list or environment.
<code>...</code>	Additional arguments (currently unused), enables backwards compatible use.
<code>sumLevel</code>	Numeric, indicating at which depth in the list of objects should the object sizes be summed (summarized). Default is <code>Inf</code> , meaning no sums. Currently, the only option other than <code>Inf</code> is <code>1</code> : <code>objSizeSession(1)</code> , which gives the size of each package.
<code>enclosingEnvs</code>	Logical indicating whether to include enclosing environments. Default <code>TRUE</code> .
<code>.prevEnvirs</code>	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(prepInputs) # only the function, without its enclosing environment
objSize(prepInputs) # 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, <code>pad</code> will be used to pad.
<code>padR</code>	numeric. Desired number of digits on right side of decimal. If not enough, <code>pad</code> 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**

obj	A character string to convert to a Path.
nParentDirs	A numeric indicating the number of parent directories starting from basename(obj) = 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 <code>rlang::quosure</code> 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	<code>rasterToMatch</code>
resolution	Yes	No	<code>rasterToMatch</code>
projection	Yes	No*	<code>rasterToMatch</code> *
alignment	Yes	No	<code>rasterToMatch</code>
mask	No**	Yes	<code>studyArea</code> **

\*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("withr", quietly = TRUE)) {
  library(reproducible)
  withr::local_dir(withr::local_tempdir())
  withr::local_options(reproducible.inputPaths = NULL)
  # 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(prepInputs(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
if (requireNamespace("sf", quietly = TRUE)) {
  studyAreaSmall <- prepInputs(localFileLuxSm, fun = "sf::st_read")
  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()
studyAreaSmall <- prepInputs(localFileLuxSm)
studyAreas[["orig"]] <- prepInputs(localFileLux)
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))
}

withr::deferred_run()
# setwd(od)
}

```

---

postProcessTo

*Transform a GIS dataset so it has the properties (extent, projection, mask) of another*

---

## 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 template 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., <code>options('reproducible.verbose' = 0)</code> to reduce t
...	Arguments passed to <code>terra::mask</code> (for <code>maskTo</code> ), <code>terra::project</code> (for <code>projectTo</code> ) or <code>terra::writeRaster</code> (for <code>writeTo</code> ) and not used for <code>cropTo</code> , as well <code>postProcess</code> 's <code>rasterToMatch</code> and <code>studyArea</code> arguments (see below). Commonly used arguments might be <code>method</code> , <code>touches</code> , and <code>datatype</code> . If <code>filename</code> is passed, it will be ignored; use <code>writeTo = .</code> If <code>reproducible.gdalwarp = TRUE</code> , then these will be passed to the <code>gdal*</code> 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 <code>cropTo</code> , so that if a reprojection has to happen on the <code>cropTo</code> prior to using it as a crop layer, then a buffer of $1.5 * \text{res}(\text{cropTo})$ will occur prior, so that no edges are cut off.
<code>isStack</code> , <code>isBrick</code> , <code>isRaster</code> , <code>isSpatRaster</code>	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 `rasterToMatch` 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**

```
if (require("terra", quietly = TRUE)) {  
  # prepare dummy data -- 3 SpatRasters, 2 SpatVectors  
  # need 2 SpatRaster  
  rf <- system.file("ex/elev.tif", package = "terra")  
  elev1 <- terra::rast(rf)
```

```

# 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
utm <- "+proj=utm +zone=28 +datum=WGS84 +units=m +no_defs"
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)
t3a <- postProcessTo(elev1, to = rInUTM, maskTo = vInUTM)

}

```

---

**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**

<b>targetFile</b>	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 *before* 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 [preProcess\(\)](#).

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 <code>CHECKSUMS.txt</code> (in <code>destinationPath</code> ), an entry will be created or appended to. This <code>CHECKSUMS.txt</code> entry will be used in subsequent calls to <code>prepInputs</code> or <code>preProcess</code> , comparing the file on hand with the ad hoc <code>CHECKSUMS.txt</code> . See table in <a href="#">preProcess()</a> .
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 <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 NULL, 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> . NA 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 <a href="#">preProcess()</a> .
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.
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"), NA (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 <a href="#">Checksums()</a> (the <code>quickCheck</code> argument), and to <a href="#">Cache()</a> (the <code>quick</code> argument). This results in faster, though less robust checking of inputs. See the respective functions.
overwrite	Logical. Passed to <code>writeTo</code> (possibly inside <code>postProcess</code> ) and <code>postProcess</code> .
purge	Logical or Integer. 0/FALSE (default) keeps existing <code>CHECKSUMS.txt</code> file and <code>prepInputs</code> will write or append to it. 1/TRUE will deleted 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 mean it will use <code>Cache</code> only up to 2 nested levels, which includes <code>preProcess</code> . <code>postProcess</code>

	and its nested *Input 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(...)</code>  > <code>Cache()</code> ).
<code>.tempPath</code>	Optional temporary path for internal file intermediate steps. Will be cleared on <code>exit</code> from this function.
<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>...</code>	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>dlFun</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 d1Fun), 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, is 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.

### Author(s)

Eliot McIntire, Jean Marchal, and Tati Micheletti

### See Also

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

### Examples

```
if (requireNamespace("terra", quietly = TRUE) &&
  requireNamespace("withr", quietly = TRUE)) {
  library(reproducible)
  withr::local_dir(withr::local_tempdir())
  # 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(), "prepInputs2.shp"))
  terra::writeVector(outerSA, tf)

  # 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(), "ecozone")
  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, "ecozone.shp")
    ecozoneFiles <- c(
      "ecozone.dbf", "ecozone.prj",
      "ecozone.sbn", "ecozone.sbx", "ecozone.shp", "ecozone.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, "ecozone.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(
      "ecozone.dbf", "ecozone.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)
}
}
withr::deferred_run()
}

## 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)
##     }
##   )

```

---

**prepInputsWithTiles**    *Alternative to prepInputs that can use Spatial Tiles stored locally or on Google Drive*

---

## Description

Downloads, processes and optionally uploads a `SpatRaster` object through a tiling intermediary. If the original `url` is for a very large object, but `to` is a relatively small subset of the area represented by the spatial file at `url`, then this function will potentially by-pass the download of the large file at `url` and instead only download the minimum number of tiles necessary to cover the `to` area. When `doUploads` is `TRUE`, then this function will potentially create and upload the tiles to `tileFolder`, prior to returning the spatial object, `postProcessed` to `to`. This function supports both Google Drive and HTTP(S) URLs.

**Usage**

```
prepInputsWithTiles(
  targetFile,
  url,
  destinationPath,
  to,
  tilesFolder = file.pathgetOption("reproducible.inputPath"), "tiles"),
  urlTiles =getOption("reproducible.prepInputsUrlTiles", NULL),
  doUploads =getOption("reproducible.prepInputsDoUploads", FALSE),
  tileGrid = "CAN",
  numTiles = NULL,
  plot.grid = FALSE,
  purge = FALSE,
  verbose =getOption("reproducible.verbose"),
  ...
)
```

**Arguments**

targetFile	Character. Name of the target file to be downloaded or processed. If missing, it will be inferred from the URL or Google Drive metadata.
url	Character. URL to the full dataset (Google Drive or HTTP/S).
destinationPath	Character. Path to the directory where files will be downloaded and processed.
to	A spatial object (e.g., <code>SpatRaster</code> , <code>SpatVector</code> , <code>sf</code> , or <code>Spatial*</code> ) defining the area of interest.
tilesFolder	A local file path to put tiles. If this is an absolute path, then that will be used; if it is a relative path, then it will be <code>file.path(destinationPath, tilesFolder)</code>
urlTiles	Character. URL to the tile source (e.g., Google Drive folder or HTTP/S endpoint). Default is <code>getOption("reproducible.prepInputsUrlTiles", NULL)</code> .
doUploads	Logical. Whether to upload processed tiles. Default is <code>getOption("reproducible.prepInputsDoUploads", FALSE)</code> .
tileGrid	Either length 3 character string, such as "CAN", to be sent to <code>geodata::gadm(...)</code> or an actual <code>SpatVector</code> object with a grid of polygons
numTiles	Integer. Number of tiles to generate. Optional.
plot.grid	Logical. Whether to plot the tile grid and area of interest. Default is FALSE.
purge	Logical or Integer. 0/FALSE (default) keeps existing <code>CHECKSUMS.txt</code> file and <code>prepInputs</code> will write or append to it. 1/TRUE will deleted the entire <code>CHECKSUMS.txt</code> file.
verbose	Logical or numeric. Controls verbosity of messages. Default is <code>getOption("reproducible.verbose")</code> .
...	Either <code>maskTo</code> , <code>cropTo</code> (which will be used if <code>to</code> is not supplied, or arguments passed to <code>writeRaster</code> , e.g., <code>datatype</code> (used when writing tiles).

## Details

This function can be triggered *inside* prepInputs if the to is supplied and both url and urlTiles are supplied. **NOTE:** urlTiles can be supplied using the option(reproducible.prepInputsUrlTiles = someGoogleDrivePath) so the original prepInputs function call can remain unaffected.

This function also uses a different checksumming procedure compared to the normal prepInputs. This function will assess the remote url for a hash. If that hash exists, then it will compare it to a local file with targetFile name, suffixed with .hash. If the two hashes differ (remote and local), then it will be redownloaded; otherwise the local one will be returned.

This function is useful for working with large spatial datasets, but where the user only requires a "relatively small" section of that dataset. This function will potentially bypass the full download and download only the tiles that are necessary for the to. It handles downloading only the required tiles based on spatial intersection with the target area, and supports resumable downloads from Google Drive or HTTP/S sources.

If targetFile is missing, the function attempts to infer it from the URL using the Content-Disposition header or the basename of the URL. For Google Drive URLs, it uses the file metadata.

## Value

A single, merged SpatRaster object postProcessed to the area of interest (to), composed of the necessary tiles. If the post-processed file already exists locally, it will be returned directly.

## See Also

[googledrive::drive\\_get\(\)](#), [terra::rast\(\)](#), [terra::crop\(\)](#), [terra::merge\(\)](#)

## Examples

```
if (FALSE) {
  to <- sf::st_as_sf(sf::st_sf(sf::st_point(c(-123.3656, 48.4284)), crs = 4326))
  result <- prepInputsWithTiles(
    url = "https://example.com/data.tif",
    destinationPath = tempdir(),
    to = to,
    urlTiles = "https://example.com/tiles/",
    tileGrid = "CAN"
  )
}
```

---

## 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,
  .callingEnv = parent.frame(),
  ...
)
```

**Arguments**

<b>n</b>	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.
<b>targetFile</b>	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> .
<b>url</b>	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> .
<b>archive</b>	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.
<b>alsoExtract</b>	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 [preProcess\(\)](#).

**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 [Checksums\(\)](#) (the `quickCheck` argument), and to [Cache\(\)](#) (the `quick` argument). This results in faster, though less robust checking of inputs. See the respective functions.

**overwrite**

Logical. Passed to `writeTo` (possibly inside `postProcess`) and `postProcess`.

**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., `options('reproducible.verbose' = 0)` to reduce t

**.tempPath**

Optional temporary path for internal file intermediate steps. Will be cleared on `exit` from this function.

**.callingEnv**

The environment where the function was called from. Used to find objects, if necessary.

**...**

Additional arguments passed to [postProcess\(\)](#) and [Cache\(\)](#). Since `...` is passed to [postProcess\(\)](#), these will `...` will also be passed into the inner functions, e.g., [cropInputs\(\)](#). Possibly useful other arguments include `dlFun` which is passed to `preProcess`. 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

---

purgeChecksums

*Purge the checksums of a single file*

---

### Description

This is a manual way of achieving `prepInputs(..., purge = 7)`, useful in cases where `prepInputs` is not called directly by the user, so it would be difficult to set `purge = 7`.

### Usage

```
purgeChecksums(checksumFile, fileToRemove)
```

### Arguments

`checksumFile` A character string indicating the absolute path to the `CHECKSUMS.txt` file.  
`fileToRemove` The filename to remove from the `checksumFile`

### Value

NULL. Run for its side effect, namely, and file removed from the ‘`CHECKSUMS.txt`’ file.

---

rasterRead

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

---

### 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 =getOption("reproducible.cachePath"), ...)
```

## Arguments

obj	(optional) object whose file path metadata will be remapped
tags	cache tags <code>data.table</code> object
cachePath	character string specifying the path to the cache directory or <code>NULL</code>
...	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()`.
- `cacheChaining` Default: `FALSE`. Used in `Cache()` in the `.cacheChaining` argument.
- `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"` (which will use `qs2` package as of reproducible version `">= 2.1.3"`), `"qs2"`, 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()`.

**dryRun** Default: FALSE.

**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?

**leaveOnDisk** Default: TRUE. Used in `postProcess()`. When there is a `SpatRaster` object, should `postProcess` force any file-backed object, to use the file-based, memory-safe tools within `terra` (by temporarily setting `terraOption(memfrac = 0)`). Alternatively, if this is set to FALSE, then `postProcess` will let `terra` decide on its own based on its internal cues (largely based on `memfrac`, `maxmem` `terraOptions`). This will be ignored, however, if the user has set the `terraOptions` away from its default of 0.5. The default increases predictability of whether the returned object is on disk or in memory.

**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. Simplifying 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`.

`reproducible.useCacheV3` Default: TRUE. If this is set to FALSE, it will use the old Cache source code. This will only be available for a short period before it is deleted from the package. See also `reproducible.digestV3`. It is not guaranteed to be identical to using a previous version of `reproducible` (<3.0).

`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_DB`I. 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, `dlGoogle` 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.

`digestV3` Default: TRUE. This uses a digest approach that includes the names of list elements and several other tweaks that were created for `reproducible 3.x`. Set this to FALSE to use *some of* the previous cache digesting to achieve some backwards compatibility with the digest algorithms of `reproducible (<3.x)`. It will not be possible to get it exact for all classes of objects, particularly those with file-backing.

## 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

<b>expr</b>	An expression to run, i.e., <code>rnorm(1)</code> , similar to what is passed to <code>try</code>
<b>envir</b>	The environment in which to evaluate the quoted expression, default to <code>parent.frame(1)</code>
<b>retries</b>	Numeric. The maximum number of retries.
<b>exponentialDecayBase</b>	Numeric > 1.0. The delay between successive retries will be <code>runif(1, min = 0, max = exponentialDecayBase ^ i - 1)</code> where i is the retry number (i.e., follows <code>seq_len(retries)</code> )
<b>silent</b>	Logical indicating whether to <code>try</code> silently.
<b>exprBetween</b>	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).
<b>messageFn</b>	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`.

---

saveToCache

*Save an object to Cache*

---

## 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"),  
  cacheSaveFormat = getOption("reproducible.cacheSaveFormat"),  
  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 <code>Cache</code> is used inside a <code>SpaDES</code> module.
cacheSaveFormat	Character string: currently either <code>qs</code> or <code>rds</code> . Defaults to <code>getOption("reproducible.cacheSaveFormat")</code> . <code>qs</code> may be faster but appears to have narrower range of conditions that work; <code>rds</code> is safer, and may be slower.
drv	If using a database backend, <code>drv</code> must be an object that inherits from <code>DBIDriver</code> (e.g., <code>RSSQLite::SQLite</code> ).
conn	an optional <code>DBIConnection</code> object, as returned by <code>dbConnect()</code> .
obj	The R object to save to the cache
userTags	A character vector with descriptions of the <code>Cache</code> function call. These will be added to the <code>Cache</code> so that this entry in the <code>Cache</code> can be found using <code>userTags</code> e.g., via <a href="#">showCache()</a> .

cacheId	The hash string representing the result of <code>.robustDigest</code>
linkToCacheId	Optional. If a <code>cacheId</code> is provided here, then a <code>file.link</code> will be made to the file with that <code>cacheId</code> 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., <code>parent.frame()</code>
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 = getDrvgetOption("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 = getDrvgetOption("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 = getDrvgetOption("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 = getDrvgetOption("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" = 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., RSQLite:::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 the amount of output.
...	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

*Get a unique name for a given study area*

### 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

studyArea	Spatial object.
...	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>
-----------	---------------------------------------

---

**Description**

The known path for unrar or 7z

**Usage**

```
.systemArchivePath
```

**Format**

An object of class `NULL` of length 0.

---

usesPointer	<i>Does an object use a pointer?</i>
-------------	--------------------------------------

---

## Description

Does an object use a pointer?

## Usage

```
usesPointer(x)
```

## Arguments

x	an object
---	-----------

## Value

logical

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writeFuture	<i>Write to cache repository, using future::future</i>
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## 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,  
  verbose = getOption("reproducible.verbose")  
)
```

### 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., RSQLite::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.
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

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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