

Package: indexr (via r-universe)

February 18, 2025

Title A Thoughtful Saver of Results

Version 0.2.2

Date 2025-02-14

Description Helps with the thoughtful saving, reading, and management of result files (using 'rds' files). The core functions take a list of parameters that are used to generate a unique hash to save results under. Then, the same parameter list can be used to read those results back in. This is helpful to avoid clunky file naming when running a large number of simulations. Additionally, helper functions are available for compiling a flat file of parameters of saved results, monitoring result usage, and cleaning up unwanted or unused results. For more information, visit the 'indexr' homepage [<https://lharris421.github.io/indexr/>](https://lharris421.github.io/indexr/).

BugReports <https://github.com/lharris421/indexr/issues>

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URL <https://lharris421.github.io/indexr/>,
<https://github.com/lharris421/indexr>

Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.3.2

Imports stringr, readr, dplyr, digest, glue, methods

Suggests testthat (>= 3.0.0)

Config/pak/sysreqs libicu-dev libx11-dev

Repository <https://lharris421.r-universe.dev>

RemoteUrl <https://github.com/lharris421/indexr>

RemoteRef HEAD

RemoteSha 15aedbca5a6d77d52aed160c3b52039e5d65dd34

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check_hash_existence *Check for the Existence of Results Under a Set of Parameters*

Description

This function checks for the existence of results saved under specified parameter list in RDS files (saved with indexr) within a given folder.

Usage

```
check_hash_existence(
  folder,
  parameters_list,
  halt = FALSE,
  hash_includes_timestamp = FALSE,
  ignore_na = TRUE,
  alphabetical_order = TRUE,
  algo = "xxhash64",
  ignore_script_name = FALSE
)
```

Arguments

folder	A string specifying the directory containing the RDS files.
parameters_list	A list of parameters for which a corresponding hash named file is checked.
halt	Logical; if TRUE, the function stops execution if an existing file is found. This may be useful as a check before running a simulation.
hash_includes_timestamp	Logical; if TRUE, timestamps are included in the hash generation process.
ignore_na	Logical; if TRUE, NA values are ignored during hash generation.

alphabetical_order
 Logical; if TRUE, parameters are sorted alphabetically before hash generation.

algo
 Character string specifying the hashing algorithm to use. Default is "xxhash64".
 See `?digest`

ignore_script_name
 Logical. If TRUE, the script name is ignored during hash generation.

Value

A logical of whether or not a file exists, unless `halt = TRUE` and a file is found, then an error is thrown.

Examples

```
## Setup
tmp_dir <- file.path(tempdir(), "example")
dir.create(tmp_dir)

## Save an object
parameters_list <- list(example = "check_hash_existence")
save_objects(folder = tmp_dir, results = 1, parameters_list = parameters_list)

## Check that an object under specified parameters is saved
check_hash_existence(folder = tmp_dir, parameters_list)

## Cleanup
unlink(tmp_dir, recursive = TRUE)
```

cleanup_from_hash_table

Remove Files Based on Hash Table

Description

Allows the user to leverage the `generate_hash` function to generate a table that is subsequently used to remove indicated results.

Usage

```
cleanup_from_hash_table(
  folder,
  hash_table,
  mode = c("manual", "all"),
  column = NULL,
  request_confirmation = TRUE
)
```

Arguments

folder	A string specifying the directory containing the RDS files.
hash_table	A data.frame from create_hash_table.
mode	A character string. When mode = "manual" (default) the function expects that the user will add a column to a hash table that indicated which files to delete. When mode = "all", any results in the hash table will be removed.
column	A character string indicating the logical column in hash_table specifying which files to delete.
request_confirmation	Logical, if TRUE will request user input before proceeding to delete files.

Details

There are a few ways to use this. When mode = "manual" (default) the function expects that the user will add a column to a hash table that indicated which files to delete. When mode = "all", any results in the hash table will be removed. This is generally only used when a filter_list is passed to create_hash_table.

Value

Nothing, this function is called for its side effects.

See Also

[create_hash_table\(\)](#)

Examples

```
## Setup
tmp_dir <- file.path(tempdir(), "example")
dir.create(tmp_dir)

## Save example objects
parameters_list1 <- list(example = "tagging1")
parameters_list2 <- list(example = "tagging2")
save_objects(folder = tmp_dir, results = 1, parameters_list = parameters_list1)
save_objects(folder = tmp_dir, results = 2, parameters_list = parameters_list2)

## See the files saved
list.files(tmp_dir)

## Create hash table (flat file of result parameters)
hash_table <- create_hash_table(folder = tmp_dir)

## Delete "all" files based on hash table, without confirmation
cleanup_from_hash_table(
  folder = tmp_dir, hash_table = hash_table, mode = "all", request_confirmation = FALSE
)

## See the files have been deleted
```

```
list.files(tmp_dir)

## Cleanup
unlink(tmp_dir, recursive = TRUE)
```

compress_incremental *Combine Results Saved by save_objects with incremental=TRUE*

Description

This function is only intended to be used after `save_objects` with `incremental=TRUE`. In this case, `save_objects` with save results under temporary hashes in a folder with the hash corresponding to the parameters. `compress_incremental` then combines the results and saves them under the corresponding hash and deletes the old directory with the temporary results.

Usage

```
compress_incremental(
  folder,
  parameters_list,
  hash_includes_timestamp = FALSE,
  ignore_na = TRUE,
  alphabetical_order = TRUE,
  algo = "xxhash64",
  ignore_script_name = FALSE,
  remove_folder = TRUE
)
```

Arguments

<code>folder</code>	Character string specifying the path to the directory where the temporary folder was saved (should be the same as supplied to <code>save_objects</code>).
<code>parameters_list</code>	The named list of arguments used with <code>save_objects</code> .
<code>hash_includes_timestamp</code>	Logical. If TRUE, the timestamp is included in the hash generation.
<code>ignore_na</code>	Logical. If TRUE, NA values in <code>parameters_list</code> are ignored during hash generation.
<code>alphabetical_order</code>	Logical. If TRUE, the names in <code>parameters_list</code> are sorted alphabetically before hash generation.
<code>algo</code>	Character string specifying the hashing algorithm to use. Default is "xxhash64". See <code>?digest</code>
<code>ignore_script_name</code>	Logical. If TRUE, the script name is ignored during hash generation.
<code>remove_folder</code>	Logical. If TRUE, the folder and the temporary results files will be discarded after the combined results are saved.

Details

If the individual results can be put into a `data.frame` they will be, otherwise they will be stored as a list.

Value

No return value. This function is called for its side effects.

See Also

[save_objects\(\)](#)

Examples

```
## Save results incrementally
params <- list(a = "1", b = "2")

tmp_dir <- file.path(tempdir(), "example")
dir.create(tmp_dir)
for (i in 1:10) {
  save_objects(tmp_dir, data.frame(idx = i, val = rnorm(1)), params, incremental = TRUE)
}

## See contents of tmp directory for incremental file
list.files(file.path(tmp_dir, generate_hash(params)))

## Compress results into a single file
compress_incremental(tmp_dir, params)
list.files(tmp_dir)

## Read in compressed file and view results
read_objects(tmp_dir, params)

## Cleanup
unlink(tmp_dir, recursive = TRUE)
```

create_hash_table

Create a Table of the Parameters for Saved Results from RDS Files

Description

Reads in all the parameter files for a give folder, flattens nested lists, and then combines the parameters into a data frame. Each row in the resulting data frame represents the arguments used for one RDS file, identified by its hash. Optionally, the function can filter the data frame based on specified criteria and save it to a file.

Usage

```
create_hash_table(folder, save_path = NULL, filter_list = NULL)
```

Arguments

folder	A string specifying the directory containing the RDS files.
save_path	An optional string specifying the path to save the resulting hash table as a CSV file. If NULL, the hash table is not saved.
filter_list	An optional list of filters to apply to the hash table. Each element of the list should be named according to a column in the hash table and contain the value to filter for in that column.

Details

Saving the hash table can be helpful for the manipulation of parameters (see ?update_hash_table) or for removal of unwanted results (see ?cleanup_from_hash_table).

Value

A data frame where each row corresponds to an parameters_list from an RDS file, with an additional column for the hash of each set of arguments.

Examples

```
## Setup
tmp_dir <- file.path(tempdir(), "example")
dir.create(tmp_dir)

## Save objects
obj1 <- rnorm(1000)
obj2 <- data.frame(
  x = runif(100),
  y = "something",
  z = rep(c(TRUE, FALSE), 50)
)
obj3 <- list(obj1, obj2)

params1 <- list(
  distribution = "normal",
  other_params = list(param1 = TRUE, param2 = 1, param3 = NA)
)
params2 <- list(
  distribution = "uniform",
  other_params = list(param1 = FALSE, param2 = 2, param3 = "1", param4 = 4)
)
params3 <- list(
  distribution = "composite",
  other_params = list(param1 = TRUE, param2 = 3, param3 = 1)
)

save_objects(tmp_dir, obj1, params1)
save_objects(tmp_dir, obj2, params2)
save_objects(tmp_dir, obj3, params3)

## Create hash table (and save it)
```

```

create_hash_table(tmp_dir, save_path = file.path(tmp_dir, "hash_table.csv"))

## Cleanup
unlink(tmp_dir, recursive = TRUE)

```

generate_hash

Generate a Consistent Hash for an Argument List

Description

This function generates a hash value for a given list of arguments. It is designed to produce a consistent hash by optionally removing NA values, ordering arguments alphabetically, handling timestamp inclusion, etc.

Usage

```

generate_hash(
  parameters_list,
  hash_includes_timestamp = FALSE,
  ignore_na = TRUE,
  alphabetical_order = TRUE,
  algo = "xxhash64",
  ignore_script_name = FALSE
)

```

Arguments

parameters_list	A named list of arguments for which the hash will be generated. Each element in the list should correspond to a parameter.
hash_includes_timestamp	Logical; if FALSE, any timestamp included in parameters_list will be removed before hash generation. If TRUE, the timestamp will be included in the hash calculation.
ignore_na	Logical; if TRUE, any NA values in parameters_list will be removed before hash generation.
alphabetical_order	Logical; if TRUE, the arguments in parameters_list will be sorted alphabetically by their names before hash generation.
algo	The hash algorithm to use (See ?digest)
ignore_script_name	Logical. If TRUE, the script name is ignored during hash generation.

Value

A character string representing the hash value of the provided argument list.

Examples

```
args <- list(param1 = "value1", param2 = 100, param3 = NA)
generate_hash(args)
```

read_objects	<i>Read Objects Based on Parameter List</i>
--------------	---

Description

Reads R objects from specified folders based on a generated hash of the provided `parameters_list`.

Usage

```
read_objects(
  folders,
  parameters_list,
  hash_includes_timestamp = FALSE,
  ignore_script_name = FALSE,
  ignore_na = TRUE,
  alphabetical_order = TRUE,
  algo = "xxhash64",
  print_hash = FALSE,
  tagging_file_name = "indexr_tagging.txt",
  silent = FALSE
)
```

Arguments

folders	Character vector specifying the paths to directories containing the saved objects. The function will check each folder in order to find the file.
parameters_list	A named list of arguments used to generate a unique hash for the file.
hash_includes_timestamp	Logical. If TRUE, the timestamp is included in the hash generation.
ignore_script_name	Logical. If TRUE, the script name is ignored during hash generation.
ignore_na	Logical. If TRUE, NA values in <code>parameters_list</code> are ignored during hash generation.
alphabetical_order	Logical. If TRUE, the names in <code>parameters_list</code> are sorted alphabetically before hash generation.
algo	Character string specifying the hashing algorithm to use. Default is "xxhash64".
print_hash	Logical. If TRUE, prints the generated hash. This is helpful for debugging.
tagging_file_name	Character string of a txt file that is being used for tagging results. See <code>?start_tagging</code> .

`silent` Logical. If TRUE, no check is done that pairs of results files (parameters and associated results) is done. This check is not necessary, but done by default to keep the user aware of a scenario that usually results from manual file manipulation.

Details

This function attempts to read an R object from files located in one of the specified folders. The file name is based on the hash of the provided arguments. If the object is successfully read and a tagging file exists and is specified, the function appends the hash and the current timestamp to the tagging file in the folder where the file was found.

Value

The data stored in the file retrieved, typically the results. Returns NULL if the file is not found in any of the specified folders.

See Also

[save_objects\(\)](#)

Examples

```
## Setup
tmp_dir <- file.path(tempdir(), "example")
dir.create(tmp_dir)

## Example using parameter list to run simulation and save results
parameters_list <- list(
  iterations = 1000,
  x_dist = "rnorm",
  x_dist_options = list(n = 10, mean = 1, sd = 2),
  error_dist = "rnorm",
  error_dist_options = list(n = 10, mean = 0, sd = 1),
  beta0 = 1,
  beta1 = 1
)

betas <- numeric(parameters_list$iterations)
for (i in 1:parameters_list$iterations) {
  x <- do.call(parameters_list$x_dist, parameters_list$x_dist_options)
  err <- do.call(parameters_list$error_dist, parameters_list$error_dist_options)
  y <- parameters_list$beta0 + parameters_list$beta1*x + err
  betas[i] <- coef(lm(y ~ x))["x"]
}

save_objects(folder = tmp_dir, results = betas, parameters_list = parameters_list)

## Read back in (consider clearing environment before running)
## Re-setup
tmp_dir <- file.path(tempdir(), "example")

parameters_list <- list(
```

```

iterations = 1000,
x_dist = "rnorm",
x_dist_options = list(n = 10, mean = 1, sd = 2),
error_dist = "rnorm",
error_dist_options = list(n = 10, mean = 0, sd = 1),
beta0 = 1,
beta1 = 1
)

betas <- read_objects(folder = tmp_dir, parameters_list = parameters_list)

## Cleanup
unlink(tmp_dir, recursive = TRUE)

```

rehash	<i>Rehash RDS Files in a Directory</i>
--------	--

Description

This function processes all RDS files in a specified directory, generating new hashes for each file's `args_list` and renaming the files accordingly. It's useful when changing the hash generation algorithm or parameters (if the parameters are manually changed for some reason).

Usage

```

rehash(
  folder,
  hash_includes_timestamp = FALSE,
  ignore_na = TRUE,
  alphabetical_order = TRUE,
  algo = "xxhash64"
)

```

Arguments

<code>folder</code>	A string specifying the directory containing the RDS files to be rehashed.
<code>hash_includes_timestamp</code>	Logical; if TRUE, includes timestamps in the hash generation.
<code>ignore_na</code>	Logical; if TRUE, NA values are ignored during hash generation.
<code>alphabetical_order</code>	Logical; if TRUE, parameters are sorted alphabetically before hash generation.
<code>algo</code>	The (potentially new) hash algorithm to use (see <code>?digest</code>)

Value

The function does not return a value but renames the RDS files in the specified directory based on new hashes.

Examples

```
## Setup
tmp_dir <- file.path(tempdir(), "example")
dir.create(tmp_dir)

# Save example objects
obj1 <- rnorm(1000)
obj2 <- data.frame(
  x = runif(100),
  y = "something",
  z = rep(c(TRUE, FALSE), 50)
)
obj3 <- list(obj1, obj2)

params1 <- list(
  distribution = "normal",
  other_params = list(param1 = TRUE, param2 = 1, param3 = NA)
)
params2 <- list(
  distribution = "uniform",
  other_params = list(param1 = FALSE, param2 = 2, param3 = "1", param4 = 4)
)
params3 <- list(
  distribution = "composite",
  other_params = list(param1 = TRUE, param2 = 3, param3 = 1)
)

save_objects(tmp_dir, obj1, params1)
save_objects(tmp_dir, obj2, params2)
save_objects(tmp_dir, obj3, params3)

## See current file names
list.files(tmp_dir)

## Rehash with new algo
rehash(tmp_dir, algo = "xxhash32")

## Observe new file names
list.files(tmp_dir)

## Cleanup
unlink(tmp_dir, recursive = TRUE)
```

save_objects

*Save Simulation Results with Names as Hashes from the Parameters
that Generated Them*

Description

Saves RDS files to a specified folder with a name that is a hash generated from a list of parameters used for the simulation. There are a number of options that control the behavior, however, the

default functionality likely covers 99% of use cases.

Usage

```
save_objects(
  folder,
  results,
  parameters_list = NULL,
  ignore_na = TRUE,
  alphabetical_order = TRUE,
  overwrite = FALSE,
  include_timestamp = TRUE,
  hash_includes_timestamp = FALSE,
  algo = "xxhash64",
  get_script_name = TRUE,
  ignore_script_name = FALSE,
  incremental = FALSE,
  silent = FALSE
)
```

Arguments

folder	Character string specifying the path to the directory where the objects will be saved.
results	The R object or list of objects to be saved.
parameters_list	A named list of arguments used to generate a unique hash for the file.
ignore_na	Logical. If TRUE, NA values in parameters_list are ignored during hash generation.
alphabetical_order	Logical. If TRUE, the names in parameters_list are sorted alphabetically before hash generation.
overwrite	Logical. If TRUE, existing files with the same hash will be overwritten. If FALSE and a conflict occurs, the results will be saved under a temporary hash.
include_timestamp	Logical. If TRUE, a timestamp is added to parameters_list.
hash_includes_timestamp	Logical. If TRUE, the timestamp is included in the hash generation.
algo	Character string specifying the hashing algorithm to use. Default is "xxhash64". See ?digest
get_script_name	Logical. If TRUE, attempts to get the script name and add it to parameters_list. Only works if script is run from command line, in an interactive session, this will always be NULL.
ignore_script_name	Logical. If TRUE, the script name is ignored during hash generation.

incremental	Logical. If TRUE, results are saved in a subfolder named after the hash and can be combined with <code>compress_incremental</code> . Note, if TRUE, no checks will be done for results that already exist, the user should check this in their script with <code>check_hash_existence</code> .
silent	Logical. If TRUE, no check is done that pairs of results files (parameters and associated results) is done. This check is not necessary, but done by default to keep the user aware of a scenario that usually results from manual file manipulation.

Details

This function saves R objects to disk with a file name based on a generated hash of the provided arguments. It supports incremental saving, where multiple results can be saved under the same hash in a subdirectory and later collected. This can be helpful for a simulation that runs and saves results in parallel for the SAME set of simulation parameters.

Value

No return value. This function is called for its side effects.

See Also

[read_objects\(\)](#)

Examples

```
## Setup
tmp_dir <- file.path(tempdir(), "example")
dir.create(tmp_dir)

## Example using parameter list to run simulation and save results
parameters_list <- list(
  iterations = 1000,
  x_dist = "rnorm",
  x_dist_options = list(n = 10, mean = 1, sd = 2),
  error_dist = "rnorm",
  error_dist_options = list(n = 10, mean = 0, sd = 1),
  beta0 = 1,
  beta1 = 1
)

betas <- numeric(parameters_list$iterations)
for (i in 1:parameters_list$iterations) {
  x <- do.call(parameters_list$x_dist, parameters_list$x_dist_options)
  err <- do.call(parameters_list$error_dist, parameters_list$error_dist_options)
  y <- parameters_list$beta0 + parameters_list$beta1*x + err
  betas[i] <- coef(lm(y ~ x))["x"]
}

save_objects(folder = tmp_dir, results = betas, parameters_list = parameters_list)

## Read back in (consider clearing environment before running)
```

```

## Re-setup
tmp_dir <- file.path(tempdir(), "example")

parameters_list <- list(
  iterations = 1000,
  x_dist = "rnorm",
  x_dist_options = list(n = 10, mean = 1, sd = 2),
  error_dist = "rnorm",
  error_dist_options = list(n = 10, mean = 0, sd = 1),
  beta0 = 1,
  beta1 = 1
)

betas <- read_objects(folder = tmp_dir, parameters_list = parameters_list)

## Cleanup
unlink(tmp_dir, recursive = TRUE)

```

start_tagging

Monitor result file usage and cleanup unused files

Description

Tagging is mainly helpful for removing unused results. `start_tagging()` initializes the tagging process by creating a txt file in the results directory which will keep a record of which results are being read by `read_objects()`. `cleanup()` removes any .rds files in the specified folder that are not listed in the tagging file. `close_tagging()` deletes the tagging file, ending the tagging session.

Usage

```

start_tagging(folder, tagging_file_name = "indxr_tagging.txt")

cleanup(
  folder,
  tagging_file_name = "indxr_tagging.txt",
  cutoff_date = NULL,
  request_confirmation = TRUE
)

close_tagging(folder, tagging_file_name = "indxr_tagging.txt")

```

Arguments

folder A character string specifying the path to the directory where the result files are saved and where the tagging file will be created.

tagging_file_name A character string for a txt file the tagging information is to be saved under.

`cutoff_date` A character string in "%Y-%m-%d %H:%M:%S" format used to specify that any tagged files before the date should also be removed.

`request_confirmation` Logical, if TRUE will request user input before proceeding to delete files.

Value

No return value. This function is called for its side effects.

Examples

```
## Setup
tmp_dir <- file.path(tempdir(), "example")
dir.create(tmp_dir)

## Save example objects
parameters_list1 <- list(example = "tagging1")
parameters_list2 <- list(example = "tagging2")
save_objects(folder = tmp_dir, results = 1, parameters_list = parameters_list1)
save_objects(folder = tmp_dir, results = 2, parameters_list = parameters_list2)

## See the files have been saved
list.files(tmp_dir)

## Start tagging
start_tagging(tmp_dir)

## Read back in one the first file, this causes this file to be tagged
res1 <- read_objects(folder = tmp_dir, parameters_list = parameters_list1)

## Remove untagged file without confirmation (that for parameters_list2)
cleanup(tmp_dir, request_confirmation = FALSE)

## See that one file was removed
list.files(tmp_dir)

## Close tagging (just removes tagging file)
close_tagging(tmp_dir)

## Cleanup
unlink(tmp_dir, recursive = TRUE)
```

update_from_hash_table

Update File Names Based on New Parameters in Adjusted Hash Table

Description

This function updates names of existing results by re-hashing each set of parameters with potentially updated values based on adjustments made to a hash table (see `?create_hash_table`) by user. It loads RDS files based on their existing hashes, compares to the corresponding entry in a hash table, generates new hashes where needed, and saves the files with the new hashes. The old files are deleted if their hashes differ from the new ones.

Usage

```
update_from_hash_table(  
  hash_table,  
  rds_folder,  
  hash_includes_timestamp = FALSE,  
  ignore_na = TRUE,  
  alphabetical_order = TRUE,  
  algo = "xxhash64"  
)
```

Arguments

<code>hash_table</code>	A file path to a modified hash table generated by <code>create_hash_table</code> .
<code>rds_folder</code>	A string specifying the directory containing the RDS files associated with the hash table.
<code>hash_includes_timestamp</code>	Logical; if TRUE, timestamps are included in the hash generation.
<code>ignore_na</code>	Logical; if TRUE, NA values are ignored during hash generation.
<code>alphabetical_order</code>	Logical; if TRUE, parameters are sorted alphabetically before hash generation.
<code>algo</code>	Character string specifying the hashing algorithm to use. Default is "xxhash64". See <code>?digest</code>

Value

The function does not return a value but saves updated RDS files and deletes old files as needed.

See Also

[create_hash_table\(\)](#)

Examples

```
## Setup  
tmp_dir <- file.path(tempdir(), "example")  
dir.create(tmp_dir)  
  
## Save objects  
obj1 <- rnorm(1000)  
obj2 <- data.frame()
```

```
x = runif(100),
y = "something",
z = rep(c(TRUE, FALSE), 50)
)
obj3 <- list(obj1, obj2)

params1 <- list(
  distribution = "normal",
  other_params = list(param1 = TRUE, param2 = 1, param3 = NA)
)
params2 <- list(
  distribution = "uniform",
  other_params = list(param1 = FALSE, param2 = 2, param3 = "1", param4 = 4)
)
params3 <- list(
  distribution = "composite",
  other_params = list(param1 = TRUE, param2 = 3, param3 = 1)
)

save_objects(tmp_dir, obj1, params1)
save_objects(tmp_dir, obj2, params2)
save_objects(tmp_dir, obj3, params3)

## Create hash table
create_hash_table(tmp_dir, save_path = file.path(tmp_dir, "hash_table.csv"))

## Read in hash table, make a change, and save
hash_table <- read.csv(file.path(tmp_dir, "hash_table.csv"))
hash_table$distribution <- "something different"
write.csv(hash_table, file.path(tmp_dir, "hash_table.csv"))

## See file names before change
list.files(tmp_dir)

update_from_hash_table(
  hash_table = file.path(tmp_dir, "hash_table.csv"),
  rds_folder = tmp_dir
)

## See difference to before running update_hash_table()
list.files(tmp_dir)

## Cleanup
unlink(tmp_dir, recursive = TRUE)
```

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