

Package ‘ravepipeline’

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Type Package

Title Reproducible Pipeline Infrastructure for Neuroscience

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Language en-US

Description Defines the underlying pipeline structure for reproducible neuroscience, adopted by 'RAVE' (reproducible analysis and visualization of intracranial electroencephalography); provides high-level class definition to build, compile, set, execute, and share analysis pipelines. Both R and 'Python' are supported, with 'Markdown' and 'shiny' dashboard templates for extending and building customized pipelines. See the full documentations at <<https://rave.wiki>>; to cite us, check out our paper by Magnotti, Wang, and Beauchamp (2020, <[doi:10.1016/j.neuroimage.2020.117341](https://doi.org/10.1016/j.neuroimage.2020.117341)>), or run `citation("`ravepipeline")` for details.

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base64-utils *Convert from and to 'base64' string*

Description

Encode or decode 'base64' raw or url-safe string

Usage

```
base64_urlencode(x)
```

```
base64_encode(x)
```

```
base64_urldecode(x)
```

```
base64_decode(x)
```

```
base64_plot(  
  expr,  
  width = 480,  
  height = 480,  
  ...,  
  quoted = FALSE,  
  envir = parent.frame()  
)
```

Arguments

x	for encoders, this is an R raw or character vectors; for decoders this is 'base64' encoded strings
expr	expression for plot, will saved to a png and converted to 'base64' string
width, height	image size in pixels
...	passed to png
quoted, envir	non-standard evaluation settings

Value

base64_encode, base64_plot returns 'base64' string in raw format; base64_urlencode returns 'base64' string url-safe format; base64_urldecode returns the original string; base64_decode returns original raw vectors.

Examples

```
# ---- For direct base64URI -----
```

```

file_raw <- as.raw(1:255)

# raw base64
base64_raw <- base64_encode(file_raw)
base64_raw

as.integer(base64_decode(base64_raw))

# ---- For URL-save base64 -----
# Can be used in URL
base64_url <- base64_urlencode(
  paste(c(letters, LETTERS, 0:9),
    collapse = ""))
base64_url

base64_urldecode(base64_url)

# ---- Convert R plots to base64 -----
img <- base64_plot({
  plot(1:10)
}, width = 320, height = 320)

# summary
print(img)

# get base64 content
img_base64 <- format(img, type = "content")

# save to png
tmppng <- tempfile(fileext = ".png")
writeBin(base64_decode(img_base64), con = tmppng)

# cleanup
unlink(tmppng)

# Format as svg
format(img, type = "html_svg")

```

dir_create2

Force creating directory with checks

Description

Force creating directory with checks

Usage

```
dir_create2(x, showWarnings = FALSE, recursive = TRUE, check = TRUE, ...)
```

Arguments

x path to create
showWarnings, recursive, ...
passed to [dir.create](#)
check whether to check the directory after creation

Value

Normalized path

Examples

```
path <- file.path(tempfile(), 'a', 'b', 'c')  
  
# The following are equivalent  
dir.create(path, showWarnings = FALSE, recursive = TRUE)  
  
dir_create2(path)
```

install_modules *Install 'RAVE' modules*

Description

Low-level function exported for down-stream 'RAVE' packages.

Usage

```
install_modules(modules, dependencies = FALSE)
```

Arguments

modules a vector of characters, repository names; default is to automatically determined from a public registry
dependencies whether to update dependent packages; default is false

Value

nothing

 logger

Logger system used by 'RAVE'

Description

Keep track of messages printed by modules or functions

Usage

```

logger(
    ...,
    level = c("info", "success", "warning", "error", "fatal", "debug", "trace"),
    calc_delta = "auto",
    .envir = parent.frame(),
    .sep = "",
    use_glue = FALSE,
    reset_timer = FALSE
)

set_logger_path(root_path, max_bytes, max_files)

logger_threshold(
    level = c("info", "success", "warning", "error", "fatal", "debug", "trace"),
    module_id,
    type = c("console", "file", "both")
)

logger_error_condition(cond, level = "error")

```

Arguments

..., .envir, .sep	passed to glue , if use_glue is true
level	the level of message, choices are 'info' (default), 'success', 'warning', 'error', 'fatal', 'debug', 'trace'
calc_delta	whether to calculate time difference between current message and previous message; default is 'auto', which prints time difference when level is 'debug'. This behavior can be changed by altering calc_delta by a logical TRUE to enable or FALSE to disable.
use_glue	whether to use glue to combine ...; default is false
reset_timer	whether to reset timer used by calc_delta
root_path	root directory if you want log messages to be saved to hard disks; if root_path is NULL, "", or nullfile , then logger path will be unset.
max_bytes	maximum file size for each logger partitions
max_files	maximum number of partition files to hold the log; old files will be deleted.

module_id	'RAVE' module identification string, or name-space; default is 'base'
type	which type of logging should be set; default is 'console', if file log is enabled through set_logger_path, type could be 'file' or 'both'. Default log level is 'info' on console and 'debug' on file.
cond	condition to log

Value

The message without time-stamps

Examples

```
logger("This is a message")

a <- 1
logger("A message with glue: a={a}")

logger("A message without glue: a={a}", use_glue = FALSE)

logger("Message A", calc_delta = TRUE, reset_timer = TRUE)
logger("Seconds before logging another message", calc_delta = TRUE)

# by default, debug and trace messages won't be displayed
logger('debug message', level = 'debug')

# adjust logger level, make sure `module_id` is a valid RAVE module ID
logger_threshold('debug', module_id = NULL)

# Debug message will display
logger('debug message', level = 'debug')

# Trace message will not display as it's lower than debug level
logger('trace message', level = 'trace')
```

module_add	<i>Add new 'RAVE' (2.0) module to current project</i>
------------	-------------------------------------------------------

Description

Creates a 'RAVE' pipeline with additional dashboard module from template.

Usage

```
module_add(
  module_id,
  module_label,
```

```

path = ".",
type = c("default", "bare", "scheduler", "python"),
...,
pipeline_name = module_id,
overwrite = FALSE
)

```

Arguments

module_id	module ID to create, must be unique; users cannot install two modules with identical module ID. We recommend that a module ID follows snake format, starting with lab name, for example, 'beauchamplab_imaging_preprocess', 'karaslab_freez', or 'upenn_ese25_foof'.
module_label	a friendly label to display in the dashboard
path	project root path; default is current directory
type	template to choose, options are 'default' and 'bare'
...	additional configurations to the module such as 'order', 'group', 'badge'
pipeline_name	the pipeline name to create along with the module; default is identical to module_id (strongly recommended); leave it default unless you know what you are doing.
overwrite	whether to overwrite existing module if module with same ID exists; default is false

Value

Nothing.

Examples

```

# For demonstrating this example only
project_root <- tempfile()
dir.create(project_root, showWarnings = FALSE, recursive = TRUE)

# Add a module
module_id <- "mylab_my_first_module"
module_add(
  module_id = module_id,
  module_label = "My Pipeline",
  path = project_root
)

# show the structure
cat(
  list.files(
    project_root,
    recursive = TRUE,
    full.names = FALSE,

```

```

        include.dirs = TRUE
    ),
    sep = "\n"
)

unlink(project_root, recursive = TRUE)

```

module_registry	<i>'RAVE' module registry</i>
-----------------	-------------------------------

Description

Create, view, or reserve the module registry

Usage

```

module_registry(
  title,
  repo,
  modules,
  authors,
  url = sprintf("https://github.com/%s", repo)
)

module_registry2(repo, description)

get_modules_registries(update = NA)

get_module_description(path)

add_module_registry(title, repo, modules, authors, url, dry_run = FALSE)

```

Arguments

title	title of the registry, usually identical to the description title in 'DESCRIPTION' or RAVE-CONFIG file
repo	'Github' repository
modules	characters of module ID, must only contain letters, digits, underscore, dash; must not be duplicated with existing registered modules
authors	a list of module authors; there must be one and only one author with 'cre' role (see person). This author will be considered maintainer, who will be in charge if editing the registry
url	the web address of the repository
update	whether to force updating the registry

path, description	path to 'DESCRIPTION' or RAVE-CONFIG file
dry_run	whether to generate and preview message content instead of opening an email link

Details

A 'RAVE' registry contains the following data entries: repository title, name, 'URL', authors, and a list of module IDs. 'RAVE' requires that each module must use a unique module ID. It will cause an issue if two modules share the same ID. Therefore 'RAVE' maintains a public registry list such that the module maintainers can register their own module ID and prevent other people from using it.

To register your own module ID, please use `add_module_registry` to validate and send an email to the 'RAVE' development team.

Value

a registry object, or a list of registries

Examples

```
library(ravepipeline)

# create your own registry
module_registry(
  repo = "rave-ieeg/rave-pipelines",
  title = "A Collection of 'RAVE' Builtin Pipelines",
  authors = list(
    list("Zhengjia", "Wang", role = c("cre", "aut"),
        email = "dipterix@rave.wiki")
  ),
  modules = "brain_viewer"
)

## Not run:

# This example will need access to Github and will open an email link

# get current registries
get_modules_registries(FALSE)

# If your repository is on Github and RAVE-CONFIG file exists
module_registry2("rave-ieeg/rave-pipelines")

# send a request to add your registry
registry <- module_registry2("rave-ieeg/rave-pipelines")
add_module_registry(registry)

## End(Not run)
```

pipeline	<i>Creates 'RAVE' pipeline instance</i>
----------	-----------------------------------------

Description

Set pipeline inputs, execute, and read pipeline outputs

Usage

```
pipeline(
  pipeline_name,
  settings_file = "settings.yaml",
  paths = pipeline_root(),
  temporary = FALSE
)

pipeline_from_path(path, settings_file = "settings.yaml")
```

Arguments

pipeline_name	the name of the pipeline, usually title field in the 'DESCRIPTION' file, or the pipeline folder name (if description file is missing)
settings_file	the name of the settings file, usually stores user inputs
paths	the paths to search for the pipeline, usually the parent directory of the pipeline; default is pipeline_root , which only search for pipelines that are installed or in current working directory.
temporary	see pipeline_root
path	the pipeline folder

Value

A [PipelineTools](#) instance

Examples

```
library(ravepipeline)

if(interactive()) {
  # ----- Set up a bare minimal example pipeline -----
  root_path <- tempdir()
  pipeline_root_folder <- file.path(root_path, "modules")

  # create pipeline folder
```

```

pipeline_path <- pipeline_create_template(
  root_path = pipeline_root_folder, pipeline_name = "raveio_demo",
  overwrite = TRUE, activate = FALSE, template_type = "rmd-bare")

# Set initial user inputs
yaml::write_yaml(
  x = list(
    n = 100,
    pch = 16,
    col = "steelblue"
  ),
  file = file.path(pipeline_path, "settings.yaml")
)

# build the pipeline for the first time
# this is a one-time setup
pipeline_build(pipeline_path)

# Temporarily redirect the pipeline project root
# to `root_path`
old_opt <- options("raveio.pipeline.project_root" = root_path)
# Make sure the options are reset
on.exit({ options(old_opt) })

# Compile the pipeline document
pipeline_render(
  module_id = "raveio_demo",
  project_path = root_path
)

## Not run:

# Open web browser to see compiled report
utils::browseURL(file.path(pipeline_path, "main.html"))

## End(Not run)

# ----- Example starts -----

# Load pipeline
pipeline <- pipeline(
  pipeline_name = "raveio_demo",
  paths = pipeline_root_folder,
  temporary = TRUE
)

# Check which pipeline targets to run
pipeline$target_table

# Run to `plot_data`, RAVE pipeline will automatically
# calculate which up-stream targets need to be updated
# and evaluate these targets

```

```

pipeline$run("plot_data")

# Customize settings
pipeline$set_settings(pch = 2)

# Run again with the new inputs, since input_data does not change,
# the pipeline will skip that target automatically
pipeline$run("plot_data")

# Read intermediate data
head(pipeline$read("input_data"))

# or use `[[]` to get results
pipeline[c("n", "pch", "col")]
pipeline[-c("input_data")]

# Check evaluating status
pipeline$progress("details")

# result summary & cache table
pipeline$result_table

# visualize the target dependency graph
pipeline$visualize(glimpse = TRUE)

# ----- Clean up -----
unlink(pipeline_path, recursive = TRUE)
}

```

pipeline-knitr-markdown

Configure 'rmarkdown' files to build 'RAVE' pipelines

Description

Allows building 'RAVE' pipelines from 'rmarkdown' files. Please use it in 'rmarkdown' scripts only. Use [pipeline_create_template](#) to create an example.

Usage

```

configure_knitr(languages = c("R", "python"))

pipeline_setup_rmd(
  module_id,
  env = parent.frame(),
  collapse = TRUE,
  comment = "#>",
  languages = c("R", "python"),

```

```

project_path = getOption("raveio.pipeline.project_root", default =
  rs_active_project(child_ok = TRUE, shiny_ok = TRUE))
)

pipeline_render(
  module_id,
  ...,
  env = new.env(parent = parent.frame()),
  entry_file = "main.Rmd",
  project_path = getOption("raveio.pipeline.project_root", default =
    rs_active_project(child_ok = TRUE, shiny_ok = TRUE))
)

```

Arguments

languages	one or more programming languages to support; options are 'R' and 'python'
module_id	the module ID, usually the name of direct parent folder containing the pipeline file
env	environment to set up the pipeline translator
collapse, comment	passed to set method of opts_chunk
project_path	the project path containing all the pipeline folders, usually the active project folder
...	passed to internal function calls
entry_file	the file to compile; default is "main.Rmd"

Value

A function that is supposed to be called later that builds the pipeline scripts

Examples

```

configure_knitr("R")

## Not run:

# Requires to configure Python
configure_knitr("python")

# This function must be called in an Rmd file setup block
# for example, see
# https://rave.wiki/posts/customize\_modules/python\_module\_01.html

pipeline_setup_rmd("my_module_id")

## End(Not run)

```

PipelineCollections *Connect and schedule pipelines*

Description

Experimental, subject to change in the future.

Value

A list containing

id the pipeline ID that can be used by deps

pipeline forked pipeline instance

target_names copy of names

depend_on copy of deps

cue copy of cue

standalone copy of standalone

Public fields

verbose whether to verbose the build

Active bindings

root_path path to the directory that contains pipelines and scheduler

collection_path path to the pipeline collections

pipeline_ids pipeline ID codes

Methods

Public methods:

- [PipelineCollections\\$new\(\)](#)
- [PipelineCollections\\$add_pipeline\(\)](#)
- [PipelineCollections\\$build_pipelines\(\)](#)
- [PipelineCollections\\$run\(\)](#)
- [PipelineCollections\\$get_scheduler\(\)](#)

Method new(): Constructor

Usage:

```
PipelineCollections$new(root_path = NULL, overwrite = FALSE)
```

Arguments:

root_path where to store the pipelines and intermediate results

overwrite whether to overwrite if root_path exists

Method `add_pipeline()`: Add pipeline into the collection

Usage:

```
PipelineCollections$add_pipeline(
  x,
  names = NULL,
  deps = NULL,
  pre_hook = NULL,
  post_hook = NULL,
  cue = c("always", "thorough", "never"),
  search_paths = pipeline_root(),
  standalone = TRUE,
  hook_envir = parent.frame()
)
```

Arguments:

`x` a pipeline name (can be found via `pipeline_list`), or a `PipelineTools`
`names` pipeline targets to execute
`deps` pipeline IDs to depend on; see 'Values' below
`pre_hook` function to run before the pipeline; the function needs two arguments: input map
 (can be edit in-place), and path to a directory that allows to store temporary files
`post_hook` function to run after the pipeline; the function needs two arguments: pipeline object,
 and path to a directory that allows to store intermediate results
`cue` whether to always run dependence
`search_paths` where to search for pipeline if `x` is a character; ignored when `x` is a pipeline
 object
`standalone` whether the pipeline should be standalone, set to TRUE if the same pipeline added
 multiple times should run independently; default is true
`hook_envir` where to look for global environments if `pre_hook` or `post_hook` contains global
 variables; default is the calling environment

Method `build_pipelines()`: Build pipelines and visualize

Usage:

```
PipelineCollections$build_pipelines(visualize = TRUE)
```

Arguments:

`visualize` whether to visualize the pipeline; default is true

Method `run()`: Run the collection of pipelines

Usage:

```
PipelineCollections$run(
  error = c("error", "warning", "ignore"),
  .scheduler = c("none", "future", "clustermq"),
  .type = c("callr", "smart", "vanilla"),
  .as_promise = FALSE,
  .async = FALSE,
  rebuild = NA,
  ...
)
```

Arguments:

error what to do when error occurs; default is 'error' throwing errors; other choices are 'warning' and 'ignore'

.scheduler, .type, .as_promise, .async, ... passed to [pipeline_run](#)

rebuild whether to re-build the pipeline; default is NA (if the pipeline has been built before, then do not rebuild)

Method `get_scheduler()`: Get scheduler object

Usage:

`PipelineCollections$get_scheduler()`

 PipelineResult

Class definition for 'RAVE' pipeline results

Description

Class definition for 'RAVE' pipeline results

Class definition for 'RAVE' pipeline results

Value

TRUE if the target is finished, or FALSE if timeout is reached

Public fields

progressor progress bar object, usually generated a progress instance

promise a [promise](#) instance that monitors the pipeline progress

verbose whether to print warning messages

names names of the pipeline to build

async_callback function callback to call in each check loop; only used when the pipeline is running in `async=TRUE` mode

check_interval used when `async=TRUE` in [pipeline_run](#), interval in seconds to check the progress

Active bindings

variables target variables of the pipeline

variable_descriptions readable descriptions of the target variables

valid logical true or false whether the result instance hasn't been invalidated

status result status, possible status are 'initialize', 'running', 'finished', 'canceled', and 'errored'. Note that 'finished' only means the pipeline process has been finished.

process (read-only) process object if the pipeline is running in 'async' mode, or NULL; see [r_bg](#).

Methods**Public methods:**

- PipelineResult\$validate()
- PipelineResult\$invalidate()
- PipelineResult\$get_progress()
- PipelineResult\$new()
- PipelineResult\$run()
- PipelineResult\$await()
- PipelineResult\$print()
- PipelineResult\$get_values()
- PipelineResult\$clone()

Method validate(): check if result is valid, raises errors when invalidated

Usage:

```
PipelineResult$validate()
```

Method invalidate(): invalidate the pipeline result

Usage:

```
PipelineResult$invalidate()
```

Method get_progress(): get pipeline progress

Usage:

```
PipelineResult$get_progress()
```

Method new(): constructor (internal)

Usage:

```
PipelineResult$new(path = character(0L), verbose = FALSE)
```

Arguments:

path pipeline path

verbose whether to print warnings

Method run(): run pipeline (internal)

Usage:

```
PipelineResult$run(  
  expr,  
  env = parent.frame(),  
  quoted = FALSE,  
  async = FALSE,  
  process = NULL  
)
```

Arguments:

expr expression to evaluate

env environment of expr

quoted whether expr has been quoted
async whether the process runs in other sessions
process the process object inherits [process](#), will be inferred from expr if process=NULL, and will raise errors if cannot be found

Method `await()`: wait until some targets get finished

Usage:

```
PipelineResult$await(names = NULL, timeout = Inf)
```

Arguments:

names target names to wait, default is NULL, i.e. to wait for all targets that have been scheduled
timeout maximum waiting time in seconds

Method `print()`: print method

Usage:

```
PipelineResult#print()
```

Method `get_values()`: get results

Usage:

```
PipelineResult$get_values(names = NULL, ...)
```

Arguments:

names the target names to read
... passed to [pipeline_read](#)

Method `clone()`: The objects of this class are cloneable with this method.

Usage:

```
PipelineResult$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

PipelineTools

Class definition for 'RAVE' pipelines

Description

Class definition for 'RAVE' pipelines

Class definition for 'RAVE' pipelines

Value

The value of the inputs, or a list if key is missing

The values of the targets

A [PipelineResult](#) instance if `as_promise` or `async` is true; otherwise a list of values for input names

An environment of shared variables

See `type`

A table of the progress

Nothing

ancestor target names (including names)

A new pipeline object based on the path given

A new pipeline object based on the path given

the saved file path

the data if file is found or a default value

A list of key-value pairs

A list of the preferences. If `simplify` is true and length of keys is 1, then returns the value of that preference

logical whether the keys exist

A job identification number, see [resolve_job](#) for querying job details

Super class

[ravepipeline::RAVESerializable](#) -> PipelineTools

Active bindings

`description` pipeline description

`settings_path` absolute path to the settings file

`extdata_path` absolute path to the user-defined pipeline data folder

`preference_path` directory to the pipeline preference folder

`target_table` table of target names and their descriptions

`result_table` summary of the results, including signatures of data and commands

`pipeline_path` the absolute path of the pipeline

`pipeline_name` the code name of the pipeline

`available_reports` available reports and their configurations

Methods**Public methods:**

- PipelineTools\$@marshal()
- PipelineTools\$@unmarshal()
- PipelineTools\$new()
- PipelineTools\$set_settings()
- PipelineTools\$get_settings()
- PipelineTools\$read()
- PipelineTools\$run()
- PipelineTools\$eval()
- PipelineTools\$shared_env()
- PipelineTools\$python_module()
- PipelineTools\$progress()
- PipelineTools\$attach()
- PipelineTools\$visualize()
- PipelineTools\$target_ancestors()
- PipelineTools\$fork()
- PipelineTools\$fork_to_subject()
- PipelineTools\$with_activated()
- PipelineTools\$clean()
- PipelineTools\$save_data()
- PipelineTools\$load_data()
- PipelineTools\$set_preferences()
- PipelineTools\$get_preferences()
- PipelineTools\$has_preferences()
- PipelineTools\$generate_report()
- PipelineTools\$clone()

Method @marshal(): Create an atomic list that can be serialized

Usage:

```
PipelineTools$@marshal(...)
```

Arguments:

... ignored

Method @unmarshal(): Restore an object from an atomic list

Usage:

```
PipelineTools$@unmarshal(object, ...)
```

Arguments:

object a list from '@marshal'

... ignored

Method new(): construction function

Usage:

```
PipelineTools$new(
  pipeline_name,
  settings_file = "settings.yaml",
  paths = pipeline_root(),
  temporary = FALSE
)
```

Arguments:

`pipeline_name` name of the pipeline, usually in the pipeline 'DESCRIPTION' file, or pipeline folder name

`settings_file` the file name of the settings file, where the user inputs are stored

`paths` the paths to find the pipeline, usually the parent folder of the pipeline; default is `pipeline_root()`

`temporary` whether not to save paths to current pipeline root registry. Set this to TRUE when importing pipelines from subject pipeline folders

Method `set_settings()`: set inputs*Usage:*

```
PipelineTools$set_settings(..., .list = NULL)
```

Arguments:

`...`, `.list` named list of inputs; all inputs should be named, otherwise errors will be raised

Method `get_settings()`: get current inputs*Usage:*

```
PipelineTools$get_settings(key, default = NULL, constraint)
```

Arguments:

`key` the input name; default is missing, i.e., to get all the settings

`default` default value if not found

`constraint` the constraint of the results; if input value is not from constraint, then only the first element of constraint will be returned.

Method `read()`: read intermediate variables*Usage:*

```
PipelineTools$read(var_names, ifnotfound = NULL, ...)
```

Arguments:

`var_names` the target names, can be obtained via `x$target_table` member; default is missing, i.e., to read all the intermediate variables

`ifnotfound` variable default value if not found

`...` other parameters passing to [pipeline_read](#)

Method `run()`: run the pipeline*Usage:*

```

PipelineTools$run(
  names = NULL,
  async = FALSE,
  as_promise = async,
  scheduler = c("none", "future", "clustermq"),
  type = c("smart", "callr", "vanilla"),
  envir = new.env(parent = globalenv()),
  callr_function = NULL,
  return_values = TRUE,
  debug = FALSE,
  ...
)

```

Arguments:

`names` pipeline variable names to calculate; default is to calculate all the targets

`async` whether to run asynchronous in another process

`as_promise` whether to return a [PipelineResult](#) instance

`scheduler`, `type`, `envir`, `callr_function`, `return_values`, `debug`, ... passed to [pipeline_run](#) if `as_promise` is true, otherwise these arguments will be passed to `pipeline_run_bare`

Method `eval()`: run the pipeline in order; unlike `$run()`, this method does not use the targets infrastructure, hence the pipeline results will not be stored, and the order of names will be respected.

Usage:

```

PipelineTools$eval(
  names,
  env = parent.frame(),
  shortcut = FALSE,
  clean = TRUE,
  ...
)

```

Arguments:

`names` pipeline variable names to calculate; must be specified

`env` environment to evaluate and store the results

`shortcut` logical or characters; default is FALSE, meaning names and all the dependencies (if missing from `env`) will be evaluated; set to TRUE if only names are to be evaluated. When `shortcut` is a character vector, it should be a list of targets (including their ancestors) whose values can be assumed to be up-to-date, and the evaluation of those targets can be skipped.

`clean` whether to evaluate without polluting `env`

... passed to [pipeline_eval](#)

Method `shared_env()`: run the pipeline shared library in scripts starting with path R/shared

Usage:

```
PipelineTools$shared_env(callr_function = callr::r)
```

Arguments:

`callr_function` either `callr::r` or `NULL`; when `callr::r`, the environment will be loaded in isolated R session and serialized back to the main session to avoid contaminating the main session environment; when `NULL`, the code will be sourced directly in current environment.

Method `python_module()`: get 'Python' module embedded in the pipeline

Usage:

```
PipelineTools$python_module(
  type = c("info", "module", "shared", "exist"),
  must_work = TRUE
)
```

Arguments:

`type` return type, choices are 'info' (get basic information such as module path, default), 'module' (load module and return it), 'shared' (load a shared sub-module from the module, which is shared also in report script), and 'exist' (returns true or false on whether the module exists or not)

`must_work` whether the module needs to be existed or not. If `TRUE`, the raise errors when the module does not exist; default is `TRUE`, ignored when type is 'exist'.

Method `progress()`: get progress of the pipeline

Usage:

```
PipelineTools$progress(method = c("summary", "details"))
```

Arguments:

`method` either 'summary' or 'details'

Method `attach()`: attach pipeline tool to environment (internally used)

Usage:

```
PipelineTools$attach(env)
```

Arguments:

`env` an environment

Method `visualize()`: visualize pipeline target dependency graph

Usage:

```
PipelineTools$visualize(
  glimpse = FALSE,
  aspect_ratio = 2,
  node_size = 30,
  label_size = 40,
  ...
)
```

Arguments:

`glimpse` whether to glimpse the graph network or render the state

`aspect_ratio` controls node spacing

`node_size`, `label_size` size of nodes and node labels

... passed to `pipeline_visualize`

Method `target_ancestors()`: a helper function to get target ancestors

Usage:

```
PipelineTools$target_ancestors(names, skip_names = NULL)
```

Arguments:

`names` targets whose ancestor targets need to be queried

`skip_names` targets that are assumed to be up-to-date, hence will be excluded, notice this exclusion is recursive, that means not only `skip_names` are excluded, but also their ancestors will be excluded from the result.

Method `fork()`: fork (copy) the current pipeline to a new directory

Usage:

```
PipelineTools$fork(path, policy = "default")
```

Arguments:

`path` path to the new pipeline, a folder will be created there

`policy` fork policy defined by module author, see text file 'fork-policy' under the pipeline directory; if missing, then default to avoid copying `main.html` and `shared` folder

Method `fork_to_subject()`: fork (copy) the current pipeline to a 'RAVE' subject

Usage:

```
PipelineTools$fork_to_subject(
  subject,
  label = "NA",
  policy = "default",
  delete_old = FALSE,
  sanitize = TRUE
)
```

Arguments:

`subject` subject ID or instance in which pipeline will be saved

`label` pipeline label describing the pipeline

`policy` fork policy defined by module author, see text file 'fork-policy' under the pipeline directory; if missing, then default to avoid copying `main.html` and `shared` folder

`delete_old` whether to delete old pipelines with the same label default is false

`sanitize` whether to sanitize the registry at save. This will remove missing folders and import manually copied pipelines to the registry (only for the pipelines with the same name)

Method `with_activated()`: run code with pipeline activated, some environment variables and function behaviors might change under such condition (for example, `targets` package functions)

Usage:

```
PipelineTools$with_activated(expr, quoted = FALSE, env = parent.frame())
```

Arguments:

`expr` expression to evaluate

`quoted` whether `expr` is quoted; default is false

`env` environment to run `expr`

Method `clean()`: clean all or part of the data store

Usage:

```
PipelineTools$clean(
  destroy = c("all", "cloud", "local", "meta", "process", "preferences", "progress",
             "objects", "scratch", "workspaces"),
  ask = FALSE
)
```

Arguments:

`destroy`, `ask` see [tar_destroy](#)

Method `save_data()`: save data to pipeline data folder

Usage:

```
PipelineTools$save_data(
  data,
  name,
  format = c("json", "yaml", "csv", "fst", "rds"),
  overwrite = FALSE,
  ...
)
```

Arguments:

`data` R object

`name` the name of the data to save, must start with letters

`format` serialize format, choices are 'json', 'yaml', 'csv', 'fst', 'rds'; default is 'json'.

To save arbitrary objects such as functions or environments, use 'rds'

`overwrite` whether to overwrite existing files; default is no

... passed to saver functions

Method `load_data()`: load data from pipeline data folder

Usage:

```
PipelineTools$load_data(
  name,
  error_if_missing = TRUE,
  default_if_missing = NULL,
  format = c("auto", "json", "yaml", "csv", "fst", "rds"),
  ...
)
```

Arguments:

`name` the name of the data

`error_if_missing` whether to raise errors if the name is missing

`default_if_missing` default values to return if the name is missing

`format` the format of the data, default is automatically obtained from the file extension

... passed to loader functions

Method `set_preferences()`: set persistent preferences from the pipeline. The preferences should not affect how pipeline is working, hence usually stores minor variables such as graphic options. Changing preferences will not invalidate pipeline cache.

Usage:

```
PipelineTools$set_preferences(..., .list = NULL)
```

Arguments:

`...`, `.list` key-value pairs of initial preference values. The keys must start with 'global' or the module ID, followed by dot and preference type and names. For example 'global.graphics.continuous_palette' for setting palette colors for continuous heat-map; "global" means the settings should be applied to all 'RAVE' modules. The module-level preference, 'power_explorer.export.default_format' sets the default format for power-explorer export dialogue.

`name` preference name, must contain only letters, digits, underscore, and hyphen, will be coerced to lower case (case-insensitive)

Method `get_preferences()`: get persistent preferences from the pipeline.

Usage:

```
PipelineTools$get_preferences(
  keys,
  simplify = TRUE,
  ifnotfound = NULL,
  validator = NULL,
  ...
)
```

Arguments:

`keys` characters to get the preferences

`simplify` whether to simplify the results when length of key is 1; default is true; set to false to always return a list of preferences

`ifnotfound` default value when the key is missing

`validator` NULL or function to validate the values; see 'Examples'

`...` passed to validator if validator is a function

Examples:

```
library(ravepipeline)
if(interactive() && length(pipeline_list()) > 0) {
  pipeline <- pipeline("power_explorer")

  # set dummy preference
  pipeline$set_preferences("global.example.dummy_preference" = 1:3)

  # get preference
  pipeline$get_preferences("global.example.dummy_preference")

  # get preference with validator to ensure the value length to be 1
  pipeline$get_preferences(
    "global.example.dummy_preference",
    validator = function(value) {
      stopifnot(length(value) == 1)
    },
    ifnotfound = 100
  )
}
```

```

    pipeline$has_preferences("global.example.dummy_preference")
}

```

Method `has_preferences()`: whether pipeline has preference keys

Usage:

```
PipelineTools$has_preferences(keys, ...)
```

Arguments:

`keys` characters name of the preferences

`...` passed to internal methods

Method `generate_report()`: generate pipeline

Usage:

```

PipelineTools$generate_report(
  name,
  subject = NULL,
  output_dir = NULL,
  output_format = "auto",
  clean = FALSE,
  ...
)

```

Arguments:

`name` report name, see field 'available_reports'

`subject` subject helps determine the output_dir and working directories

`output_dir` parent folder where output will be stored

`output_format` output format

`clean` whether to clean the output; default is false

`...` passed to 'rmarkdown' render function

Method `clone()`: The objects of this class are cloneable with this method.

Usage:

```
PipelineTools$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

See Also

[pipeline](#)

Examples

```

## -----
## Method `PipelineTools$get_preferences`
## -----

```

```
library(ravepipeline)
if(interactive() && length(pipeline_list()) > 0) {
  pipeline <- pipeline("power_explorer")

  # set dummy preference
  pipeline$set_preferences("global.example.dummy_preference" = 1:3)

  # get preference
  pipeline$get_preferences("global.example.dummy_preference")

  # get preference with validator to ensure the value length to be 1
  pipeline$get_preferences(
    "global.example.dummy_preference",
    validator = function(value) {
      stopifnot(length(value) == 1)
    },
    ifnotfound = 100
  )

  pipeline$has_preferences("global.example.dummy_preference")
}
```

pipeline_collection *Combine and execute pipelines*

Description

Experimental, subject to change in the future.

Usage

```
pipeline_collection(root_path = NULL, overwrite = FALSE)
```

Arguments

root_path	directory to store pipelines and results
overwrite	whether to overwrite if root_path exists; default is false, and raises an error when root_path exists

Value

A [PipelineCollections](#) instance

pipeline_install *Install 'RAVE' pipelines*

Description

Install 'RAVE' pipelines

Usage

```
pipeline_install_local(
  src,
  to = c("default", "custom", "workdir", "tempdir"),
  upgrade = FALSE,
  force = FALSE,
  set_default = NA,
  ...
)

pipeline_install_github(
  repo,
  to = c("default", "custom", "workdir", "tempdir"),
  upgrade = FALSE,
  force = FALSE,
  set_default = NA,
  ...
)
```

Arguments

src	pipeline directory
to	installation path; choices are 'default', 'custom', 'workdir', and 'tempdir'. Please specify pipeline root path via pipeline_root when 'custom' is used.
upgrade	whether to upgrade the dependence; default is FALSE for stability, however, it is highly recommended to upgrade your dependencies
force	whether to force installing the pipelines
set_default	whether to set current pipeline module folder as the default, will be automatically set when the pipeline is from the official 'Github' repository.
...	other parameters not used
repo	'Github' repository in user-repository combination, for example, 'rave-ieeg/rave-pipeline'

Value

nothing

Examples

```
## Not run:

pipeline_install_github("rave-ieeg/pipelines")

# or download github.com/rave-ieeg/pipelines repository, extract
# to a folder, and call
pipeline_install_local("path/to/pipeline/folder")

## End(Not run)
```

```
pipeline_settings_get_set
```

Get or change pipeline input parameter settings

Description

Get or change pipeline input parameter settings

Usage

```
pipeline_settings_set(
  ...,
  pipeline_path = Sys.getenv("RAVE_PIPELINE", "."),
  pipeline_settings_path = file.path(pipeline_path, "settings.yaml")
)

pipeline_settings_get(
  key,
  default = NULL,
  constraint = NULL,
  pipeline_path = Sys.getenv("RAVE_PIPELINE", "."),
  pipeline_settings_path = file.path(pipeline_path, "settings.yaml")
)
```

Arguments

`pipeline_path` the root directory of the pipeline

`pipeline_settings_path` the settings file of the pipeline, must be a 'yaml' file; default is 'settings.yaml' in the current pipeline

`key, ...` the character key(s) to get or set

default	the default value is key is missing
constraint	the constraint of the resulting value; if not NULL, then result must be within the constraint values, otherwise the first element of constraint will be returned. This is useful to make sure the results stay within given options

Value

`pipeline_settings_set` returns a list of all the settings. `pipeline_settings_get` returns the value of given key.

Examples

```

root_path <- tempfile()
pipeline_root_folder <- file.path(root_path, "modules")

# create pipeline folder
pipeline_path <- pipeline_create_template(
  root_path = pipeline_root_folder, pipeline_name = "raveio_demo",
  overwrite = TRUE, activate = FALSE, template_type = "rmd-bare")

# Set initial user inputs
yaml::write_yaml(
  x = list(
    n = 100,
    pch = 16,
    col = "steelblue"
  ),
  file = file.path(pipeline_path, "settings.yaml")
)

# build the pipeline for the first time
# this is a one-time setup
pipeline_build(pipeline_path)

# get pipeline settings
pipeline_settings_get(
  key = "n",
  pipeline_path = pipeline_path
)

# get variable with default if missing
pipeline_settings_get(
  key = "missing_variable",
  default = "missing",
  pipeline_path = pipeline_path
)

pipeline_settings_set(
  missing_variable = "A",
  pipeline_path = pipeline_path

```

```
)

pipeline_settings_get(
  key = "missing_variable",
  default = "missing",
  pipeline_path = pipeline_path
)

unlink(root_path, recursive = TRUE)
```

rave-pipeline *Low-level 'RAVE' pipeline functions*

Description

Utility functions for 'RAVE' pipelines, currently designed for internal development use. The infrastructure will be deployed to 'RAVE' in the future to facilitate the "self-expanding" aim. Please check the official 'RAVE' website.

Usage

```
pipeline_root(root_path, temporary = FALSE)

pipeline_list(root_path = pipeline_root())

pipeline_find(name, root_path = pipeline_root())

pipeline_attach(name, root_path = pipeline_root())

pipeline_run(
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),
  scheduler = c("none", "future", "clustermq"),
  type = c("smart", "callr", "vanilla"),
  envir = new.env(parent = globalenv()),
  callr_function = NULL,
  names = NULL,
  async = FALSE,
  check_interval = 0.5,
  progress_quiet = !async,
  progress_max = NA,
  progress_title = "Running pipeline",
  return_values = TRUE,
  debug = FALSE,
  ...
)
```

```
pipeline_clean(  
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),  
  destroy = c("all", "cloud", "local", "meta", "process", "preferences", "progress",  
    "objects", "scratch", "workspaces"),  
  ask = FALSE  
)  
  
pipeline_run_bare(  
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),  
  scheduler = c("none", "future", "clustermq"),  
  type = c("smart", "callr", "vanilla"),  
  envir = new.env(parent = globalenv()),  
  callr_function = NULL,  
  names = NULL,  
  return_values = TRUE,  
  debug = FALSE,  
  ...  
)  
  
load_targetgets(..., env = NULL)  
  
pipeline_target_names(pipe_dir = Sys.getenv("RAVE_PIPELINE", "."))  
  
pipeline_debug(  
  quick = TRUE,  
  env = parent.frame(),  
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),  
  skip_names  
)  
  
pipeline_dep_targetgets(  
  names,  
  skip_names = NULL,  
  pipe_dir = Sys.getenv("RAVE_PIPELINE", ".")  
)  
  
pipeline_eval(  
  names,  
  env = new.env(parent = parent.frame()),  
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),  
  settings_path = file.path(pipe_dir, "settings.yaml"),  
  shortcut = FALSE  
)  
  
pipeline_visualize(  
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),  
  glimpse = FALSE,
```

```
    targets_only = TRUE,
    shortcut = FALSE,
    zoom_speed = 0.1,
    ...
)

pipeline_progress(
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),
  method = c("summary", "details", "custom"),
  func = targets::tar_progress_summary
)

pipeline_fork(
  src = Sys.getenv("RAVE_PIPELINE", "."),
  dest = tempfile(pattern = "rave_pipeline_"),
  policy = "default",
  activate = FALSE,
  ...
)

pipeline_build(pipe_dir = Sys.getenv("RAVE_PIPELINE", "."))

pipeline_read(
  var_names,
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),
  branches = NULL,
  ifnotfound = NULL,
  dependencies = c("none", "ancestors_only", "all"),
  simplify = TRUE,
  ...
)

pipeline_vartable(
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),
  targets_only = TRUE,
  complete_only = FALSE,
  ...
)

pipeline_hasname(var_names, pipe_dir = Sys.getenv("RAVE_PIPELINE", "."))

pipeline_watch(
  pipe_dir = Sys.getenv("RAVE_PIPELINE", "."),
  targets_only = TRUE,
  ...
)

pipeline_create_template(
```

```
    root_path,
    pipeline_name,
    overwrite = FALSE,
    activate = TRUE,
    template_type = c("rmd", "r", "rmd-bare", "rmd-scheduler", "rmd-python")
  )

pipeline_create_subject_pipeline(
  subject,
  pipeline_name,
  overwrite = FALSE,
  activate = TRUE,
  template_type = c("rmd", "r", "rmd-python")
)

pipeline_description(file)

pipeline_load_extdata(
  name,
  format = c("auto", "json", "yaml", "csv", "fst", "rds"),
  error_if_missing = TRUE,
  default_if_missing = NULL,
  pipe_dir = Sys.getenv("RAVE_PIPELINES", "."),
  ...
)

pipeline_save_extdata(
  data,
  name,
  format = c("json", "yaml", "csv", "fst", "rds"),
  overwrite = FALSE,
  pipe_dir = Sys.getenv("RAVE_PIPELINES", "."),
  ...
)

pipeline_shared(
  pipe_dir = Sys.getenv("RAVE_PIPELINES", "."),
  callr_function = callr::r
)

pipeline_set_preferences(
  ...,
  .list = NULL,
  .pipe_dir = Sys.getenv("RAVE_PIPELINES", "."),
  .preference_instance = NULL
)

pipeline_get_preferences(
```

```

    keys,
    simplify = TRUE,
    ifnotfound = NULL,
    validator = NULL,
    ...,
    .preference_instance = NULL
)

pipeline_has_preferences(keys, ..., .preference_instance = NULL)

```

Arguments

root_path	the root directory for pipeline templates
temporary	whether not to save paths to current pipeline root registry. Set this to TRUE when importing pipelines from subject pipeline folders
name, pipeline_name	the pipeline name to create; usually also the folder
pipe_dir, .pipe_dir	where the pipeline directory is; can be set via system environment <code>Sys.setenv("RAVE_PIPELINE"=...)</code>
scheduler	how to schedule the target jobs: default is 'none', which is sequential. If you have multiple heavy-weighted jobs that can be scheduled at the same time, you can choose 'future' or 'clustermq'
type	how the pipeline should be executed; current choices are "smart" to enable 'future' package if possible, 'callr' to use <code>r</code> , or 'vanilla' to run everything sequentially in the main session.
callr_function	function that will be passed to <code>tar_make</code> ; will be forced to be NULL if type='vanilla', or <code>r</code> if type='callr'
names	the names of pipeline targets that are to be executed; default is NULL, which runs all targets; use <code>pipeline_target_names</code> to check all your available target names.
async	whether to run pipeline without blocking the main session
check_interval, progress_title, progress_max, progress_quiet	when running in background (non-blocking mode), how often to check the pipeline control the progress
return_values	whether to return pipeline target values; default is true; only works in <code>pipeline_run_bare</code> and will be ignored by <code>pipeline_run</code>
debug	whether to debug the process; default is false
..., .list	other parameters, targets, etc.
destroy	what part of data repository needs to be cleaned
ask	whether to ask
env, envir	environment to execute the pipeline
quick	whether to skip finished targets to save time
skip_names	hint of target names to fast skip provided they are up-to-date; only used when quick=TRUE. If missing, then skip_names will be automatically determined

settings_path	path to settings file name within subject's pipeline path
shortcut	whether to display shortcut targets
glimpse	whether to hide network status when visualizing the pipelines
targets_only	whether to return the variable table for targets only; default is true
zoom_speed	zoom speed when visualizing the pipeline dependence
method	how the progress should be presented; choices are "summary", "details", "custom". If custom method is chosen, then func will be called
func	function to call when reading customized pipeline progress; default is tar_progress_summary
src, dest	pipeline folder to copy the pipeline script from and to
policy	fork policy defined by module author, see text file 'fork-policy' under the pipeline directory; if missing, then default to avoid copying main.html and shared folder
activate	whether to activate the new pipeline folder from dest; default is false
var_names	variable name to fetch or to check
branches	branch to read from; see tar_read
ifnotfound	default values to return if variable is not found
dependencies	whether to load dependent targets, choices are 'none' (default, only load targets specified by names), 'ancestors_only' (load all but the ancestors targets), and 'all' (both targets and ancestors)
simplify	whether to simplify the output
complete_only	whether only to show completed and up-to-date target variables; default is false
overwrite	whether to overwrite existing pipeline; default is false so users can double-check; if true, then existing pipeline, including the data will be erased
template_type	which template type to create; choices are 'r' or 'rmd'
subject	character indicating valid 'RAVE' subject ID, or a RAVESubject instance
file	path to the 'DESCRIPTION' file under the pipeline folder, or pipeline collection folder that contains the pipeline information, structures, dependencies, etc.
format	format of the extended data, default is 'json', other choices are 'yaml', 'fst', 'csv', 'rds'
error_if_missing, default_if_missing	what to do if the extended data is not found
data	extended data to be saved
.preference_instance	internally used
keys	preference keys
validator	NULL or function to validate values

Value

pipeline_root the root directories of the pipelines
 pipeline_list the available pipeline names under pipeline_root
 pipeline_find the path to the pipeline
 pipeline_run a `PipelineResult` instance
 load_targets a list of targets to build
 pipeline_target_names a vector of characters indicating the pipeline target names
 pipeline_visualize a widget visualizing the target dependence structure
 pipeline_progress a table of building progress
 pipeline_fork a normalized path of the forked pipeline directory
 pipeline_read the value of corresponding var_names, or a named list if var_names has more than one element
 pipeline_vartable a table of summaries of the variables; can raise errors if pipeline has never been executed
 pipeline_hasname logical, whether the pipeline has variable built
 pipeline_watch a basic shiny application to monitor the progress
 pipeline_description the list of descriptions of the pipeline or pipeline collection

rave-pipeline-jobs	<i>Run a function (job) in another session</i>
--------------------	------------------------------------------------

Description

Run a function (job) in another session

Usage

```

start_job(
  fun,
  fun_args = list(),
  packages = NULL,
  workdir = NULL,
  method = c("callr", "rs_job", "mirai"),
  name = NULL,
  ensure_init = TRUE,
  digest_key = NULL,
  envvars = NULL
)

check_job(job_id)

resolve_job(

```

```

    job_id,
    timeout = Inf,
    auto_remove = TRUE,
    must_init = TRUE,
    unresolved = c("warning", "error", "silent")
)

remove_job(job_id)

```

Arguments

fun	function to evaluate
fun_args	list of function arguments
packages	list of packages to load
workdir	working directory; default is temporary path
method	job type; choices are 'rs_job' (only used in 'RStudio' environment), 'mirai' (when package 'mirai' is installed), and 'callr' (default).
name	name of the job
ensure_init	whether to make sure the job has been started; default is true
digest_key	a string that will affect how job ID is generated; used internally
envvars	additional environment variables to set; must be a named list of environment variables
job_id	job identification number
timeout	timeout in seconds before the resolve ends; jobs that are still running are subject to unresolved policy
auto_remove	whether to automatically remove the job if resolved; default is true
must_init	whether the resolve should error out if the job is not initialized: typically meaning either the resolving occurs too soon (only when ensure_init=FALSE) or the job files are corrupted; default is true
unresolved	what to do if the job is still running after timing-out; default is 'warning' and return NULL, other choices are 'error' or 'silent'

Value

For `start_job`, a string of job identification number; `check_job` returns the job status; `resolve_job` returns the function result.

Examples

```

## Not run:

# Basic use
job_id <- start_job(function() {
  Sys.sleep(1)
  Sys.getpid()
})

```

```
check_job(job_id)

result <- resolve_job(job_id)

# As promise
library(promises)
as.promise(
  start_job(function() {
    Sys.sleep(1)
    Sys.getpid()
  })
) %...>%
print()

## End(Not run)
```

rave-serialize-refhook

Serialization reference hook generic functions

Description

Serialization reference hook generic functions

Usage

```
rave_serialize_refhook(object)

rave_serialize_impl(object)

## Default S3 method:
rave_serialize_impl(object)

## S3 method for class 'RAVEserializable'
rave_serialize_impl(object)

## S3 method for class '`rave-brain`'
rave_serialize_impl(object)

rave_unserialize_refhook(x)

rave_unserialize_impl(x)

## Default S3 method:
```

```

rave_unserialize_impl(x)

## S3 method for class 'rave_serialized'
rave_unserialize_impl(x)

## S3 method for class 'rave_serialized_r6'
rave_unserialize_impl(x)

## S3 method for class '`rave_serialized_rave-brain`'
rave_unserialize_impl(x)

```

Arguments

object	Object to serialize (environment or external pointers)
x	raw or string objects that will be passed to <code>unserialize</code> function before further reconstruction

Value

`rave_serialize_refhook` returns either serialized objects in string (raw vector converted to char via `rawToChar`), or NULL indicating the object undergoing default serialization; `rave_unserialize_refhook` returns the reconstructed object.

Examples

```

# This example requires additional `filearray` package
# If you are an RAVE user (installed RAVE via rave.wiki)
# then this package was installed

x0 <- array(rnorm(240000), c(200, 300, 4))
x1 <- filearray::as_filearray(x0)
x2 <- RAVEFileArray$new(x1, temporary = TRUE)

r0 <- serialize(x0, NULL, refhook = rave_serialize_refhook)
r1 <- serialize(x1, NULL, refhook = rave_serialize_refhook)
r2 <- serialize(x2, NULL, refhook = rave_serialize_refhook)

# Compare the serialization sizes
c(length(r0), length(r1), length(r2))

y0 <- unserialize(r0, refhook = rave_unserialize_refhook)
y1 <- unserialize(r1, refhook = rave_unserialize_refhook)
y2 <- unserialize(r2, refhook = rave_unserialize_refhook)

all(y0 == x0)
all(y1[] == x0)
all(y2[] == x0)

## Not run:

# 3D Brain, this example needs RAVE installation, not included in

```

```
# this package, needs extra installations available at rave.wiki

# 4 MB
brain <- ravecore::rave_brain("demo/DemoSubject")

# 52 KB
rbrain <- serialize(brain, NULL, refhook = rave_serialize_refhook)

brain2 <- unserialize(rbrain, refhook = rave_unserialize_refhook)

brain2$plot()

## End(Not run)
```

rave-snippet *'RAVE' code snippets*

Description

Run snippet code

Usage

```
update_local_snippet(force = TRUE)

install_snippet(path)

list_snippets()

load_snippet(topic, local = TRUE)
```

Arguments

force	whether to force updating the snippets; default is true
path	for installing code snippets locally only; can be an R script, a zip file, or a directory
topic	snippet topic
local	whether to use local snippets first before requesting online repository

Value

load_snippet returns snippet as a function, others return nothing

Examples

```

# This example script requires running in an interactive session

if(interactive()){

# ---- Example 1: Install built-in pipeline snippets -----
update_local_snippet(force = TRUE)

# ---- Example 2: Install customized pipeline snippets -----
snippets <- file.path(
  "https://github.com/rave-ieeg/rave-gists",
  "archive/refs/heads/main.zip",
  fsep = "/"
)
tempf <- tempfile(fileext = ".zip")
utils::download.file(url = snippets, destfile = tempf)

install_snippet(tempf)
}

# ---- List snippets -----

# list all topics
list_snippets()

# ---- Run snippets as functions -----

topic <- "image-burn-contacts-to-t1"

# check whether this example can run
# This snippet requires installing package `raveio`
# which is currently not on CRAN (soon it will)

condition_met <- topic %in% list_snippets() &&
  (system.file(package = "raveio") != "")

if( interactive() && condition_met ) {

  snippet <- load_snippet(topic)

  # Read snippet documentation
  print(snippet)

  results <- snippet(
    subject_code = "DemoSubject",
    project_name = "demo",
    save_path = NA,
    blank_underlay = FALSE
  )
}

```

```

    plot(results)
  }

```

RAVEFileArray	'R6' wrapper for 'FileArray'
---------------	------------------------------

Description

Wrapper for better serialization (check 'See also')

Super class

`ravepipeline::RAVESerializable` -> RAVEFileArray

Public fields

`temporary` whether this file array is to be upon garbage collection; default is false. The file array will be deleted if the temporary flag is set to true and the array mode is 'readwrite'

Active bindings

`valid` whether the array is valid and ready to read

`@impl` the underlying array object

Methods

Public methods:

- `RAVEFileArray$@marshal()`
- `RAVEFileArray$@unmarshal()`
- `RAVEFileArray$new()`
- `RAVEFileArray$clone()`

Method `@marshal()`: Serialization helper, convert the object to a descriptive list

Usage:

`RAVEFileArray$@marshal(...)`

Arguments:

... ignored

Method `@unmarshal()`: Serialization helper, convert the object from a descriptive list

Usage:

`RAVEFileArray$@unmarshal(object, ...)`

Arguments:

object serialized list

... ignored

Method `new()`: Constructor

Usage:

```
RAVEFileArray$new(x, temporary = FALSE)
```

Arguments:

`x` file array or can be converted to [as_filearray](#)

`temporary` whether this file array is to be deleted once the object is out-of-scope; default is `false`

Method `clone()`: The objects of this class are cloneable with this method.

Usage:

```
RAVEFileArray$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

See Also

[RAVESerializable](#) [rave-serialize-refhook](#)

raveio-option

Set/Get 'RAVE' option

Description

Persist settings on local configuration file

Usage

```
raveio_setopt(key, value, .save = TRUE)
```

```
raveio_resetopt(all = FALSE)
```

```
raveio_getopt(key, default = NA, temp = TRUE)
```

```
raveio_confpath(cfile = "settings.yaml")
```

Arguments

<code>key</code>	character, option name
<code>value</code>	character or logical of length 1, option value
<code>.save</code>	whether to save to local drive, internally used to temporary change option. Not recommended to use it directly.
<code>all</code>	whether to reset all non-default keys

default	is key not found, return default value
temp	when saving, whether the key-value pair should be considered temporary, a temporary settings will be ignored when saving; when getting options, setting temp to false will reveal the actual settings.
cfile	file name in configuration path

Details

`raveio_setopt` stores key-value pair in local path. The values are persistent and shared across multiple sessions. There are some read-only keys such as "session_string". Trying to set those keys will result in error.

The following keys are reserved by 'RAVE':

`data_dir` Directory path, where processed data are stored; default is at home directory, folder `~/rave_data/data_dir`

`raw_data_dir` Directory path, where raw data files are stored, mainly the original signal files and imaging files; default is at home directory, folder `~/rave_data/raw_dir`

`max_worker` Maximum number of CPU cores to use; default is one less than the total number of CPU cores

`mni_template_root` Directory path, where 'MNI' templates are stored

`raveio_getopt` returns value corresponding to the keys. If key is missing, the whole option will be returned.

If set `all=TRUE`, `raveio_resetopt` resets all keys including non-standard ones. However "session_string" will never reset.

Value

`raveio_setopt` returns modified value; `raveio_resetopt` returns current settings as a list; `raveio_confpath` returns absolute path for the settings file; `raveio_getopt` returns the settings value to the given key, or `default` if not found.

Side-Effects

The following options will alter other packages and might cause changes in behaviors:

'`disable_fork_clusters`' This option will change the `options` '`dipsaus.no.fork`' and '`dipsaus.cluster.backup`', which handles the parallel computing

'`threeBrain_template_subject`' This option will set and persist option '`threeBrain.template_subject`', which changes the default group-level template brain.

See Also

`R_user_dir`

Examples

```
# get one RAVE option
ncore <- raveio_getopt("max_worker")
print(ncore)

# get all options
raveio_getopt()

# set option
raveio_setopt("disable_fork_clusters", FALSE)
```

ravepipeline-constants

Constant variables used in 'RAVE' pipeline

Description

Regular expression PIPELINE_FORK_PATTERN defines the file matching rules when forking a pipeline; see [pipeline_fork](#) for details.

Usage

PIPELINE_FORK_PATTERN

Format

An object of class character of length 1.

ravepipeline_finalize_installation

Download 'RAVE' built-in pipelines and code snippets

Description

The official built-in pipeline repository is located at <https://github.com/rave-ieeg/rave-pipelines>;
The code snippet repository is located at <https://github.com/rave-ieeg/rave-gists>.

Usage

```
ravepipeline_finalize_installation(
  upgrade = c("ask", "always", "never", "config-only", "data-only"),
  async = FALSE,
  ...
)
```

Arguments

upgrade	rules to upgrade dependencies; default is to ask if needed
async	whether to run in the background; ignore for now
...	ignored; reserved for external calls.

Value

A list built-in pipelines will be installed, the function itself returns nothing.

Examples

```
## Not run:

# This function requires connection to the Github, and must run
# under interactive session since an user prompt will be displayed

ravepipeline_finalize_installation()

## End(Not run)
```

RAVESerializable *Abstract class for 'RAVE' serialization*

Description

For package inheritance only; do not instantiate the class directly.

Methods**Public methods:**

- [RAVESerializable\\$new\(\)](#)
- [RAVESerializable\\$@marshal\(\)](#)
- [RAVESerializable\\$@unmarshal\(\)](#)
- [RAVESerializable\\$@compare\(\)](#)
- [RAVESerializable\\$clone\(\)](#)

Method `new()`: Abstract constructor

Usage:

`RAVESerializable$new(...)`

Arguments:

... ignored

Method `@marshal()`: Create an atomic list that can be serialized

Usage:

```
RAVESerializable$@marshal(...)
```

Arguments:

```
... ignored
```

Method @unmarshal(): Restore an object from an atomic list

Usage:

```
RAVESerializable$@unmarshal(object, ...)
```

Arguments:

```
object a list from '@marshal'
```

```
... ignored
```

Method @compare(): How two object can be compared to each other

Usage:

```
RAVESerializable$@compare(other)
```

Arguments:

```
other another object to compare with self
```

Method clone(): The objects of this class are cloneable with this method.

Usage:

```
RAVESerializable$clone(deep = FALSE)
```

Arguments:

```
deep Whether to make a deep clone.
```

See Also

[RAVEFileArray rave-serialize-refhook](#)

```
rave_progress
```

```
'RAVE' progress
```

Description

Automatically displays 'shiny' progress when shiny is present, or text messages to track progress

Usage

```
rave_progress(
  title,
  max = 1,
  ...,
  quiet = FALSE,
  session = get_shiny_session(),
  shiny_auto_close = FALSE,
  log = NULL
)
```

Arguments

title	progress title
max	maximum steps
...	passed to shiny progress
quiet	whether to suppress the progress
session	shiny session
shiny_auto_close	whether to automatically close the progress bar when the parent function is closed
log	alternative log function if not default (message)

Value

A list of functions to control the progress bar

Examples

```
# Naive example
progress <- rave_progress(title = "progress", max = 10)
progress$inc("job 1")
progress$inc("job 2")
progress$close()

# Within function
slow_sum <- function(n = 11) {
  p <- rave_progress(title = "progress", max = n,
                    shiny_auto_close = TRUE)

  s <- 0
  for( i in seq(1, n) ) {
    Sys.sleep(0.1)
    p$inc(sprintf("adding %d", i))
    s <- s + i
  }
  invisible(s)
}

slow_sum()
```

read-write-yaml *Read write 'YAML' format*

Description

supports reading data into a map object, and write the map to files with names sorted for consistency

Usage

```
load_yaml(file, ..., map = NULL)
```

```
save_yaml(x, file, ..., sorted = FALSE)
```

Arguments

file	file to read from or write to
...	passed to as.list
map	a fastmap object; can be generated by fastmap or dipsaus package; default is to create a new map internally
x	list or map to write
sorted	whether to sort the list by name; default is false

Value

A map object

Examples

```
tfile <- tempfile(fileext = ".yaml")
save_yaml(list(b = 2, a = 1), tfile, sorted = TRUE)
cat(readLines(tfile), sep = "\n")
load_yaml(tfile)
unlink(tfile)
```

with_rave_parallel *Internal parallel functions*

Description

Experimental parallel functions, intended for internal use now. The goal is to allow 'RAVE' functions to gain the potential benefit from parallel computing, but allow users to control whether to do it.

Usage

```
with_rave_parallel(expr, .workers = 0)
```

```
lapply_jobs(
  x,
  fun,
  ...,
  .globals = list(),
  .workers = 0,
  .always = FALSE,
  callback = NULL
)
```

Arguments

expr	expression to evaluate with parallel workers
.workers	number of workers: note the actual numbers may differ, depending on the options and number of input elements
x	a list, vector, array of R objects
fun	function to apply to each element of x
...	additional arguments to be passed to fun
.globals	global variables to be serialized
.always	whether always use workers, only considered when number of workers is one; default is false, then run jobs in the main process when only one worker is required
callback	callback function, input is each element of x and should return a string, for progress bar
workers	number of workers

Details

By default, `lapply_jobs` is almost identical to `lapply`. It only runs in parallel when running inside of `with_rave_parallel`.

The hard max-limit number of workers are determined by the 'RAVE' option `raveio_getopt('max_worker')`. Users can lower this number for memory-intensive tasks manually, via argument `.workers`. The

actual number of workers might be less than the requested ones, this is often a result of sort input `x`. If the number of input iterations has fewer than the max worker size, then the number of workers automatically shrinks to the length of input list. All workers will be a child process running separate from the main session, except for when only one worker is needed and `.always=FALSE`: the only task will be executed in the main session.

Each worker session will run a completely isolated new process. There is a ramp-up serialization that is needed for global objects (objects that are defined elsewhere or outside of the function). Please make sure the global objects are specified explicitly in `.globals`, a named list. Unlike future package, users must specify the global objects.

The global objects might be large to serialize. Please optimize the code to avoid serializing big objects, especially environments or functions. All objects inheriting `RAVESerializable` class with `@marshal` and `@unmarshal` methods implemented correctly will be serialized with reference hook `rave_serialize_refhook`, making them extremely efficient.

Examples

```
# Run without `with_rave_parallel`
res <- lapply_jobs(1:5, function(x, ...) {
  c(child = Sys.getpid(), ...)
}, main = Sys.getpid())

simplify2array(res)

# Comparison
f <- function(n = 5, workers = 0) {
  system.time({
    ravepipeline::lapply_jobs(seq_len(n), function(x, ...) {
      Sys.sleep(1)
      c(child = Sys.getpid(), ...)
    }, main = Sys.getpid(), .workers = workers, callback = I)
  })
}

## Not run:

# Without parallel
f()
#>   user  system elapsed
#> 0.022  0.019  5.010

# with parallel
with_rave_parallel({
  f()
})
#>   user  system elapsed
#> 0.729  0.190  1.460

## End(Not run)
```


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