Package ‘reproducible’

May 30, 2024

Type Package

Title Enhance Reproducibility of R Code

Description A collection of high-level, machine- and OS-independent tools for making reproducible and reusable content in R. The two workhorse functions are Cache() and prepInputs(). Cache() allows for nested caching, is robust to environments and objects with environments (like functions), and has deals with some classes of file-backed R objects e.g., from terra and raster packages. Both functions have been developed to be foundational components of data retrieval and processing in continuous workflow situations. In both functions, efforts are made to make the first and subsequent calls of functions have the same result, but faster at subsequent times by way of checksums and digesting. Several features are still under development, including cloud storage of cached objects, allowing for sharing between users. Several advanced options are available, see ‘reproducibleOptions()’.

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

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

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Depends R (>= 4.1)

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

Suggests archive, covr, crayon, DBI, future, geodata, glue, googledrive, httr, knitr, parallel, qs, raster (>= 3.5-15), RCpp (>= 1.0.4.8), rlang, rmarkdown, RSQLite, R.utils, sp, terra (>= 1.4-2), testthat, withr

Additional_repositories https://predictiveecology.r-universe.dev/

Encoding UTF-8

Language en-CA
License GPL-3

VignetteBuilder knitr, rmarkdown

BugReports https://github.com/PredictiveEcology/reproducible/issues

ByteCompile yes

RoxygenNote 7.3.1

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

NeedsCompilation no

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R topics documented:

reproducible-package .................................................... 4
    .debugCache ...................................................... 5
    .file.move .......................................................... 6
    .isMemoised .......................................................... 6
    .prefix .............................................................. 7
    .prepareFileBackedRaster ......................................... 8
    .removeCacheAtts .................................................. 9
    .requireNamespace ................................................ 9
    .setSubAttrInList ................................................. 10
    .wrap .............................................................. 10
    assessDataType ...................................................... 13
    basename2 .......................................................... 16
    Cache .............................................................. 16
    CacheDigest ........................................................ 25
    CacheGeo ........................................................... 26
    checkAndMakeCloudFolderID ...................................... 29
R topics documented:

<table>
<thead>
<tr>
<th>Function</th>
<th>Page</th>
</tr>
</thead>
<tbody>
<tr>
<td>checkPath</td>
<td>30</td>
</tr>
<tr>
<td>checkRelative</td>
<td>31</td>
</tr>
<tr>
<td>Checksums</td>
<td>32</td>
</tr>
<tr>
<td>cloudDownload</td>
<td>34</td>
</tr>
<tr>
<td>compareNA</td>
<td>35</td>
</tr>
<tr>
<td>convertPaths</td>
<td>36</td>
</tr>
<tr>
<td>Copy</td>
<td>37</td>
</tr>
<tr>
<td>copySingleFile</td>
<td>39</td>
</tr>
<tr>
<td>createCache</td>
<td>40</td>
</tr>
<tr>
<td>determineFilename</td>
<td>44</td>
</tr>
<tr>
<td>downloadFile</td>
<td>45</td>
</tr>
<tr>
<td>downloadRemote</td>
<td>47</td>
</tr>
<tr>
<td>extractFromArchive</td>
<td>49</td>
</tr>
<tr>
<td>fastMask</td>
<td>50</td>
</tr>
<tr>
<td>Filenames</td>
<td>51</td>
</tr>
<tr>
<td>fixErrorsIn</td>
<td>52</td>
</tr>
<tr>
<td>gdalProject</td>
<td>53</td>
</tr>
<tr>
<td>getRelative</td>
<td>55</td>
</tr>
<tr>
<td>internetExists</td>
<td>56</td>
</tr>
<tr>
<td>isUpdated</td>
<td>57</td>
</tr>
<tr>
<td>keepOrigGeom</td>
<td>57</td>
</tr>
<tr>
<td>linkOrCopy</td>
<td>58</td>
</tr>
<tr>
<td>listNamed</td>
<td>60</td>
</tr>
<tr>
<td>loadFile</td>
<td>60</td>
</tr>
<tr>
<td>mergeCache</td>
<td>61</td>
</tr>
<tr>
<td>messageDF</td>
<td>62</td>
</tr>
<tr>
<td>minFn</td>
<td>64</td>
</tr>
<tr>
<td>movedCache</td>
<td>65</td>
</tr>
<tr>
<td>normPath</td>
<td>66</td>
</tr>
<tr>
<td>objSize</td>
<td>68</td>
</tr>
<tr>
<td>paddedFloatToChar</td>
<td>69</td>
</tr>
<tr>
<td>Path-class</td>
<td>70</td>
</tr>
<tr>
<td>postProcess</td>
<td>71</td>
</tr>
<tr>
<td>postProcessTo</td>
<td>74</td>
</tr>
<tr>
<td>prepInputs</td>
<td>79</td>
</tr>
<tr>
<td>preProcessParams</td>
<td>85</td>
</tr>
<tr>
<td>rasterRead</td>
<td>88</td>
</tr>
<tr>
<td>remapFilenames</td>
<td>88</td>
</tr>
<tr>
<td>reproducibleOptions</td>
<td>89</td>
</tr>
<tr>
<td>retry</td>
<td>92</td>
</tr>
<tr>
<td>saveToCache</td>
<td>93</td>
</tr>
<tr>
<td>searchFull</td>
<td>94</td>
</tr>
<tr>
<td>set.randomseed</td>
<td>95</td>
</tr>
<tr>
<td>showCache</td>
<td>96</td>
</tr>
<tr>
<td>studyAreaName</td>
<td>100</td>
</tr>
<tr>
<td>tempdir2</td>
<td>101</td>
</tr>
<tr>
<td>tempfile2</td>
<td>102</td>
</tr>
<tr>
<td>unrarPath</td>
<td>102</td>
</tr>
</tbody>
</table>
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.

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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

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

Usage

/debugCache(obj, preDigest, ...)

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>obj</td>
<td>An arbitrary R object.</td>
</tr>
<tr>
<td>preDigest</td>
<td>A list of hashes.</td>
</tr>
<tr>
<td>...</td>
<td>Dots passed from Cache</td>
</tr>
</tbody>
</table>

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

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

Value

Logical indicating whether operation succeeded.

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

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

Copy the file-backing of a file-backed Raster* object

Description

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

Usage

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

Arguments

- **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, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite
- **conn**: an optional DBIConnection object, as returned by dbConnect().
- **...**: Not used

Value

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

Author(s)

Eliot McIntire
.removeCacheAtts

Remove attributes that are highly varying

Description
Remove attributes that are highly varying

Usage
.removeCacheAtts(x)

Arguments
x Any arbitrary R object that could have attributes

.requireNamespace

Provide standard messaging for missing package dependencies

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

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

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

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

*Set subattributes within a list by reference*

**Description**

Sets only a single element within a list attribute.

**Usage**

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

**Arguments**

- **object**
  
  An arbitrary object

- **attr**
  
  The attribute name (that is a list object) to change

- **subAttr**
  
  The list element name to change

- **value**
  
  The new value

**Value**

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

### .wrap

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

**Description**

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

**Usage**

```
.wrap(
    obj,
    cachePath,
    preDigest,
    drv = getDrv(getOption("reproducible.drv", NULL)),
    conn = getOption("reproducible.conn", NULL),
    verbose = getOption("reproducible.verbose"),
    outputObjects = NULL,
    ...
)
```
## S3 method for class 'list'
\[\text{\texttt{\textbackslash wrap}}(\text{\texttt{obj}}, \text{\texttt{cachePath}}, \text{\texttt{preDigest}}, \text{\texttt{drv}} = \text{\texttt{getDrv(getOption("reproducible.drv", NULL))}}, \text{\texttt{conn}} = \text{\texttt{getOption("reproducible.conn", NULL)}}, \text{\texttt{verbose}} = \text{\texttt{getOption("reproducible.verbose")}}, \text{\texttt{outputObjects}} = \text{\texttt{NULL}}, \text{\texttt{...}})\]

## S3 method for class 'environment'
\[\text{\texttt{\textbackslash wrap}}(\text{\texttt{obj}}, \text{\texttt{cachePath}}, \text{\texttt{preDigest}}, \text{\texttt{drv}} = \text{\texttt{getDrv(getOption("reproducible.drv", NULL))}}, \text{\texttt{conn}} = \text{\texttt{getOption("reproducible.conn", NULL)}}, \text{\texttt{verbose}} = \text{\texttt{getOption("reproducible.verbose")}}, \text{\texttt{outputObjects}} = \text{\texttt{NULL}}, \text{\texttt{...}})\]

## Default S3 method:
\[\text{\texttt{\textbackslash wrap}}(\text{\texttt{obj}}, \text{\texttt{cachePath}}, \text{\texttt{preDigest}}, \text{\texttt{drv}} = \text{\texttt{getDrv(getOption("reproducible.drv", NULL))}}, \text{\texttt{conn}} = \text{\texttt{getOption("reproducible.conn", NULL)}}, \text{\texttt{verbose}} = \text{\texttt{getOption("reproducible.verbose")}}, \text{\texttt{...}})\]

## Default S3 method:
\[\text{\texttt{\textbackslash unwrap}}(\text{\texttt{obj}}, \text{\texttt{cachePath}}, \text{\texttt{cacheId}}, \text{\texttt{drv}} = \text{\texttt{getDrv(getOption("reproducible.drv", NULL))}}, \text{\texttt{conn}} = \text{\texttt{getOption("reproducible.conn", NULL)}}, \text{\texttt{...}})\]

\[\text{\texttt{\textbackslash unwrap}}(\text{\texttt{obj}})\]
cachePath,
cacheId,
drv = getDrv(getOption("reproducible.drv", NULL)),
conn = getOption("reproducible.conn", NULL),
...
)

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

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

Arguments

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

Value

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

Examples

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

assessDataType  Assess the appropriate raster layer data type

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
      rasOrg <- terra::rast(ncols = 10, nrows = 10)
      ras <- rasOrg
      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[] <- c(0, NA, rep(c(0,1),49))
assessDataType(ras)

## INT5
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,
  ..., 
  notOlderThan = NULL,
  .objects = NULL,
  .cacheExtra = NULL,
  .functionName = NULL,
  outputObjects = NULL,
  algo = "xxhash64",
)
cacheRepo = NULL,
cachePath = NULL,
length = getOption("reproducible.length", Inf),
compareRasterFileLength,
userTags = c(),
omitArgs = NULL,
classOptions = list(),
debugCache = character(),
sideEffect = FALSE,
makeCopy = FALSE,
quick = getOption("reproducible.quick", FALSE),
verb = getOption("reproducible.verb", 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., rnorm), a function call (e.g., rnorm(1)), or an unevaluated function call (e.g., using quote).

... Arguments passed to FUN, if FUN is not an expression.

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.

outputObjects Optional character vector indicating which objects to return. This is only relevant for list, environment (or similar) objects

algo The algorithms to be used; currently available choices are md5, which is also the default, sha1, crc32, sha256, sha512, xxhash32, xxhash64, murmur32, spookyhash, blake3, crc32c, xxh3_64, and xxh3_128.

.cachePath Same as cachePath, but kept for backwards compatibility.

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 is getOption("reproducible.length"), which is set to Inf.

compareRasterFileLength
Being deprecated; use length.

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 showCache().

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(...).

sideEffect Now deprecated. Logical or path. Determines where the function will look for new files following function completion. See Details. NOTE: this argument is experimental and may change in future releases.

makeCopy Now deprecated. Ignored if used.

quick Logical or character. If TRUE, no disk-based information will be assessed, i.e., only memory content. See Details section about quick in Cache().

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 to minimal

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.

useCache Logical, numeric or "overwrite" or "devMode". See details.

useCloud Logical. See Details.

cloudFolderID A googledrive dribble of a folder, e.g., using drive_mkdir(). If left as NULL, the function will create a cloud folder with name from last two folder levels of the cachePath path: paste0(basename(dirname(cachePath)), "_", basename(cachePath)). This cloudFolderID will be added to options("reproducible.cloudFolderId")
but this will not persist across sessions. If this is a character string, it will treat this as a folder name to create or use on GoogleDrive.

**showSimilar**  
A logical or numeric. Useful for debugging. If `TRUE` or 1, then if the Cache does not find an identical archive in the `cachePath`, it will report (via message) the next most recent similar archive, and indicate which argument(s) is/are different. If a number larger than 1, then it will report the N most recent similar archived objects.

**drv**  
if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite

**conn**  
an optional DBIConnection object, as returned by `dbConnect()`.

**Details**

There are other similar functions in the R universe. This version of Cache has been used as part of a robust continuous workflow approach. As a result, we have tested it with many "non-standard" R objects (e.g., RasterLayer, terra 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 terra* 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 then 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 = "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 <- file.path(tempdir())
opts <- options(reproducible.cachePath = tmpDir)

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

a[[1]] <- Cache(rnorm(1)) # no evaluation prior to Cache
a[[2]] <- Cache(rnorm, 1) # no evaluation prior to Cache
a[[3]] <- Cache(do.call, rnorm, list(1))
a[[4]] <- Cache(do.call(rnorm, list(1)))
a[[5]] <- Cache(do.call(b$fun, list(1)))
a[[6]] <- Cache(do.call, b$fun, list(1))
a[[7]] <- Cache(b$fun, 1)
a[[8]] <- Cache(b$fun(1))
a[[9]] <- Cache(quote(rnorm(1)))
a[[10]] <- Cache(stats::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
Cache

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

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

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

#################################################
# Advanced examples
#################################################
# .cacheExtra -- add something to digest
Cache(rnorm(1), .cacheExtra = "sfesee11") # adds something other than fn args
Cache(rnorm(1), .cacheExtra = "nothing") # even though fn is same, the extra is different

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

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

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

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

#################################################
# devMode -- enables cache database to stay
# small even when developing code
#################################################
opt <- options("reproducible.useCache" = "devMode")
clearCache(tmpDir, ask = FALSE)
centralTendency <- function(x) {
  mean(x)
}
funnyData <- c(1, 1, 1, 1, 10)
```
uniqueUserTags <- c("thisIsUnique", "reallyUnique")
ranNumsB <- Cache(centralTendency, funnyData, cachePath = tmpDir, 
                   userTags = uniqueUserTags) # sets new value to Cache
showCache(tmpDir) # 1 unique cacheId -- cacheId is 71cd24ec3b0d0cac

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

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

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

---

**CacheDigest**

The exact digest function that Cache uses

**Description**

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

**Usage**

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

**Arguments**

- `objsToDigest` A list of all the objects (e.g., arguments) to be digested
- `...` passed to `.robustDigest`. 
The algorithms to be used; currently available choices are md5, which is also the default, sha1, crc32, sha256, sha512, xxhash32, xxhash64, mmur32, spookyhash, blake3, crc32c, xxh3_64, and xxh3_128.

calledFrom

A character string, length 1, with the function to compare with. Default is "Cache". All other values may not produce robust CacheDigest results.

.functionName

A an arbitrary character string that provides a name that is different than the actual function name (e.g., "rnorm") which will be used for messaging. This can be useful when the actual function is not helpful for a user, such as do.call.

quick

Logical or character. If TRUE, no disk-based information will be assessed, i.e., only memory content. See Details section about quick in Cache().

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

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>targetFile</td>
<td>The (optional) local file (or path to file)</td>
</tr>
<tr>
<td>url</td>
<td>The (optional) url of the object on Google Drive (the only option currently). This is only for downloading and uploading to.</td>
</tr>
<tr>
<td>domain</td>
<td>An sf polygon object that is the spatial area of interest. If NULL, then this will return the whole object in targetFile.</td>
</tr>
<tr>
<td>FUN</td>
<td>A function call that will be called if there is the domain is not already contained within the sf object at url or targetFile. This function call MUST return either a sf class object or a data.frame class object that has a geometry column (which can then be converted to sf with st_as_sf)</td>
</tr>
<tr>
<td>destinationPath</td>
<td>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(&quot;reproducible.inputPaths&quot;) 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.</td>
</tr>
<tr>
<td>useCloud</td>
<td>A logical.</td>
</tr>
<tr>
<td>cloudFolderID</td>
<td>If this is specified, then it must be either 1) a googledrive url to a folder where the targetFile will be read from or written to, or 2) a googledrive id or 3) an absolute path to a (possibly non-existent yet) folder on your google drive.</td>
</tr>
<tr>
<td>purge</td>
<td>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.</td>
</tr>
<tr>
<td>useCache</td>
<td>Passed to Cache in various places. Defaults togetOption(&quot;reproducible.useCache&quot;, 2L) in prepInputs, and getOption(&quot;reproducible.useCache&quot;, FALSE) if calling any of the inner functions manually. For prepInputs, this mean it will use Cache only up to 2 nested levels, which includes preProcess, postProcess and its nested *Input functions (e.g., cropInputs, projectInputs, maskInputs) are no longer internally cached, as terra processing speeds mean internal caching is more time consuming. We recommend caching the full prepInputs call instead (e.g. prepInputs(…)</td>
</tr>
<tr>
<td>overwrite</td>
<td>Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.</td>
</tr>
<tr>
<td>action</td>
<td>A character string, with one of c(&quot;nothing&quot;, &quot;update&quot;, &quot;replace&quot;, &quot;append&quot;). Partial matching is used (&quot;n&quot; is sufficient). nothing will prevent any updating of the targetFile, i.e., &quot;read only&quot;. append will add the spatial elements in domain to targetFile (and writing it back to disk), update will do the same as append, but will also remove any identical geometries before appending. replace does nothing currently.</td>
</tr>
</tbody>
</table>
...

Any named objects that are needed for FUN

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 passed is within the geographic limits of the `targetFile` or file located at the url. This is similar to a geospatial operation on a remote GIS server, with 2 differences:

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

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

**Value**

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

**Examples**

```r
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")))),
    destinationPath = dPath,
    action = "update"
    #, cloudFolderID = "cachedObjects" # to upload/download from cloud
  )
}

checkAndMakeCloudFolderID

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

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

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

Arguments
cloudFolderID The google folder ID where cloud caching will occur.

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

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 team_drive Logical indicating whether to check team drives.

Value

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

checkPath path

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

```r
## 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)
```

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

```r
cHECKRelative(
  files,
  absolutePrefix,
  knownRelativeFiles,
  verbose =getOption("reproducible.verbose")
)
```
Arguments

- **files**: A character vector of files to check to see if they are the same as **knownRelativeFiles**, once the **absolutePrefix** is removed.
- **absolutePrefix**: A directory to "remove" from files to compare to **knownRelativeFiles**.
- **knownRelativeFiles**: A character vector of relative filenames, that could have sub-folder structure.
- **verbose**: Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., `options('reproducible.verbose' = 0)` to reduce the information output.

Checksums

**Calculate checksum**

Description

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

Usage

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

### S4 method for signature 'character,logical'

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

### S4 method for signature 'character,missing'

```r
Checksums(path, write,  
```
Checksums

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

Arguments

path Character string giving the directory path containing CHECKSUMS.txt file, or
where it will be written if checksumFile = TRUE.
write Logical indicating whether to overwrite CHECKSUMS.txt. Default is FALSE, as
users should not change this file. Module developers should write this file prior
to distributing their module code, and update accordingly when the data change.
quickCheck Logical. If TRUE, then this will only use file sizes, rather than a digest::digest
hash. This is generally faster, but will be much 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 messag-
ing, 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 digest::digest() and utils::write.table(). 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 be-
tween 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

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

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

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

## End(Not run)
```

cloudDownload

*Download from cloud, if necessary*

Description

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

Usage

```r
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.
compareNA

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: \texttt{paste0(basename(dirname(cachePath)), ", \", basename(cachePath))}. This cloudFolderID will be added to options("reproducible.cloudFolderID") but this will not persist across sessions. If this is a character string, it will treat this as a folder name to create or use on GoogleDrive.

drv  if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite

conn  an optional DBIConnection object, as returned by dbConnect().

verbose  Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce this.

---

<table>
<thead>
<tr>
<th>compareNA</th>
<th>NA-aware comparison of two vectors</th>
</tr>
</thead>
</table>

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

\texttt{compareNA(v1, v2)}

Arguments

- \texttt{v1} A vector
- \texttt{v2} A vector

Value

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

Examples

\texttt{a <- c(NA, 1, 2, NA)}
\texttt{b <- c(1, NA, 2, NA)}
\texttt{compareNA(a, b)}
convertPaths

Change the absolute path of a file

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)
Recursive copying of nested environments, and other "hard to copy"
objects

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

Usage
Copy(object, ...)

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

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

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

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

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

Arguments
object An R object (likely containing environments) or an environment.
... Only used for custom Methods
filebackedDir A directory to copy any files that are backing R objects, currently only valid for Raster classes. Defaults to .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.
**Copy**

**drv**  if using a database backend, drv must be an object that inherits from DBIDriver
e.g., from package RSQLite, e.g., SQLite

**conn**  an optional DBIConnection object, as returned by dbConnect().

**verbose**  Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce the

**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

**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$one
f$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
copySingleFile

## 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**

```r
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**

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

Low-level functions to create and work with a cache

Description

These are intended for advanced use only.

Usage

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

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

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

extractFromCache(sc, elem, ifNot = NULL)

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

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

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

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

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

CacheIsACache(
createCache

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

### Arguments

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>cachePath</td>
<td>A path describing the directory in which to create the database file(s)</td>
</tr>
<tr>
<td>drv</td>
<td>A driver, passed to <code>dbConnect</code></td>
</tr>
<tr>
<td>conn</td>
<td>an optional DBIConnection object, as returned by <code>dbConnect()</code>.</td>
</tr>
<tr>
<td>force</td>
<td>Logical. Should it create a cache in the <code>cachePath</code>, even if it already exists, overwriting.</td>
</tr>
<tr>
<td>verbose</td>
<td>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 to minimal.</td>
</tr>
<tr>
<td>cacheId</td>
<td>The cacheId or otherwise digested hash value, as character string.</td>
</tr>
<tr>
<td>preDigest</td>
<td>The list of <code>preDigest</code> that comes from <code>CacheDigest</code> of an object</td>
</tr>
<tr>
<td>fullCacheTableForObj</td>
<td>The result of <code>showCache</code>, but subsetted for only the cacheId being loaded or selected</td>
</tr>
<tr>
<td>format</td>
<td>The text string representing the file extension used normally by different save formats; currently only &quot;rds&quot; or &quot;qs&quot;. Defaults to <code>getOption(&quot;reproducible.cacheSaveFormat&quot;, &quot;rds&quot;)</code>.</td>
</tr>
<tr>
<td>.functionName</td>
<td>Optional. Used for messaging when this function is called from Cache</td>
</tr>
<tr>
<td>.dotsFromCache</td>
<td>Optional. Used internally.</td>
</tr>
<tr>
<td>sc</td>
<td>a cache tags data.table object</td>
</tr>
<tr>
<td>elem</td>
<td>character string specifying a tagKey value to match</td>
</tr>
<tr>
<td>ifNot</td>
<td>character (or NULL) specifying the return value to use if <code>elem</code> not matched</td>
</tr>
<tr>
<td>obj</td>
<td>The optional object that is of interest; it may have an attribute &quot;saveRawFile&quot; that would be important.</td>
</tr>
<tr>
<td>create</td>
<td>Logical. Currently only affects non RSQLite default drivers. If TRUE and there is no Cache database, the function will create one.</td>
</tr>
</tbody>
</table>

### 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;
- `CacheStoreFile()` 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
determineFilename

Determine filename, either automatically or manually

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

Arguments

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 .prefix(basename(filename1), prefix). If a character string, it will use this as its file name. See determineFilename().

filename1 Character strings giving the file paths of the input 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., options('reproducible.verbose' = 0) to reduce the

prefix The character string to prepend to filename1, if filename2 not provided.

... Passed into writeTo()
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 straightforward.

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 refered to as targetFile) and the filename2 is the file name of the of post-processed file.

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

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

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

downloadFile

A wrapper around a set of downloading functions

Description

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

Usage

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

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 preProcess(). If it is NA, then it will not 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 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().

neededFiles  Character string giving the name of the file(s) to be extracted.

destinationPath  Character string of a directory in which to download and save the file that comes from 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.

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.

checksumFile  A character string indicating the absolute path to the CHECKSUMS.txt file.

dlFun  Optional "download function" name, such as "raster::getData", which does custom downloading, in addition to loading into R. Still experimental.

checkSums  A checksums file, e.g., created by Checksums(..., write = TRUE)

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 preProcess().

needCheckums  A numeric, with 0 indicating do not write a new checksums, 1 write a new one, 2 append new information to existing one.

preDigest  The list of preDigest that comes from CacheDigest of an object
**overwrite**
Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.

**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 verbosity.

**purge**
Logical or Integer. `0/FALSE` (default) keeps existing `CHECKSUMS.txt` file and `prepInputs` will write or append to it. `1/TRUE` will deleted the entire `CHECKSUMS.txt` file. Other options, see details.

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

**...**
Passed to `dlFun`. Still experimental. Can be e.g., `type` for google docs.

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

**Author(s)**
Eliot McIntire

---

**downloadRemote**

*Download a remote file*

**Description**
Download a remote file

**Usage**

```r
downloadRemote(
  url,  
  archive,  
  targetFile,  
  checkSums,  
  dlFun = NULL,  
  fileToDownload,  
  messSkipDownload,  
  destinationPath,  
  overwrite,  
  needChecksums,  
  .tempPath,  
  preDigest,  
  verbose = optarg("reproducible.verbose", 1),  
  ...  
)
```


Arguments

url
Optional character string indicating the URL to download from. If not specified, then no download will be attempted. If not entry exists in the CHECKSUMS.txt (in destinationPath), an entry will be created or appended to. This CHECKSUMS.txt entry will be used in subsequent calls to prepInputs or preProcess, comparing the file on hand with the ad hoc CHECKSUMS.txt. See table in preProcess().

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 preProcess(). If it is NA, then it will \textit{not} 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 \textit{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().

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. \textbf{NOTE} (still experimental): To prevent repeated downloads in different locations, the user can also set options("reproducible.inputPaths") to one or more local file paths to search for the file before attempting to download. Default for that option is NULL meaning do not search locally.

overwrite
Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.

needChecksums
Logical indicating whether to generate checksums. ## TODO: add overwrite arg to the function?

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

preDigest
The list of preDigest that comes from CacheDigest of an object

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 ...
extractFromArchive  

Extract files from archive

Description

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

Usage

```r
evaluateFromArchive(
  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),
  ...,
)
```

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`: Character string giving the name of the file(s) to be extracted.
- `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 this.

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

...  Passed to unzip or untar, e.g., `overwrite`

**Value**

A character vector listing the paths of the extracted archives.

**Author(s)**

Jean Marchal and Eliot McIntire

---

**fastMask**  

*Faster operations on rasters (DEPRECATED because terra::mask is fast)*

**Description**

Deprecated. Use `maskTo()`.

**Usage**

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

**Arguments**

- **x**  A Raster* object.
- **y**  A SpatialPolygons object. If it is not in the same projection as x, it will be reprojected on the fly to that of x
- **cores**  An integer* or 'AUTO'. This will be used if gdalwarp is triggered. 'AUTO' will calculate 90% of the total number of cores in the system, while an integer or rounded float will be passed as the exact number of cores to be used.
Filenames

**useGDAL**  
Deprecated. Logical or "force". This is defunct; internals now can use terra if `options("reproducible.useTerra" = TRUE)`, which is not (yet) the default.

**verbose**  
Numeric. -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., `options('reproducible.verbose' = 0)` to reduce this.

... Currently unused.

**messageSkipDeprecated**  
Logical. If TRUE, then the message about this function being deprecated will be suppressed.

**Value**

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

**Author(s)**

Eliot McIntire

---

**Filenames**  
*Return the filename(s) from a Raster* object

**Description**

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

**Usage**

Filenames(obj, allowMultiple = TRUE, returnList = FALSE)

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

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

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

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

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

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 is the object is not file-backed or does not have a method to recover the file-backed filename.

Author(s)

Eliot McIntire

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**

- **x**
  The SpatStat or SpatVect object to try to fix.

- **error**
  The error message, e.g., coming from `try(...)`

- **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 to minimal.

- **fromFnName**
  The function name that produced the error, e.g., `maskTo`

**Value**

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

---

**Description**

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

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

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

**Usage**

```r

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 postProcessTo(), but can only be a SpatRaster.
toRas see to argument from postProcessTo(), but can only be a SpatRaster.
filenameDest A filename with an appropriate extension (e.g., .tif) for gdal to write the output to. Since this function is conceived to be part of a chain, and not the final step, this function does not use writeTo, which is reserved for the final step in the chain.
verbose Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce to minimal ...

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 maskTo(), but can only be a SpatVector
writeTo Optional character string of a filename to use writeRaster to save the final object. Default is NULL, which means there is no writeRaster

Details

These three functions are used within postProcessTo, in the sequence: gdalProject, gdalResample and gdalMask, when from and projectTo are SpatRaster and maskTo is a SpatVector, but only if options(reproducible.gdalwarp = TRUE) is set.

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

See Also
gdalResample(), and gdalMask() and the overarching postProcessTo()

Examples

# prepare dummy data -- 3 SpatRasters, 2 SpatVectors
# need 2 SpatRaster
rf <- system.file("ex/elev.tif", package = "terra")
elev1 <- terra::rast(rf)
#'
ras2 <- terra::deepcopy(elev1)
ras2[ras2 > 200 & ras2 < 300] <- NA_integer_
terra::values(elev1) <- rep(1L, terra::ncell(ras2))

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

#
utt <- terra::crs("epsg:23028") # $wkt
vInUTM <- terra::project(vOrig, utm)
vAsRasInLongLat <- terra::rast(vOrig, resolution = 0.008333333)
res100 <- 100
rInUTM <- terra::rast(vInUTM, resolution = res100)

# crop, reproject, mask, crop a raster with a vector in a different projection
# --> gives message about not enough information
t1 <- postProcessTo(elev1, to = vInUTM)

# crop, reproject, mask a raster to a different projection, then mask
t2a <- postProcessTo(elev1, to = vAsRasInLongLat, maskTo = vInUTM)

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

---

getRelative  Relative paths

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

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

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

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

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

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

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

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

**internetExists**

Checks for existed of a url or the internet using [https://CRAN.R-project.org](https://CRAN.R-project.org)

**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

**Usage**

- `isUpdated(x)`

**Arguments**

- **x**
  - cached object

**Value**

- logical

---

### keepOrigGeom

**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

```r
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`  
**Hardlink, symlink, or copy a file**

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

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

Arguments

- **from, to**: Character vectors, containing file names or paths. `to` can alternatively be the path to a single existing directory.
- **symlink**: Logical indicating whether to use symlink (instead of hardlink). Default `FALSE`.
- **overwrite**: Logical. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.
- **verbose**: Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messing, 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 to minimal.

Value

This function is called for its side effects, which will be a `file.link` if 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
}
listNamed

Create a list with names from object names

Description

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

Usage

```r
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

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

loadFile

Load a file from the cache

Description

Load a file from the cache

Usage

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

Arguments

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

Value

the object loaded from file

mergeCache | *Merge two cache repositories together*

Description

Usage

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

```
## S4 method for signature 'ANY'
mergeCache(
  cacheTo,
  cacheFrom,
  drvTo = getDrv(getOption("reproducible.drv", NULL)),
  drvFrom = getDrv(getOption("reproducible.drv", NULL)),
  connTo = NULL,
  connFrom = NULL,
  verbose = getOption("reproducible.verbose")
)
```

Arguments

- **cacheTo** The cache repository (character string of the file path) that will become larger, i.e., merge into this
- **cacheFrom** The cache repository (character string of the file path) from which all objects will be taken and copied from
- **drvTo** The database driver for the cacheTo.
messageDF

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>drvFrom</td>
<td>The database driver for the cacheFrom</td>
</tr>
<tr>
<td>connTo</td>
<td>The connection for the cacheTo. If not provided, then a new one will be made</td>
</tr>
<tr>
<td>connFrom</td>
<td>The database for the cacheFrom. If not provided, then a new one will be made</td>
</tr>
<tr>
<td>verbose</td>
<td>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 verbose output.</td>
</tr>
</tbody>
</table>

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

Use message with a consistent use of verbose

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
messageDF

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 crayon
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 to minimal 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**

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

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

```r
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

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

Arguments

- **new**: Either the path of the new cachePath where the cache was moved or copied to, or the new DB Table Name.
- **old**: Optional, if there is only one table in the new cache path. Either the path of the previous cachePath where the cache was moved or copied from, or the old DB Table Name.
- **drv**: if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite.
- **conn**: an optional DBIConnection object, as returned by dbConnect().
- **verbose**: Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., `options('reproducible.verbose' = 0)` to reduce the verbosity.
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::.setDThreads(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 a absolute paths, whereas `normPathRel()` maintains relative paths.

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

Value

Character vector of cleaned up filepaths.

Examples

## normalize file paths
paths <- list("./aaa/zzz",
              "./aaa/zzz/",
              ".//aaa//zzz",
              ".//aaa//zzz/",
              ".\\aaa\\zzz",
              ".\\aaa\\zzz\\",
              ".\\\\aaa\\\\zzz",
              ".\\\\aaa\\\\zzz\\\"
"
checked <- normPath(paths)
length(unique(checked)) # all of the above are equivalent

## check to see if a path exists
tmpdir <- file.path(tempdir(), "example_checkPath")
dir.exists(tmpdir) # FALSE
tryCatch(checkPath(tmpdir, create = FALSE), error = function(e) FALSE) # FALSE

cHECKPath(tmpdir, create = TRUE)
dir.exists(tmpdir) # TRUE
unlink(tmpdir, recursive = TRUE)

---

**objSize**  
*Wrapper around lobstr::obj_size*

**Description**

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

**Usage**

```r
objSize(x, quick = FALSE, ...)
objSizeSession(sumLevel = Inf, enclosingEnvs = TRUE, .prevEnvirs = list())
```

**Arguments**

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

```r
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

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

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

Value

Character string representing the filename.

Author(s)

Eliot McIntire and Alex Chubaty

Examples

```r
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

```r
asPath(obj, nParentDirs = 0)
```

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

## S3 method for class 'null'
```r
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

Description

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

Usage

postProcess(x, ...)

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

postProcess(x, ...)

### Arguments

- **x**: A GIS object of postProcessing, e.g., Spat* or sf*. This can be provided as a `rlang::quosure` or a normal R object.
- **...**: Additional arguments passed to methods. For spatialClasses, these are: `cropTo()`, `fixErrorsIn()`, `projectTo()`, `maskTo()`, `determineFilename()`, and `writeTo()`. Each of these may also pass `...` into other functions, like `writeTo()`. This might include potentially important arguments like `datatype`, `format`. Also passed to `terra::project`, with likely important arguments such as `method = "bilinear"`. See details.

### Value

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

### Post processing sequence

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

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

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

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

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

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

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

<table>
<thead>
<tr>
<th></th>
<th>rasterToMatch</th>
<th>studyArea</th>
<th>Both</th>
</tr>
</thead>
<tbody>
<tr>
<td>extent</td>
<td>Yes</td>
<td>Yes</td>
<td></td>
</tr>
<tr>
<td>resolution</td>
<td>Yes</td>
<td>No</td>
<td></td>
</tr>
</tbody>
</table>
### postProcess

<table>
<thead>
<tr>
<th></th>
<th>Yes</th>
<th>No*</th>
<th>rasterToMatch*</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>projec</strong>tion</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>alignment</td>
<td>Yes</td>
<td>No</td>
<td>rasterToMatch</td>
</tr>
<tr>
<td>mask</td>
<td>No**</td>
<td>Yes</td>
<td>studyArea**</td>
</tr>
</tbody>
</table>

*C*an be overridden with useSAcrs.

**Will mask with NAS from rasterToMatch if maskWithRTM.

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

<table>
<thead>
<tr>
<th></th>
<th>Yes</th>
<th>Yes</th>
<th>rasterToMatch</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>rasterToMatch</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>studyArea</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Both</td>
<td></td>
<td></td>
<td>rasterToMatch*</td>
</tr>
</tbody>
</table>

*Can be overridden with useSAcrs

**See Also**

prepInputs

**Examples**

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

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

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

  # 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)

if (!all(terra::is.valid(studyArea2))) studyArea2 <- terra::makeValid(studyArea2)

tf <- tempfile(fileext = ".tif")
download.file(url = remoteTifUrl, destfile = tf, mode = "wb")

Checksums(dPath, write = TRUE, files = tf)
elevOrig <- terra::rast(tf)
elevForStudy2 <- terra::project(elevOrig, terra::crs(studyArea2), res = 250) |>
   terra::crop(studyArea2) |>
   terra::mask(studyArea2)

isTRUE(all.equal(studyArea, studyArea2))  # Yes!

# sf class
studyAreaSmall <- prepInputs(localFileLuxSm)

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

# SpatVector-- note: doesn't matter what class the "to" object is, only the "from"
studyAreas <- list()
studyAreas[["orig"]]<- prepInputs(localFileLux, fun = "terra::vect")
studyAreas[["reprojected"]]<- projectTo(studyAreas[["orig"]], studyAreaSmall)
studyAreas[["cropped"]]<- suppressWarnings(cropTo(studyAreas[["orig"]], studyAreaSmall))
studyAreas[["masked"]]<- suppressWarnings(maskTo(studyAreas[["orig"]], studyAreaSmall))

if (interactive()) {
  par(mfrow = c(2,2));
  out <- lapply(studyAreas, function(x) terra::plot(x))
}

setwd(od)

---

postProcessTo  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. 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"),
  ...
)
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. Attention. Conflicts may arise when projectTo is a Vector/CRS object with a distinct CRS from to. Because to is used for masking after from is re-projected using projectTo, the extents of to and from may no longer overlap (as in align) perfectly leading to failure during the masking step. We recommend passing a raster templates to projectTo whose extent and CRS are both compatible with the object used later for masking (either to or maskTo).
maskTo Optional Gridded or Vector dataset which, if supplied, will supply the extent with which to mask from. If Gridded, it will mask with the NA values on the maskTo; if Vector, it will mask on the terra::aggregate(maskTo). To omit masking completely, set this to NA. If supplied, this will override to for the masking step. Defaults to NULL, which means use to
writeTo Optional character string of a filename to use writeRaster to save the final object. Default is NULL, which means there is no writeRaster
overwrite Logical. Used if writeTo is not NULL; also if terra determines that the object requires writing to disk during a crop, mask or project call e.g., because it is too large.
postProcessTo

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 to minimal

Arguments passed to `terra::mask` (for `maskTo`), `terra::project` (for `projectTo`) or `terra::writeRaster` (for `writeTo`) and not used for `cropTo`, as well postProcess's `rasterToMatch` and `studyArea` arguments (see below). Commonly used arguments might be method, touches, and datatype. If `filename` is passed, it will be ignored; use `writeTo = .` If `reproducible.gdalwarp = TRUE`, then these will be passed to the `gdal*` functions. See them for details.

needBuffer Logical. Defaults to FALSE, meaning nothing is done out of the ordinary. If TRUE, then a buffer around the `cropTo`, so that if a reprojection has to happen on the `cropTo` prior to using it as a crop layer, then a buffer of 1.5 * res(cropTo) will occur prior, so that no edges are cut off.

isStack, isBrick, isRaster, isSpatRaster Logical. Default NULL. Used to convert from back to these classes prior to writing, if provided.

Details

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

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

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

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

Use Cases

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

<table>
<thead>
<tr>
<th>from</th>
<th>to</th>
<th>from will have:</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gridded</td>
<td>Gridded</td>
<td>the extent, projection, origin, resolution and masking where there are NA from the to</td>
</tr>
<tr>
<td>Gridded</td>
<td>Vector</td>
<td>the projection, origin, and mask from to, and extent will be a round number of pixels that fit within the</td>
</tr>
<tr>
<td>Vector</td>
<td>Vector</td>
<td>the projection, origin, extent and mask from to</td>
</tr>
</tbody>
</table>

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 * res(from)[1]), so that edge cells still have a complete set of neighbours. The second crop is at the end, after projecting and masking. After the projection step, the crop is no longer tight. Under some conditions, masking will effectively mask and crop in one step, but under some conditions, this is not true, and the mask leaves padded NAs out to the extent of the from (as it is after crop, project, mask). Thus the second crop removes all NA cells so they are tight to the mask.

See Also

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

Examples

# prepare dummy data -- 3 SpatRasters, 2 SpatVectors
# need 2 SpatRaster
rf <- system.file("ex/elev.tif", package = "terra")
elev1 <- terra::rast(rf)
#
ras2 <- terra::deepcopy(elev1)
ras2[ras2 > 200 & ras2 < 300] <- NA_integer_
terra::values(elev1) <- rep(1L, terra::ncell(ras2))
#
# a polygon vector
f <- system.file("ex/lux.shp", package = "terra")
vOrig <- terra::vect(f)
v <- vOrig[1:2,]
#
utm <- terra::crs("epsg:23028") # $wkt
vInUTM <- terra::project(vOrig, utm)
vAsRsInLongLat <- terra::rast(vOrig, resolution = 0.008333333)
res100 <- 100
rInUTM <- terra::rast(vInUTM, resolution = res100)
# 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 = vAsRsInLongLat, maskTo = vInUTM)

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

prepInputs

Download and optionally post-process files

Description

Usage

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

**targetFile** Character string giving the filename (without relative or absolute path) to the eventual file (raster, shapefile, csv, etc.) after downloading and extracting from a zip or tar archive. This is the file 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 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 `preProcess()`.

**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 `preProcess()`. If it is NA, then it will not attempt to see it as an archive, even if it has archive-like file extension (e.g., .zip). This may be useful when an R function is expecting an archive directly.

**alsoExtract** Optional character string naming files other than targetFile that must be extracted from the archive. If NULL, the default, then it will extract all files. Other options: "similar" will extract all files with the same filename without file extension as targetFile. NA will extract nothing other than targetFile. A character string of specific file names will cause only those to be extracted. See table in `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.

**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. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.

**purge** Logical or Integer. 0/FALSE (default) keeps existing CHECKSUMS.txt file and prepInputs will write or append to it. 1/TRUE will deleted the entire CHECKSUMS.txt file. Other options, see details.
prepInputs

useCache Passed to Cache in various places. Defaults to option("reproducible.useCache", 2L) in prepInputs, and option("reproducible.useCache", FALSE) if calling any of the inner functions manually. For prepInputs, this means it will use Cache only up to 2 nested levels, which includes preProcess and its nested *Input functions (e.g., cropInputs, projectInputs, maskInputs) are no longer internally cached, as terra processing speeds mean internal caching is more time consuming. We recommend caching the full prepInputs call instead (e.g., prepInputs(...)|> Cache()).

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

verbose Numeric, -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., options('reproducible.verbose' = 0) to reduce to minimal

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.

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. Determine file name determineFilename() via filename2;
6. Optionally, write that file name to disk via writeTo().

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

NOTE: sf objects are still very experimental.

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

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

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

fun

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

fun can be supplied in three ways: a function, a character string (i.e., a function name as a string), or an expression. If a character string or function, 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.
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)
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See Also
postProcessTo(), downloadFile(), extractFromArchive(), postProcess().

Examples

if (requireNamespace("terra", quietly = TRUE) &&
requireNamespace("sf", quietly = TRUE)) {
  library(reproducible)

  # Make a dummy study area map -- user would supply this normally
  coords <- structure(c(-122.9, -116.1, -99.2, -106, -122.9, 59.9, 65.7, 63.6, 54.8, 59.9),
    Dim = c(5L, 2L))
  studyArea <- terra::vect(coords, "polygons")
  terra::crs(studyArea) <- "+proj=longlat +datum=WGS84 +no_defs +ellps=WGS84 +towgs84=0,0,0"

  # Make dummy "large" map that must be cropped to the study area
  outerSA <- terra::buffer(studyArea, 50000)
  terra::crs(outerSA) <- "+proj=longlat +datum=WGS84 +no_defs +ellps=WGS84 +towgs84=0,0,0"
  tf <- normPath(file.path(tempdir2("prepInputsEx"), "prepInputs2.shp"))
  terra::writeVector(outerSA, tf, overwrite = TRUE)

  # run prepInputs -- load file, postProcess it to the studyArea
  studyArea2 <- prepInputs(}
targetFile = tf, to = studyArea,
fun = "terra::vect",
destinationPath = tempdir2()
)
  |> suppressWarnings() # not relevant warning here

# clean up
unlink("CHECKSUMS.txt")

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

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

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

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

# specify targetFile, alsoExtract, and fun, wrap with Cache
ecozoneFilename <- file.path(dPath, "ecozones.shp")
# Note, you don’t need to "alsoExtract" the archive... if the archive is not there, but the
# targetFile is there, it will not redownload the archive.
ecozoneFiles <- c("ecozones.dbf", "ecozones.prj",
                  "ecozones.sbn", "ecozones.sbx", "ecozones.shp", "ecozones.shx")

shpEcozoneSm <- Cache(prepInputs,
                      url = shpUrl,
                      targetFile = reproducible::asPath(ecozoneFilename),
                      alsoExtract = reproducible::asPath(ecozoneFiles),
                      studyArea = studyArea,
                      fun = "terra::vect",
                      destinationPath = dPath,
                      filename2 = "EcozoneFile.shp"
)  # passed to determineFilename

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

## Using quoted dlFun and fun -- this is not intended to be run but used as a template
## prepInputs(..., fun = customFun(x = targetFile), customFun = customFun)
## # or more complex
## test5 <- prepInputs(
##   targetFile = targetFileLuxRDS,
##   dlFun = getDataFn(name = "GADM", country = "LUX", level = 0) # preProcess keeps file from this!
##   ,
##   fun = {
##     out <- readRDS(targetFile)
##     sf::st_as_sf(out)}
##   )

preProcessParams
Download, Checksum, Extract files
Description

This does downloading (via downloadFile), checksumming (Checksums), and extracting from archives (extractFromArchive), plus cleaning up of input arguments (e.g., paths, function names).

This is the first stage of three used in prepInputs.

Usage

preProcessParams(n = NULL)

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),
... )

Arguments

n

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

targetFile

Character string giving the filename (without relative or absolute path) to the eventual file (raster, shapefile, csv, etc.) after downloading and extracting from a zip or tar archive. This is the file 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 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 preProcess().

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 preProcess(). If it is NA, then it will not attempt to see it as an archive, even if it has archive-like file extension (e.g., .zip). This may be useful when an R function is expecting an archive directly.
alsoExtract  Optional character string naming files other than targetFile that must be extracted from the archive. If NULL, the default, then it will extract all files. Other options: "similar" will extract all files with the same filename without file extension as targetFile. NA will extract nothing other than targetFile. A character string of specific file names will cause only those to be extracted. See table in `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. Should downloading and all the other actions occur even if they pass the checksums or the files are all there.  

purge  Logical or Integer. 0/FALSE (default) keeps existing CHECKSUMS.txt file and `prepInputs` will write or append to it. 1/TRUE will deleted the entire CHECKSUMS.txt file. Other options, see details.  

verbose  Numeric. -1 silent (where possible), 0 being very quiet, 1 showing more messaging, 2 being more messaging, etc. Default is 1. Above 3 will output much more information about the internals of Caching, which may help diagnose Caching challenges. Can set globally with an option, e.g., `options(‘reproducible.verbose’ = 0)` to reduce to minimal.  

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

...  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

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**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**

_Remap file names_

---

**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, ...)`
reproducibleOptions

Arguments

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

Description

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

Usage

reproducibleOptions()

Details

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

The following options are likely of interest to most users:

ask Default: TRUE. Used in clearCache() and keepCache().
cachePath Default: .reproducibleTempCacheDir. Used in Cache() and many others. The default path for repositories if not passed as an argument.
cacheSaveFormat Default: "rds". What save format to use; currently, "qs" or "rds".
cacheSpeed Default "slow". One of "slow" or "fast" (1 or 2). "slow" uses digest::digest internally, which is transferable across operating systems, but much slower than digest::digest(algo = "spooky"). So, if all caching is happening on a single machine, "fast" would be a good setting.
conn Default: NULL. Sets a specific connection to a database, e.g., dbConnect(drv = RSQLite::SQLite()) or dbConnect(drv = RPostgres::Postgres()). For remote database servers, setting one connection may be far faster than using drv which must make a new connection every time.
destinationPath Default: NULL. Used in prepInputs() and preProcess(). Can be set globally here.
drv Default: RSQLite::SQLite(). Sets the default driver for the backend database system. Only tested with RSQLite::SQLite() and RPostgres::Postgres().
futurePlan Default: FALSE. On Linux OSes, Cache and cloudCache have some functionality that uses the future package. Default is to not use these, as they are experimental. They may, however, be very effective in speeding up some things, specifically, uploading cached elements via googledrive in cloudCache.
gdalwarp Default: FALSE. Experimental. During postProcessTo the standard approach is to use terra functions directly, with several strategic uses of sf. However, in the special case when from is a SpatRaster or Raster, maskTo is a SpatVector or SFC_POLYGON and projectTo is a SpatRaster or Raster, setting this option to TRUE will use sf::gdal_utils("warp"). In many test cases, this is much faster than the terra sequence. The resulting SpatRaster is not identical, but it is very similar.

gdalwarpThreads Default: 2. This will set -wo NUM_THREADS= to this number. Default is now 2, meaning gdalwarp will use 2 threads with gdalProject. To turn off threading, set to 0, 1 or NA.

inputPaths Default: NULL. Used in prepInputs() and preProcess(). If set to a path, this will cause these functions to save their downloaded and preprocessed file to this location, with a hardlink (via file.link) to the file created in the destinationPath. This can be used so that individual projects that use common data sets can maintain modularity (by placing downloaded objects in their destinationPath, but also minimize re-downloading the same (perhaps large) file over and over for each project. Because the files are hardlinks, there is no extra space taken up by the apparently duplicated files.

inputPathsRecursive Default: FALSE. Used in prepInputs() and preProcess(). Should the reproducible.inputPaths be searched recursively for existence of a file?

memoisePersist Default: FALSE. Used in Cache(). Should the memoised copy of the Cache objects persist even if reproducible reloads e.g., via devtools::load_all? This is mostly useful for developers of reproducible. If TRUE, a object named paste0(".reproducibleMemoise_", cachePath) will be placed in the .GlobalEnv, i.e., one for each cachePath.

nThreads Default: 1. The number of threads to use for reading/writing cache files.

objSize Default: TRUE. Logical. If TRUE, then object sizes will be included in the cache database. Simplying calculating object size of large objects can be time consuming, so setting this to FALSE will make caching up to 10% faster, depending on the objects.

overwrite Default: FALSE. Used in prepInputs(), preProcess(), downloadFile(), and postProcess(). 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.

timeout Default 1200. Used in preProcess when downloading occurs. If a user has R.utils package installed, R.utils::withTimeout( , timeout = getOption("reproducible.timeout")) will be wrapped around the download so that it will timeout (and error) after this many seconds.

useCache Default: TRUE. Used in Cache(). If FALSE, then the entire Cache machinery is skipped and the functions are run as if there was no Cache occurring. Can also take 2 other values: 'overwrite' and 'devMode'. 'overwrite' will cause no recovery of objects from the cache repository, only new ones will be created. If the hash is identical to a previous one, then this will overwrite the previous one. 'devMode' will function as normally Cache except it will use
the userTags to determine if a previous function has been run. If the userTags are identical, but the digest value is different, the old value will be deleted from the cache repository and this new value will be added. This addresses a common situation during the development stage: functions are changing frequently, so any entry in the cache repository will be stale following changes to functions, i.e., they will likely never be relevant again. This will therefore keep the cache repository clean of stale objects. If there is ambiguity in the userTags, i.e., they do not uniquely identify a single entry in the cachePath, then this option will default back to the non-dev-mode behaviour to avoid deleting objects. This, therefore, is most useful if the user is using unique values for userTags.

useCloud Default FALSE. Passed to Cache.

useDBI Default: TRUE if DBI is available. Default value can be overridden by setting environment variable R_REPRODUCIBLE_USE_DBI. As of version 0.3, the backend is now DBI instead of archivist.

useGdown Default: FALSE. If a user provides a Google Drive url to preProcess/prepInputs, reproducible will use the googledrive package. This works reliably in most cases. However, for large files on unstable internet connections, it will stall and stop the download with no error. If a user is finding this behaviour, they can install the gdown package, making sure it is available on the PATH. This call to gdown will only work for files that do not need authentication. If authentication is needed, 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.

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


---

**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**

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

exprBetween  Another expression that should be run after a failed attempt of the expr. This should return a named list, where the names indicate the object names to update in the main expr, 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 expr) will be updated prior to running the expr again. For backwards compatibility, this still works).

messageFn  A function for messaging to console. Defaults to message

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

Arguments

  cachePath       A repository used for storing cached objects. This is optional if Cache is used inside a SpaDES module.
  drv             if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite
  conn            an optional DBIConnection object, as returned by dbConnect().
  obj             The R object to save to the cache
userTags  A character vector with descriptions of the Cache function call. These will be added to the Cache so that this entry in the Cache can be found using userTags e.g., via showCache().
cacheId  The hash string representing the result of .robustDigest
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 this.

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

searchFull  

Search up the full scope for functions

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

Usage
searchFull(env = parent.frame(), simplify = TRUE)
searchFullEx()

Arguments
env  The environment to start searching at. Default is calling environment, i.e., parent.frame()
simplify  Logical. Should the output be simplified to character, if possible (usually it is not possible because environments don’t always coerce correctly)

Details
searchFullEx can be used to show an example of the use of searchFull.
Value
A list of environments that is the actual search path, unlike search() which only prints from .GlobalEnv up to base through user attached packages.

See Also
base::search()

Examples

```r
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

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

Arguments

```r
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
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,
showCache

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

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

keepCache

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

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

\texttt{x} \quad A \texttt{simList} or a directory containing a valid Cache repository. Note: For compatibility with Cache argument, \texttt{cachePath} can also be used instead of \texttt{x}, though \texttt{x} will take precedence.

\texttt{userTags} \quad Character vector. If used, this will be used in place of the \texttt{after} and \texttt{before}. Specifying one or more \texttt{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 \texttt{showCache()}, i.e., \texttt{artifact}, \texttt{tagValue} or \texttt{tagName}. Also, if \texttt{length(userTags) > 1}, then matching is by and. For or matching, use | in a single character string. See examples.

\texttt{after} \quad A time (POSIX, character understandable by \texttt{data.table}). Objects cached after this time will be shown or deleted.

\texttt{before} \quad A time (POSIX, character understandable by \texttt{data.table}). Objects cached before this time will be shown or deleted.

\texttt{fun} \quad An optional character vector describing the function name to extract. Only functions with this/these functions will be returned.

\texttt{cacheId} \quad An optional character vector describing the cacheIds to extract. Only entries with this/these cacheIds will be returned. If \texttt{useDBI(FALSE)}, this will also be dramatically faster than using \texttt{userTags}, for a large cache.

\texttt{ask} \quad Logical. If \texttt{FALSE}, then it will not ask to confirm deletions using \texttt{clearCache} or \texttt{keepCache}. Default is \texttt{TRUE}.

\texttt{useCloud} \quad Logical. If \texttt{TRUE}, then every object that is deleted locally will also be deleted in the \texttt{cloudFolderID}, if it is non-NULL.

\texttt{cloudFolderID} \quad A googledrive dribble of a folder, e.g., using \texttt{drive_mkdir()}. If left as \texttt{NULL}, the function will create a cloud folder with name from last two folder levels of the \texttt{cachePath} path: \texttt{paste0(basename(dirname(cachePath)), ",", basename(cachePath))}. This \texttt{cloudFolderID} will be added to \texttt{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.

\texttt{drv} \quad if using a database backend, \texttt{drv} must be an object that inherits from \texttt{DBIDriver}, e.g., from package RSQLite, e.g., \texttt{SQLite}.

\texttt{conn} \quad an optional \texttt{DBIConnection} object, as returned by \texttt{dbConnect()}.

\texttt{verbose} \quad 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., \texttt{options('reproducible.verbose' = 0)} to reduce to minimal...

\ldots

Other arguments. Can be in the form of \texttt{tagKey = tagValue}, such as, \texttt{class = "numeric"} to find all entries that are numerics in the cache. Note: the special cases of \texttt{cacheId} and \texttt{fun} have their own named arguments in these functions. Also can be \texttt{regexp = xx}, where \texttt{xx} is \texttt{TRUE} if the user is passing a regular expression. Otherwise, \texttt{userTags} will need to be exact matches. Default is missing, which is the same as \texttt{TRUE}. If there are errors due to regular expression...
showCache

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.

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”.

cc(secs) is just a shortcut for clearCache(repo = currentRepo, after = secs).

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 at cachePath of the sim object, if sim is provided, or 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
studyAreaName <- 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
cacheAfter <- showCache(tmpDir, userTags = c("runif")) # Only the small one is left

clearCache(tmpDir, ask = FALSE)

cache1 <- showCache(tmpDir, userTags = c("runif"))
toRemove <- cache1[tagKey == "object.size"[as.numeric(tagValue) > 700]]$cacheId
clearCache(tmpDir, userTags = toRemove, ask = FALSE)

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**

```r
studyAreaName(studyArea, ...)  
```

## S4 method for signature 'character'

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

## Arguments

- **studyArea**: Spatial object.
- **...**: Other arguments (not currently used)

## Value

A character string using the \texttt{.robustDigest} of the \texttt{studyArea}. This is only intended for use with spatial objects.

## Examples

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

---

## Description

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

## Usage

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

## Arguments

- **sub**: Character string, length 1. Can be a result of `file.path("smth", "smth2")` 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 `getOption("reproducible.tempPath")`.
- **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

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

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 file.path("smth", "smth2")
  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. De-
 faults to getOption("reproducible.tempPath").

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

The known path for unrar or 7z

Description

The known path for unrar or 7z

Usage

.unrarPath

Format

An object of class NULL of length 0.
writeFuture

Write to cache repository, using future::future

Description

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

Usage

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

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>written</td>
<td>Integer. If zero or positive then it needs to be written still. Should be 0 to start.</td>
</tr>
<tr>
<td>outputToSave</td>
<td>The R object to save to repository</td>
</tr>
<tr>
<td>cachePath</td>
<td>The file path of the repository</td>
</tr>
<tr>
<td>userTags</td>
<td>Character string of tags to attach to this outputToSave in the CacheRepo</td>
</tr>
<tr>
<td>drv</td>
<td>if using a database backend, drv must be an object that inherits from DBIDriver e.g., from package RSQLite, e.g., SQLite</td>
</tr>
<tr>
<td>conn</td>
<td>an optional DBIConnection object, as returned by dbConnect().</td>
</tr>
<tr>
<td>cacheId</td>
<td>Character string. If passed, this will override the calculated hash of the inputs, and return the result from this cacheld 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.</td>
</tr>
<tr>
<td>linkToCacheId</td>
<td>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.</td>
</tr>
</tbody>
</table>

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.
Index

* datasets
  * unrarPath, 102
  .debugCache, 5
  .digest(), 32
  .file.move, 6
  .isMemoised, 6
  .messageFunctionFn (messageDF), 62
  .prefix, 7
  .prepareFileBackedRaster, 8
  .removeCacheAtts, 9
  .requireNamespace, 9
  .robustDigest(), 22, 23, 38
  .setSubAttrInList, 10
  .suffix (.prefix), 7
  .unrarPath (unrarPath), 102
  .unwrap (.wrap), 10
  .wrap, 10

  asPath (Path-class), 70
  assessDataType, 13

  base::basename(), 16
  base::search(), 95
  basename2, 16

  Cache, 16
  Cache(), 4, 18, 26, 46, 49, 80, 81, 87, 89–92, 99
  CacheDBFile (createCache), 40
  CacheDBTableName (createCache), 40
  CacheDigest, 25
  CacheDigest(), 23, 92
  CacheGeo, 26
  CacheIsACache (createCache), 40
  CacheStorageDir (createCache), 40
  CacheStorageDir(), 23
  CacheStoredFile (createCache), 40
  cc (showCache), 96
  checkAndMakeCloudFolderID, 29
  checkPath, 30
  checkPath, character, logical-method
    (checkPath), 30
  checkPath, character, missing-method
    (checkPath), 30
  checkPath, missing, ANY-method
    (checkPath), 30
  checkPath, NULL, ANY-method (checkPath), 30
  checkRelative, 31
  Checksums, 32
  Checksums(), 46, 80, 87
  Checksums, character, logical-method
    (Checksums), 32
  Checksums, character, missing-method
    (Checksums), 32
  clearCache (showCache), 96
  clearCache(), 19, 23, 89
  clearCache, ANY-method (showCache), 96
  cloudDownload, 34
  compareNA, 35
  convertPaths, 36
  convertRasterPaths (convertPaths), 36
  Copy, 37
  Copy, ANY-method (Copy), 37
  Copy, data.frame-method (Copy), 37
  Copy, data.table-method (Copy), 37
  Copy, list-method (Copy), 37
  Copy, refClass-method (Copy), 37
  copyFile (copySingleFile), 39
  copySingleFile, 39
  createCache, 40
  cropInputs(), 49, 74, 81, 87
  cropTo (postProcessTo), 74
  cropTo(), 71, 72, 77, 78, 82

  dataType2 (minFn), 64
  determineFilename, 44
  determineFilename(), 44, 72, 82
  digest::digest(), 19, 33
  dir.create(), 31
downloadFile, 45
downloadFile(), 83, 90
downloadRemote, 47

extractFromArchive, 49
extractFromArchive(), 83
extractFromCache (createCache), 40

fastMask, 50
file.copy(), 59
file.exists(), 31
file.link(), 59
file.symlink(), 59
Filenames, 51
Filenames,ANY-method (Filenames), 51
Filenames, data.table-method (Filenames), 51
Filenames, environment-method (Filenames), 51
Filenames, list-method (Filenames), 51
Filenames, Path-method (Filenames), 51
fixErrorsIn, 52
fixErrorsIn(), 71, 72, 78

gdalMask (gdalProject), 53
gdalMask(), 54, 78
gdalProject, 53
gdalProject(), 78
gdalResample (gdalProject), 53
gdalResample(), 54, 78
getRelative, 55

internetExists, 56
isUpdated, 57

keepCache (showCache), 96
keepCache(), 23, 89
keepCache, ANY-method (showCache), 96
keepOrigGeom, 57

linkOrCopy, 58
listNamed, 60
loadFile, 60
loadFile(), 43
loadFromCache (createCache), 40

makeRelative (getRelative), 55
maskTo (postProcessTo), 74
maskTo(), 50, 54, 71, 72, 77, 78, 82

maxFn (minFn), 64
mergeCache, 61
mergeCache(), 99
mergeCache, ANY-method (mergeCache), 61
messageCache (messageDF), 62
messageColoured (messageDF), 62
messageDF, 62
messagePrepInputs (messageDF), 62
messagePreProcess (messageDF), 62
messageQuestion (messageDF), 62
minFn, 64
movedCache, 65
movedCache(), 23

nlayers2 (minFn), 64
normPath, 66
normPath(), 31
normPath, character-method (normPath), 66
normPath, list-method (normPath), 66
normPath, logical-method (normPath), 66
normPath, missing-method (normPath), 66
normPath, NULL-method (normPath), 66
normPathRel (normPath), 66

objSize, 68
objSizeSession (objSize), 68
options(), 4

paddedFloatToChar, 69
Path-class, 70
postProcess, 71
postProcess(), 49, 74, 77, 78, 81–83, 87, 90
postProcessTerra (postProcessTo), 74
postProcessTo, 74
postProcessTo(), 54, 57, 72, 81–83
prepInputs, 79
prepInputs(), 4, 45, 89, 90
preProcess (preProcessParams), 85
preProcess(), 46, 48, 80, 81, 86, 87, 89, 90
preProcessParams, 85
projectTo (postProcessTo), 74
projectTo(), 71, 72, 77, 78, 82

rasterRead, 88
remapFilenames, 88
reproducible (reproducible-package), 4
reproducible-package, 4
reproducibleOptions, 89
reproducibleOptions(), 4
retry, 92
rmFromCache (createCache), 40rmFromCache(), 23

saveToCache, 93
searchFull, 94
searchFullEx (searchFull), 94
set.randomseed, 95
showCache, 96
showCache(), 18, 23, 94
showCache, ANY-method (showCache), 96
studyAreaName, 100
studyAreaName, ANY-method
(studyAreaName), 100
studyAreaName, character-method
(studyAreaName), 100
suffix (.prefix), 7

tempdir2, 101, 102
tempfile2, 101, 102

unrarPath, 102
untar(), 81
unzip(), 81
urlExists (internetExists), 56
utils::download.file(), 45, 81
utils::write.table(), 33

values2 (minFn), 64

writeFuture, 103
writeTo (postProcessTo), 74
writeTo(), 44, 71, 72, 77, 78, 82