## Package 'rfacts'

October 14, 2022

Title R Interface to 'FACTS' on Unix-Like Systems

Description The 'rfacts' package is an R interface to the Fixed and Adaptive Clinical Trial Simulator ('FACTS') on Unix-like systems. It programmatically invokes 'FACTS' to run clinical trial simulations, and it aggregates simulation output data into tidy data frames. These capabilities provide end-to-end automation for large-scale simulation pipelines, and they enhance computational reproducibility. For more information on 'FACTS' itself, please visit <https://www.berryconsultants.com/software/>.

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URL https://elilillyco.github.io/rfacts/,
 https://github.com/EliLillyCo/rfacts

BugReports https://github.com/EliLillyCo/rfacts/issues

SystemRequirements FACTS Linux engines (>= 6.2.4), FLFLL (>= 6.2.4), Mono (>= 5.20.1.19)

**Depends** R (>= 3.6.0)

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rfacts-package

rfacts: interface to FACTS on Unix-like systems

## Description

Call FACTS from R.

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts") # example FACTS file
out <- run_facts(
   facts_file,
   n_sims = 4,
   verbose = FALSE
)
```

#### facts\_engines

```
# What results files do we have?
head(get_csv_files(out))
# Read all the "patients*.csv" files with `read_patients(out)`.
# For each scenario, we have files named
# patients00001.csv, patients00002.csv, patients00003.csv,
# and patients00004.csv.
read_patients(out)
}
```

facts\_engines Engine-specific trial simulation functions

## Description

These functions are the inner functions called by run\_engine(). In this help file, only the most common engine functions are listed. To identify the appropriate engine function for your FACTS file, call get\_facts\_engine().

- run\_engine\_aipf\_contin(): Enrichment continuous.
- run\_engine\_aipf\_dichot(): Enrichment dichotomous.
- run\_engine\_aipf\_tte(): Enrichment time to event.
- run\_engine\_contin(): Core continuous.
- run\_engine\_crm(): continual reassessment method (CRM).
- run\_engine\_dichot(): Core dichotomous.
- run\_engine\_multep(): Multiple endpoint.
- run\_engine\_tte(): Time to event.

## Usage

```
run_engine_aipf_contin(
   param_files,
   n_sims = 1L,
   mode = c("", "r"),
   seed = NULL,
   analysis_data = NULL,
   analysis_mode = NULL,
   current_week = NULL,
   current_week = NULL,
   final = NULL,
   interim = NULL,
   interim = NULL,
   verbose = FALSE,
   version = NULL
)
```

run\_engine\_aipf\_dichot(

facts\_engines

```
param_files,
  n_sims = 1L,
 mode = c("", "r"),
  seed = NULL,
  analysis_data = NULL,
  analysis_mode = NULL,
  current_week = NULL,
  execdata = NULL,
  final = NULL,
  interim = NULL,
 mcmc_num = NULL,
 verbose = FALSE,
  version = NULL
)
run_engine_aipf_tte(
  param_files,
  n_sims = 1L,
 mode = c("", "r"),
  seed = NULL,
  analysis_data = NULL,
  analysis_mode = NULL,
  current_week = NULL,
  execdata = NULL,
  final = NULL,
  interim = NULL,
 mcmc_num = NULL,
 verbose = FALSE,
  version = NULL
)
run_engine_contin(
 param_files,
  n_sims = 1L,
 mode = c("s", "r", "p"),
  seed = NULL,
  analysis_data = NULL,
  analysis_mode = NULL,
  arm_selection = NULL,
  armsdropped = NULL,
  complete_data_analysis = NULL,
  current_week = NULL,
  execdata = NULL,
  final = NULL,
  fsimdata = NULL,
  fsimexp = NULL,
  fsimparam = NULL,
  interim = NULL,
```

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```
keepfiles = NULL,
 mcmc_num = NULL,
 noadapt = NULL,
  s2_aux_paramfile = NULL,
  stage = NULL,
 verbose = FALSE,
 version = NULL
)
run_engine_crm(
  param_files,
  n_sims = 1L,
 mode = c("s", ""),
 directory = ".",
  allocator = NULL,
  charting_info = NULL,
  estimator = NULL,
  force_cohort = NULL,
  reduced_priority = NULL,
  version = NULL.
  verbose = FALSE
)
run_engine_dichot(
  param_files,
 n_sims = 1L,
 mode = c("s", "r", "p"),
  seed = NULL,
  analysis_data = NULL,
  analysis_mode = NULL,
  arm_selection = NULL,
  armsdropped = NULL,
  complete_data_analysis = NULL,
  current_week = NULL,
  execdata = NULL,
  final = NULL,
  fsimdata = NULL,
  fsimexp = NULL,
  fsimparam = NULL,
  interim = NULL,
  keepfiles = NULL,
 mcmc_num = NULL,
  noadapt = NULL,
  s2_aux_paramfile = NULL,
  stage = NULL,
  verbose = FALSE,
  version = NULL
)
```

```
run_engine_multep(
  param_files,
  n_sims = 1L,
 mode = c("s", "r", "p"),
  seed = NULL,
  analysis_data = NULL,
  analysis_mode = NULL,
  arm_selection = NULL,
  armsdropped = NULL,
  complete_data_analysis = NULL,
  current_week = NULL,
  execdata = NULL,
  final = NULL,
  fsimdata = NULL,
  fsimexp = NULL,
  fsimparam = NULL,
  interim = NULL,
  keepfiles = NULL,
 mcmc_num = NULL,
 noadapt = NULL,
  s2_aux_paramfile = NULL,
  stage = NULL,
  verbose = FALSE,
  version = NULL
)
run_engine_tte(
  param_files,
  n_sims = 1L,
 mode = c("s", "r", "p"),
  seed = NULL,
  analysis_data = NULL,
  analysis_mode = NULL,
  arm_selection = NULL,
  armsdropped = NULL,
  complete_data_analysis = NULL,
  current_week = NULL,
  execdata = NULL,
  final = NULL,
  fsimdata = NULL,
  fsimexp = NULL,
  fsimparam = NULL,
  interim = NULL,
  keepfiles = NULL,
 mcmc_num = NULL,
  noadapt = NULL,
  s2_aux_paramfile = NULL,
```

## facts\_engines

```
stage = NULL,
verbose = FALSE,
version = NULL
)
```

## Arguments

| param_files                    | Character vector of file paths or the output of prep_param_files(). If a character vector, the elements can be directories containing *.param files or the paths to the *.param files themselves. Such a directory is returned by run_flfll().  |  |
|--------------------------------|---|--|
| n_sims                         | Positive integer, number of simulations per param file.   |  |
| mode                           | Character scalar: "s" for simulation mode in non-enrichment designs, "" for simulation mode in enrichment designs, "r" for execution mode, and "p" for prediction mode. For the CRM engine, mode needs to be "s" or "".   |  |
| seed                           | Positive integer, random number generator seed for the actual trial simulations.<br>Use this seed argument instead of flfll_seed (run_facts(), run_flfll())<br>to control pseudo-randomness in the actual trial simulations. flfll_seed only<br>controls how the *.param files are generated.   |  |
| analysis_data                  | Character, analysis mode patient data file name.  |  |
| analysis_mode                  | Logical, whether to activate analysis mode.   |  |
| current_week                   | Numeric, current time in weeks.   |  |
| execdata                       | Character, name of the execution mode patient file.   |  |
| final                          | Logical, whether to do the final analysis. For execution mode only.   |  |
| interim                        | Integer, interim number.  |  |
| mcmc_num                       | Integer, MCMC file number. For analysis mode only.  |  |
| verbose                        | Logical, whether to print progress information to the R console.  |  |
| version                        | Character scalar, version of FACTS corresponding to the FACTS file. Get by calling get_facts_version() on your FACTS file. See possible versions with get_facts_versions(). Do not supply version to run_engine(). run_engine() detects the version automatically from the FACTS file and passes it to the appropriate engine function. |  |
| arm_selection                  | Logical, whether to activate arm selection.   |  |
| armsdropped<br>complete_data_a | Character, a comma-separated collection of integers indicating dropped arms.<br>analysis  |  |
|                                | Logical, whether to do a complete data analysis.  |  |
| fsimdata                       | Character, prediction mode patient data file name.  |  |
| fsimexp                        | Logical. For expert use only.   |  |
| fsimparam                      | Character, name of the prediction mode *.param file.  |  |
| keepfiles                      | Logical, whether to deactivate cleanup of extraneous staged design files.   |  |
| noadapt                        | Logical, whether to deactivate adaptive actions in prediction mode.   |  |
| s2_aux_paramfile               |   |  |
|                                | Character, name of the stage 2 execution auxiliary *.param file.  |  |

| stage            | Integer, trial design stage. For staged designs only.           |  |
|------------------|---|--|
| directory        | Character, working directory. CRM only.                         |  |
| allocator        | Logical, allocator/execution/recommender mode. CRM only.        |  |
| charting_info    | Logical, unused.  |  |
| estimator        | Logical, use estimator. CRM only.                               |  |
| force_cohort     | Logical, whether to force small cohort run-in to end. CRM only. |  |
| reduced_priority |   |  |
|                  | Logical, whether to run at reduced priority. CRM only.          |  |

## Details

If you need to repeatedly invoke an engine, as with most trial execution mode workflows, these engine functions may be slow on their own. To avoid the most severe sources of slowness, consider running prep\_param\_files() and then passing the result to one of the individual engine functions (such as run\_engine\_contin()).

#### Value

Nothing.

## See Also

run\_engine(),get\_facts\_file\_example(),get\_facts\_engine(),run\_facts(),run\_flfll().

## Examples

```
facts_file <- get_facts_file_example("contin.facts")
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
  out <- run_flfll(facts_file, verbose = FALSE) # Generate param files.
  # Identify which engine you need.
  get_facts_engine(facts_file)
  # Run the sims with the engine function or `run_engine()`.
  run_engine_contin(out, n_sims = 1, verbose = FALSE, version = "6.2.5")
  read_patients(out)
}</pre>
```

facts\_results k

#### facts\_results

#### Description

These functions read trial simulation results. The results were computed by FACTS (via run\_facts() or run\_engine() or one of the engine functions such as run\_engine\_contin()) and are stored in CSV files. Different functions read different types of output. The functions are named according to the CSV files they read. For example, read\_patients() reads all files named patients00001.csv, patients00002.csv, etc. The most important functions are read\_patients() and read\_weeks(). The read\_s1\*(), read\_s2\*(), and read\_master\*() functions are for staged designs. The read\_csv\_special() function allows you to supply a custom file name prefix such as "patients", but be warned: not every kind of CSV output file is tested in rfacts.

#### Usage

```
read_patients(csv_files)
```

read\_weeks(csv\_files)

read\_mcmc(csv\_files)

read\_s1\_mcmc(csv\_files)

read\_s1\_weeks(csv\_files)

read\_s1\_patients(csv\_files)

read\_s2\_patients(csv\_files)

read\_s2\_weeks(csv\_files)

read\_s2\_mcmc(csv\_files)

read\_master\_mcmc(csv\_files)

read\_master\_patients(csv\_files)

read\_master\_weeks(csv\_files)

read\_cohorts(csv\_files)

read\_simulations(csv\_files)

read\_csv\_special(csv\_files, prefix, numbered = TRUE)

#### Arguments

| csv_files | Character vector of file paths. Either the directories containing the trial simula-<br>tion results or the actual CSV file files themselves. |
|-----------|--|
| prefix    | Character, name of the prefix for read csy special(), read weeks(x) is   |

equivalent to read\_csv\_special(x, prefix = "weeks"). Be careful: not all

kinds of CSV output are tested. We can only guarantee the file types with special functions will be read correctly, e.g. read\_patients() and read\_weeks(). Logical. If TRUE, only read the numbered files like patients00001.csv, weeks00017.csv, etc. If FALSE, only list the non-numbered files like simulations.csv and simulations\_freq\_locf.csv. Avoid summary.csv files. They are not reliable on Linux.

## Value

A data frame of trial simulation data. Each read\_\*() function returns different information, but all the read\_\*() functions support the following columns:

- facts\_file: character, the base name of the FACTS file.
- facts\_scenario: character, the name of the simulation scenario from FACTS. Usually, this factors in the virtual subject response (VSR) profile, accrual profile (how fast do patients enroll?) and dropout profile (how fast do they drop out?).
- facts\_sim: integer, numeric index of the CSV file name. For example, the facts\_sim of patients00012.csv is 12. In trial execution mode, all these indices are 00000, so facts\_id is much safer than facts\_sim for packetized trial execution mode.
- facts\_id: character, random unique id of each CSV file being read. Different for every call to read\_patients() etc. Safer than facts\_sim for aggregation over simulations.
- facts\_output: character, type of output is in the data frame: "patients" for patients files, "weeks" for weeks files, "mcmc" for MCMC files, etc. These names adhere to established conventions in FACTS.
- facts\_csv: character, full path to the original CSV files where FACTS stored the simulation output. Required for overwrite\_csv\_files().
- facts\_header: a character vector of \n-delimited CSV file headers. Required for overwrite\_csv\_files().

## See Also

get\_facts\_file\_example(), run\_facts(), run\_flfll(), run\_engine()

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts")</pre>
out <- run_facts(</pre>
 facts_file,
 n_sims = 4,
 verbose = FALSE
)
# What results files do we have?
head(get_csv_files(out))
# Read all the "patients*.csv" files with `read_patients(out)`.
# For each scenario, we have files named
# patients00001.csv, patients00002.csv, patients00003.csv,
# and patients00004.csv.
read_patients(out)
}
```

get\_csv\_files

## Description

List output CSV files in a directory or directories.

## Usage

```
get_csv_files(csv_files, numbered = TRUE)
```

## Arguments

| csv_files | Character vector of directories containing numbered CSV files   |
|-----------|---|
| numbered  | Logical. If TRUE, only list the numbered files like patients00001.csv, weeks00017.csv, etc. If FALSE, only list the non-numbered files like simulations.csv and simulations_freq_locf.csv. Avoid summary.csv files. They are not reliable on Linux. |

## Value

A character vector of names of CSV files.

## Examples

```
facts_file <- get_facts_file_example("contin.facts")
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
  out <- run_facts(
    facts_file,
    n_sims = 2L,
    verbose = FALSE
)
get_csv_files(out)
}</pre>
```

get\_facts\_engine Get the FACTS engine function matching your FACTS file

## Description

Identify the correct run\_engine\_\*() function for your FACTS file.

### Usage

get\_facts\_engine(facts\_file)

#### Arguments

facts\_file Character, name of a FACTS file. Usually has a \*. facts file extension.

## Details

For most cases, it is sufficient to call run\_facts(), or to call run\_flfll() followed by run\_engine(). But either way, you will need to know the arguments of the run\_engine\_\*() function that corresponds to your FACTS file. Even if you are not calling this run\_engine\_\*() directly, you will need to pass the arguments to ... in run\_facts() or run\_engine(). get\_facts\_engine() identifies the correct run\_engine\_\*() function so you can open the help file and read about the arguments, e.g. ?run\_engine\_contin.

## Value

Character, the name of a FACTS engine function.

#### See Also

run\_facts(), run\_engine()

#### Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts")</pre>
out <- run_flfll(facts_file, verbose = FALSE) # Generate param files.</pre>
# Find the appropriate FACTS engine function.
get_facts_engine(facts_file)
# Read about the function arguments.
# You can pass these arguments to `...` in `run_facts()`
# or `run_engine()` or just call `run_engine_contin()` directly.
# ?run_engine_contin
# Call the FACTS engine function to run simulations.
# Alternatively, you could just call `run_engine()`.
run_engine_contin(out, n_sims = 1, verbose = FALSE, version = "6.2.5")
# See the results.
read_patients(out)
}
```

get\_facts\_file\_example

Locate an example FACTS file

#### Description

Get the path to an example FACTS file inside rfacts itself.

#### Usage

```
get_facts_file_example(facts_file)
```

## Arguments

facts\_file Character, name of a FACTS file. Usually has a \*.facts file extension. Does not include the directory name. Possible choices:

- "aipf\_contin.facts" Enrichment continuous.
- "aipf\_dichot.facts" Enrichment dichotomous.
- "aipf\_tte.facts" Enrichment time to event.
- "broken.facts" A broken FACTS file.
- "contin.facts" Core continuous.
- "crm.facts" N-CRM design.
- "dichot.facts" Core dichotomous.
- "multep.facts" Multiple endpoints.
- "staged.facts" Staged design.
- "tte.facts" Time to event.
- "unsupported.facts" FACTS file with an unsupported engine type.

#### Details

The rfacts package comes with some example FACTS files. Use the get\_facts\_file\_example() function to get the full path to an example FACTS file. Use this file to try out run\_flfll(), run\_engine\_contin(), etc.

#### Value

Character, the path to a FACTS file included with rfacts.

## See Also

run\_facts(), run\_flfll(), run\_engine(), run\_engine\_contin()

```
# Only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts")
facts_file
out <- run_facts(
  facts_file,
   n_sims = 1,
   verbose = FALSE
)
read_patients(out)
}
```

get\_facts\_scenarios List the names of simulation scenarios

## Description

Get the names of the simulation scenarios of a FACTS file. without actually running any simulations. These names usually come from the virtual subject response (VSR) scenarios, the accrual profiles, and the dropout profiles.

## Usage

```
get_facts_scenarios(facts_file, verbose = FALSE)
```

#### Arguments

| facts_file | Character, name of a FACTS file. Usually has a *.facts file extension. |
|------------|--|
| verbose    | Logical, whether to print progress to the R console.                   |

## Value

Character vector of FACTS simulation scenarios.

#### See Also

get\_param\_dirs(), run\_facts(), run\_flfll(), run\_engine(), run\_engine\_contin()

## Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts")
get_facts_scenarios(facts_file)
}
```

get\_facts\_version Get FACTS version matching your FACTS file

## Description

Get the version of FACTS compatible with your \*.facts file.

## Usage

get\_facts\_version(facts\_file)

#### Arguments

facts\_file Character, name of a FACTS file. Usually has a \*. facts file extension.

## Value

A version string.

## See Also

get\_facts\_versions()

## Examples

```
facts_file <- get_facts_file_example("contin.facts")
facts_file
get_facts_version(facts_file)</pre>
```

get\_facts\_versions List supported FACTS versions

## Description

List versions of FACTS supported by rfacts. You can supply any of these versions to functions engine-specific functions such as run\_engine\_contin().

## Usage

get\_facts\_versions()

#### Details

If your FACTS file does not perfectly agree with one of the supported versions, rfacts will try to find the best version for you, either

- 1. The greatest supported version less than or equal to the one in the FACTS file, or
- 2. The lowest supported version if (1) does not exist.

## Value

A character vector of supported FACTS versions.

### See Also

get\_facts\_version(), run\_engine\_contin()

## Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
get_facts_versions()
}
```

get\_param\_dirs List the directories containing param files

#### Description

Get the directory paths containing param files. This helps us run FACTS simulation scenarios one at a time.

### Usage

```
get_param_dirs(param_files)
```

## Arguments

| param_files | Character, path to a top-level directory containing param files. run_flfll() and       |
|-------------|--|
|             | <pre>run_facts() return paths you can supply to param_files in get_param_dirs().</pre> |

#### Details

When you run run\_flfll() or run\_facts(), rfacts creates a directory. This directory has a bunch of subdirectories, each corresponding to a single simulation scenario (VSR profile x accrual profile x dropout profile, etc).

## Value

Character vector of FACTS simulation scenario directories.

#### See Also

get\_facts\_scenarios(), run\_facts(), run\_flfll(), run\_engine(), run\_engine\_contin()

#### Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts")
param_files <- run_flfll(facts_file, verbose = FALSE)
scenarios <- get_param_dirs(param_files)
scenarios
scenario <- scenarios[1]
run_engine_contin(scenario, n_sims = 2, verbose = FALSE, version = "6.2.5")
read_patients(scenario)
}
```

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### Description

List the paths to the all the param files in a directory or directories.

#### Usage

```
get_param_files(param_files)
```

## Arguments

param\_files Character vector of directories containing param files.

## Value

Character vector of paths to param files.

#### Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts")
dir <- run_flfll(facts_file, verbose = FALSE)
get_param_files(dir)
}
```

overwrite\_csv\_files Overwrite FACTS CSV output files

## Description

read\_patients() and friends read CSV output files from FACTS and return special aggregated data frames. overwite\_csv\_files() accepts such an aggregated data frame and writes the content to the original CSV files it came from.

## Usage

```
overwrite_csv_files(x)
```

### Arguments

Х

An aggregated data frame from read\_patients() or similar function.

#### Value

Nothing.

#### Examples

```
facts_file <- get_facts_file_example("contin.facts")
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
  out <- run_facts(facts_file, n_sims = 2)
    pats <- read_patients(out)
    head(pats$visit_1)
    pats$visit_1 <- 0
    overwrite_csv_files(pats)
    pats2 <- read_patients(out)
    head(pats2$visit_1)
}</pre>
```

prep\_param\_files Arrange the param files for the engines ahead of time.

#### Description

If you call prep\_param\_files() ahead of time, subsequent calls to the engines will initialize much faster. This is useful in situations like trial execution mode, which require calling an engine function on each new simulation. This function does not actually modify the param files themselves on disk.

#### Usage

```
prep_param_files(param_files)
```

#### Arguments

param\_files A character vector of param files and/or directories containing param files.

## Details

prep\_param\_files() searches for the required \*.param files groups them by directory, sorts them, and returns the result as a list of special param\_files objects. (It does not modify the actual contents of the \*.param`` files.) This preprocessing step is fast when executed once, but slow when executed the second state of the second state

## Value

A list of special "params\_files" objects that the engine functions can process fast.

## See Also

run\_flfll(), run\_engine(), run\_engine\_contin()

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## read\_facts

## Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts")
out <- run_flfll(facts_file, verbose = FALSE)
param_files <- prep_param_files(out) # For speed.
param_files # Shows where the param files live and how they are organized.
run_engine_contin(
    param_files,
    n_sims = 2,
    verbose = FALSE,
    version = "6.2.5"
)
# Slower: run_engine_contin(out, n_sims = 2, verbose = FALSE) # nolint
}
```

read\_facts

Read parts of FACTS files.

## Description

Read specific fields of a FACTS file.

#### Usage

```
read_facts(facts_file, fields)
```

#### Arguments

| facts_file | Character of length 1, path to FACTS XML file to read.  |
|------------|---|
| fields     | Data frame defining the kind of XML data to be read. It must have one row per field definition and the following columns: |
|            | 1. field: custom name of the field.   |
|            | 2. type: value of the "type" attribute of the <parametersets> tag.</parametersets>  |
|            | 3. set: value of the "name" attribute of the <parameterset> tag.</parameterset>   |
|            | 4. property: value of the "name" attribute of the <property> tag.</property>  |

#### Details

A FACTS file has a special kind of XML format. Most of the content sits in an overarching <facts> tag, then a <parameterSets> tag, then a <parameterSet> tag, then a <property> tag. For example, here is the part of a FACTS file that controls the weeks between interims.

#### <facts>

```
<parameterSets type="NucleusParameterSet">
  <parameterSet name="nucleus">
   <property name="update_freq_save">4</property></property>
```

To use the read\_facts() function, you must first identify the parts of the FACTS file you want to read using the fields argument. To read the above part of the XML, you would first define the update\_freq\_save field.

```
fields <- tibble::tibble(
   field = "my_interval",
   type = "NucleusParameterSet",
   set = "nucleus",
   property = "update_freq_save"
)</pre>
```

and then call read\_facts(input = "your\_file.facts", fields = fields).

#### Value

A one-row tibble with the requested fields from the FACTS file.

#### Examples

```
facts_file <- get_facts_file_example("contin.facts")
fields <- data.frame(
    field = c("my_subjects", "my_vsr"),
    type = c("NucleusParameterSet", "EfficacyParameterSet"),
    set = c("nucleus", "resp2"),
    property = c("max_subjects", "true_endpoint_response")
)
read_facts(facts_file = facts_file, fields = fields)</pre>
```

reset\_rfacts\_paths Reset system dependency info

#### Description

Reset system dependency information based on the current value of the RFACTS\_PATHS environment variable.

#### Usage

```
reset_rfacts_paths()
```

#### Dependencies

rfacts has strict system requirements, and the installations vary from system to system. You need to specify the locations of system executables in a CSV file that lists the path and metadata of each executable. This file must have one row per executable and the following columns.

• executable\_type: Must be "mono", "flfll", or "engine" to denote the general type of the executable.

- facts\_version: The version of FACTS with which this executable is compatible.
- path: File path to the executable.
- engine\_name: For engines only. Name of the engine. Must be one of the engine types in the example CSV file at system.file("example\_paths.csv", package = "rfacts").
- param\_set: For engines only. Parameter set designation listed in the XML code of FACTS files for that engine. See system.file("example\_paths.csv", package = "rfacts") for examples.
- param\_type: For engines only. Parameter type designation listed in the XML code of FACTS files for that engine. See system.file("example\_paths.csv", package = "rfacts") for examples.

When you call a trial simulation function in rfacts, the package automatically reads this file and memorizes the contents for later use. The file at system.file("example\_paths.csv", package = "rfacts") (inst/example\_paths.csv in the package source.) has an example of such a file. All the columns in that file are required, and you may, remove, or modify rows to fit your specific system.

To enable rfacts to find this CSV file, you need to set the RFACTS\_PATHS environment variable to the path to this file. The easiest way to do this is call usethis::edit\_r\_environ() to edit your .Renviron file and then add a new line with something like RFACTS\_PATHS=/path/to/file/paths.csv. Then, restart your R session and call Sys.getenv("RFACTS\_PATHS") to verify that this environment variable was set correctly.

The rfacts\_sitrep() function inspects the current system dependency info and ensures each executable exists and has the correct permissions.

If you change the RFACTS\_PATHS environment variable, you need to call reset\_rfacts\_paths() or restart R for the changes to take effect.

#### See Also

rfacts\_paths, rfacts\_sitrep

#### Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
  reset_rfacts_paths()
}
```

rfacts\_paths Read paths to rfacts system dependencies

## Description

Read the file specified by the RFACTS\_PATHS environment variable.

#### Usage

rfacts\_paths()

A data frame with paths and other metadata about rfacts system dependencies

#### Dependencies

rfacts has strict system requirements, and the installations vary from system to system. You need to specify the locations of system executables in a CSV file that lists the path and metadata of each executable. This file must have one row per executable and the following columns.

- executable\_type: Must be "mono", "flfll", or "engine" to denote the general type of the executable.
- facts\_version: The version of FACTS with which this executable is compatible.
- path: File path to the executable.
- engine\_name: For engines only. Name of the engine. Must be one of the engine types in the example CSV file at system.file("example\_paths.csv", package = "rfacts").
- param\_set: For engines only. Parameter set designation listed in the XML code of FACTS files for that engine. See system.file("example\_paths.csv", package = "rfacts") for examples.
- param\_type: For engines only. Parameter type designation listed in the XML code of FACTS files for that engine. See system.file("example\_paths.csv", package = "rfacts") for examples.

When you call a trial simulation function in rfacts, the package automatically reads this file and memorizes the contents for later use. The file at system.file("example\_paths.csv", package = "rfacts") (inst/example\_paths.csv in the package source.) has an example of such a file. All the columns in that file are required, and you may, remove, or modify rows to fit your specific system.

To enable rfacts to find this CSV file, you need to set the RFACTS\_PATHS environment variable to the path to this file. The easiest way to do this is call usethis::edit\_r\_environ() to edit your .Renviron file and then add a new line with something like RFACTS\_PATHS=/path/to/file/paths.csv. Then, restart your R session and call Sys.getenv("RFACTS\_PATHS") to verify that this environment variable was set correctly.

The rfacts\_sitrep() function inspects the current system dependency info and ensures each executable exists and has the correct permissions.

If you change the RFACTS\_PATHS environment variable, you need to call reset\_rfacts\_paths() or restart R for the changes to take effect.

#### See Also

rfacts\_sitrep

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
rfacts_paths()
}
```

rfacts\_sitrep

#### Description

Examine the file paths to executables and check that they exist and have the correct permissions.

#### Usage

```
rfacts_sitrep()
```

#### Value

A data frame of information on the status of each executable.

#### Dependencies

rfacts has strict system requirements, and the installations vary from system to system. You need to specify the locations of system executables in a CSV file that lists the path and metadata of each executable. This file must have one row per executable and the following columns.

- executable\_type: Must be "mono", "fifil", or "engine" to denote the general type of the executable.
- facts\_version: The version of FACTS with which this executable is compatible.
- path: File path to the executable.
- engine\_name: For engines only. Name of the engine. Must be one of the engine types in the example CSV file at system.file("example\_paths.csv", package = "rfacts").
- param\_set: For engines only. Parameter set designation listed in the XML code of FACTS files for that engine. See system.file("example\_paths.csv", package = "rfacts") for examples.
- param\_type: For engines only. Parameter type designation listed in the XML code of FACTS files for that engine. See system.file("example\_paths.csv", package = "rfacts") for examples.

When you call a trial simulation function in rfacts, the package automatically reads this file and memorizes the contents for later use. The file at system.file("example\_paths.csv", package = "rfacts") (inst/example\_paths.csv in the package source.) has an example of such a file. All the columns in that file are required, and you may, remove, or modify rows to fit your specific system.

To enable rfacts to find this CSV file, you need to set the RFACTS\_PATHS environment variable to the path to this file. The easiest way to do this is call usethis::edit\_r\_environ() to edit your .Renviron file and then add a new line with something like RFACTS\_PATHS=/path/to/file/paths.csv. Then, restart your R session and call Sys.getenv("RFACTS\_PATHS") to verify that this environment variable was set correctly.

The rfacts\_sitrep() function inspects the current system dependency info and ensures each executable exists and has the correct permissions. If you change the RFACTS\_PATHS environment variable, you need to call reset\_rfacts\_paths() or restart R for the changes to take effect.

#### See Also

rfacts\_paths

## Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
rfacts_sitrep()
}
```

run\_engine

Run trial simulations

## Description

For fine control over trial simulations, you must first call run\_flfll() and then call either run\_engine() or one of the specific engine functions (such as run\_engine\_contin()). The engines read the \*.param files generated by run\_flfll(), run the trial simulations, and save output to a bunch of CSV files. You can find these CSV output files next to the \*.param files.

#### Usage

```
run_engine(facts_file, ...)
```

## Arguments

| facts_file | Character, name of a FACTS file. Usually has a *.facts file extension.                 |
|------------|--|
|            | Named arguments to the appropriate inner engine function, such as run_engine_contin(). |
|            | Use get_facts_engine() to identify the appropriate engine function for your            |
|            | FACTS file. Then, open the help file of that function to read about the argu-          |
|            | ments.   |

## Details

If you need to repeatedly invoke an engine, as with most trial execution mode workflows, run\_engine() is slow. Instead, consider running prep\_param\_files() and then passing the result to one of the individual engine functions (such as run\_engine\_contin()).

## Value

Nothing.

## See Also

```
get_facts_file_example(), run_flfll(), get_facts_engine(), prep_param_files()
```

## run\_facts

## Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts") # example FACTS file
out <- run_flfll(facts_file, verbose = FALSE) # Generate param files.
# Run the simulations.
run_engine(
    facts_file,
    param_files = out,
    n_sims = 1,
    verbose = FALSE
)
read_patients(out)
}
```

run\_facts

Run FACTS

## Description

Run FACTS trial simulations.

## Usage

```
run_facts(
   facts_file,
   output_path = tempfile(),
   log_path = output_path,
   n_burn = NULL,
   n_mcmc = NULL,
   n_weeks_files = 10000,
   n_patients_files = 10000,
   n_mcmc_files = 0,
   n_mcmc_thin = NULL,
   flfll_seed = NULL,
   flfll_offset = NULL,
   n_sims,
   ...
)
```

#### Arguments

| facts_file  | Character, name of a FACTS file. Usually has a *.facts file extension. |
|-------------|--|
| output_path | Character, directory path to the files to generate.                    |
| log_path    | Character, path to the log file generated by FLFLL.                    |
| n_burn      | Number of burn-in iterations for the MCMC.                             |
| n_mcmc      | Number of MCMC iterations used in inference.                           |

| n_weeks_files   | Number of weeks*.csv files to save in output_path.   |
|-----------------|--|
| n_patients_file | es   |
|                 | Number of patients*.csv files to save in output_path.  |
| n_mcmc_files    | Number of mcmc*.csv files to save in output_path.  |
| n_mcmc_thin     | Number of thinning iterations for the MCMC.  |
| flfll_seed      | Positive integer, random number generator seed for FLFLL. This seed is only<br>used for stochastic preprocessing steps for generating the *.param files. It is<br>not the random number generator seed for the actual trial simulations. To set the<br>trial simulation seed, use the seed argument of run_facts(), run_engine(),<br>or one of the specific run_engine*() functions. |
| flfll_offset    | Integer, offset for the random number generator.   |
| n_sims          | Positive integer, number of simulations per param file.  |
|                 | Named arguments to the appropriate FACTS engine function. Use get_facts_engine() to identify the appropriate engine function and then open the help file of that function to read about the arguments, e.g. ?run_engine_contin.  |

## Details

run\_facts() calls run\_flfll() and then run\_engine(). For finer control over trial simulation, you can call these latter two functions individually.

## Value

Character, path to the directory with FACTS output.

## See Also

run\_flfll(), run\_engine(), get\_facts\_engine()

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts") # example FACTS file</pre>
out <- run_facts(</pre>
  facts_file,
 n_sims = 4,
  verbose = FALSE
)
# What results files do we have?
head(get_csv_files(out))
# Read all the "patients*.csv" files with `read_patients(out)`.
# For each scenario, we have files named
# patients00001.csv, patients00002.csv, patients00003.csv,
# and patients00004.csv.
read_patients(out)
}
```

run\_flfll

## Description

Generate the preparatory files required for simulation.

## Usage

```
run_flfll(
  facts_file,
  output_path = tempfile(),
  log_path = output_path,
  n_burn = NULL,
  n_mcmc = NULL,
  n_weeks_files = 10000,
  n_patients_files = 10000,
  n_mcmc_files = 0,
  n_mcmc_thin = NULL,
  flfll_seed = NULL,
  flfll_offset = NULL,
  verbose = FALSE,
  max_sims = 99999L
)
```

## Arguments

| facts_file      | Character, name of a FACTS file. Usually has a *.facts file extension.   |
|-----------------|--|
| output_path     | Character, directory path to the files to generate.  |
| log_path        | Character, path to the log file generated by FLFLL.  |
| n_burn          | Number of burn-in iterations for the MCMC.   |
| n_mcmc          | Number of MCMC iterations used in inference.   |
| n_weeks_files   | Number of weeks*.csv files to save in output_path.   |
| n_patients_file | es   |
|                 | Number of patients*.csv files to save in output_path.  |
| n_mcmc_files    | Number of mcmc*.csv files to save in output_path.  |
| n_mcmc_thin     | Number of thinning iterations for the MCMC.  |
| flfll_seed      | Positive integer, random number generator seed for FLFLL. This seed is only used for stochastic preprocessing steps for generating the *.param files. It is not the random number generator seed for the actual trial simulations. To set the trial simulation seed, use the seed argument of run_facts(), run_engine(), or one of the specific run_engine*() functions. |
| flfll_offset    | Integer, offset for the random number generator.   |
| verbose         | Logical, whether to print progress information to the R console.   |

max\_sims Positive integer of length 1, maximum number of simulations that will be allowed to run for certain engines like CRM in subsequent calls to the engine. If the n\_sims argument of the engine is larger than max\_sims, only max\_sims simulations will be run. The max\_sims argument only applies to FLFLL >= 6.4.1 and only needs to be set manually if you are manually calling run\_flfll() and then the engine instead of just run\_facts().

## Details

For advanced control over trial simulations, you must first call run\_flfll() and then call one of the engine functions such as run\_engine\_contin(). run\_flfll() generates the preparatory \*.param files that the run\_engine\_\*() functions understand. You will pass these \*.param files or their parent directory to param\_files argument of run\_engine\_contin() etc.

#### Value

Character, the value of output\_path. output\_path is the directory path to the files generated by run\_flfll().

#### See Also

get\_facts\_file\_example(), run\_engine(), run\_engine\_contin()

#### Examples

```
# Can only run if system dependencies are configured:
if (file.exists(Sys.getenv("RFACTS_PATHS"))) {
facts_file <- get_facts_file_example("contin.facts") # example FACTS file
out <- run_flfll(facts_file, verbose = FALSE) # Generate param files.
# Run the simulations.
run_engine(
    facts_file,
    param_files = out,
    n_sims = 1,
    verbose = FALSE
)
read_patients(out)
}
```

write\_facts

Write modified FACTS files.

#### Description

Write modified versions of existing FACTS files. This function can be used to tweak properties of a FACTS file such as maximum sample size, number of weeks between interims, allocation ratios, data generation parameters, and analysis priors.

#### write\_facts

#### Usage

write\_facts(fields, values, default\_dir = "\_facts")

#### Arguments

| fields      | Data frame defining the kind of XML data to be replaced. It must have one row per field definition and the following columns:   |
|-------------|---|
|             | 1. field: custom name of the field.   |
|             | 2. type: value of the "type" attribute of the <parametersets> tag.</parametersets>  |
|             | 3. set: value of the "name" attribute of the <parameterset> tag.</parameterset>   |
|             | 4. property: value of the "name" attribute of the <property> tag.</property>  |
| values      | Data frame defining the FACTS files to generate. Must have one row per FACTS file and a column called facts_file with the names of the input FACTS files. An output column with the names of the output FACTS files is recommended but not required. (If output is not specified, the output FACTS files will be written to automatically generated paths inside default_dir.) Other columns must have names corresponding to elements of fields\$field and contain values to insert into the FACTS files. These columns could be vectors or lists of vectors. In the former case, each element is a scalar replacement to a property. In the latter case, an XML property receives an entire vector as an item list, and the vector must be the same length as the original item list. |
| default_dir | Directory to write the output FACTS files if values has no output column.   |

## Details

A FACTS file has a special kind of XML format. Most of the content sits in an overarching <facts> tag, then a <parameterSet> tag, then a <parameterSet> tag, then a <property> tag. For example, here is the part of a FACTS file that controls the weeks between interims.

#### <facts>

```
<parameterSets type="NucleusParameterSet">
   <parameterSet name="nucleus">
    <property name="update_freq_save">4</property></property></property>
```

To use the write\_facts() function, you must first identify the parts of the FACTS file you want to modify (the fields argument) then the values that should be substituted in (the values argument). Given the XML above, to create new FACTS files with intervals 5 and 6 instead of 4, you would set

```
fields <- tibble::tibble(
   field = "my_interval",
   type = "NucleusParameterSet",
   set = "nucleus",
   property = "update_freq_save"
)
values <- tibble::tibble(
   facts_file = "your_facts_file.facts",
   output = "output_file.facts",</pre>
```

```
my_interval = c(5, 6)
```

and then call write\_facts(fields = fields, values = values).

#### Value

The function writes FACTS XML files and returns a character vector with the paths to those files.

```
# Identify a source FACTS file.
facts_file <- get_facts_file_example("contin.facts")</pre>
# Create 4 new FACTS files with different numbers of max patients.
fields <- data.frame(</pre>
 field = "my_subjects",
 type = "NucleusParameterSet",
  set = "nucleus",
  property = "max_subjects"
)
values <- data.frame(</pre>
  facts_file = facts_file,
  output = c("_facts/out1000.facts", "_facts/out2000.facts"),
 my_subjects = c(1000, 2000)
)
default_dir <- tempfile()</pre>
write_facts(fields = fields, values = values, default_dir = default_dir)
list.files("_facts")
unlink("_facts", recursive = TRUE)
```

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