Package 'tern.rbmi'

July 21, 2025

Title Create Interface for 'RBMI' and 'tern'

Version 0.1.6

Date 2025-07-18

Description 'RBMI' implements standard and reference based multiple
imputation methods for continuous longitudinal endpoints (Gower-Page
et al. (2022) <doi:10.21105/joss.04251>). This package provides an
interface for 'RBMI' uses the 'tern'
<https://cran.r-project.org/package=tern> framework by Zhu et al.
(2023) and tabulate results easily using 'rtables'
<https://cran.r-project.org/package=rtables> by Becker et al. (2023).

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URL https://github.com/insightsengineering/tern.rbmi,
https://insightsengineering.github.io/tern.rbmi/

BugReports https://github.com/insightsengineering/tern.rbmi/issues

Depends R (>= 3.6), rbmi (>= 1.2.5), tern (>= 0.9.7)

Imports broom (>= 0.5.4), checkmate (>= 2.1.0), formatters (>=
0.5.10), lifecycle (>= 0.2.0), magrittr (>= 1.5), rtables (>=
0.6.11)

Suggests BH, dplyr (>= 1.0.3), knitr (>= 1.42), Matrix, RcppEigen,
rmarkdown (>= 2.23), rstan, testthat (>= 3.0.4), tidyr (>=
0.8.3), V8

VignetteBuilder knitr, rmarkdown

Config/Needs/verdepcheck insightsengineering/rbmi,
insightsengineering/tern, tidymodels/broom, mllg/checkmate,
insightsengineering/formatters, r-lib/lifecycle,
tidyverse/magrittr, insightsengineering/rtables,
tidyverse/dplyr, yihui/knitr, rstudio/rmarkdown,
r-lib/testthat, tidyverse/tidyr

Config/Needs/website insightsengineering/nesttemplate

Config/testthat/edition 3

Encoding UTF-8

Language en-US

LazyData true

RoxygenNote 7.3.2

Collate 'tabulate_rbmi.R' 'tern.rbmi-package.R'

NeedsCompilation no

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Contents

a_rbmi_lsmeans	*Formatted Analysis function which can be further customized by calling* [rtables::make_afun()] *on it. It is used as* afun *in* [rtables::analyze()].

Description

[Experimental]

Usage

```
a_rbmi_lsmeans(df, .in_ref_col, show_relative = c("reduction", "increase"))
```

Arguments

df	input dataframe
.in_ref_col	boolean variable, if reference column is specified
show_relative	"reduction" if (control - treatment, default) or "increase" (treatment - control) of relative change from baseline?

Value

Formatted Analysis function

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h_tidy_pool	*Helper function to produce data frame with results of pool for a single visit*

Description

[Experimental]

Usage

```
h_tidy_pool(x)
```

Arguments

x   (pool) is a list of pooled object from rbmi analysis results. This list includes
           analysis results, confidence level, hypothesis testing type.

Value

Data frame with results of pool for a single visit.

Examples

```
data("rbmi_test_data")
pool_obj <- rbmi_test_data

h_tidy_pool(pool_obj$pars[1:3])
```

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rbmi_test_data	*Example dataset for* tern.rbmi *package. This is an pool object from the rbmi analysis, see* browseVignettes(package = "tern.rbmi")

Description

[Experimental]

Usage

```
rbmi_test_data
```

Format

An object of class pool of length 5.

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summarize_rbmi   *Analyze function for tabulating LS means estimates from tidied* rbmi
                        pool *results.*

Description

[Experimental]

Usage

```
summarize_rbmi(
  lyt,
  ...,
  table_names = "rbmi_summary",
  .stats = NULL,
  .formats = NULL,
  .indent_mods = NULL,
  .labels = NULL
)
```

Arguments

lyt	(layout) input layout where analyses will be added to.
...	additional argument.
table_names	(character) this can be customized in case that the same vars are analyzed multiple times, to avoid warnings from rtables.
.stats	(character) statistics to select for the table.
.formats	(named character or list) formats for the statistics.
.indent_mods	(named integer) indent modifiers for the labels.
.labels	(named character) labels for the statistics (without indent).

Value

rtables layout for tabulating LS means estimates from tidied rbmi pool results.

Examples

```
library(rtables)
library(dplyr)
library(broom)

data("rbmi_test_data")
pool_obj <- rbmi_test_data

df <- tidy(pool_obj)

basic_table() %>%
  split_cols_by("group", ref_group = levels(df$group)[1]) %>%
  split_rows_by("visit", split_label = "Visit", label_pos = "topleft") %>%
  summarize_rbmi() %>%
  build_table(df)
```

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s_rbmi_lsmeans | *Statistics function which is extracting estimates from a tidied LS means data frame.*

Description

[Experimental]

Usage

```
s_rbmi_lsmeans(df, .in_ref_col, show_relative = c("reduction", "increase"))
```

Arguments

df	input dataframe
.in_ref_col	boolean variable, if reference column is specified
show_relative	"reduction" if (control - treatment, default) or "increase" (treatment - control) of relative change from baseline?

Value

A list of statistics extracted from a tidied LS means data frame.

Examples

```
library(rtables)
library(dplyr)
library(broom)

data("rbmi_test_data")
pool_obj <- rbmi_test_data
```

```
df <- tidy(pool_obj)

s_rbmi_lsmeans(df[1, ], .in_ref_col = TRUE)

s_rbmi_lsmeans(df[2, ], .in_ref_col = FALSE)
```

| tidy.pool | *Helper method (for* broom::tidy() *) to prepare a data frame from an* pool rbmi *object containing the LS means and contrasts and multiple visits* |

Description

[Experimental]

Usage

```
## S3 method for class 'pool'
tidy(x, ...)
```

Arguments

x   (pool) is a list of pooled object from rbmi analysis results. This list includes
            analysis results, confidence level, hypothesis testing type.

...   Additional arguments. Not used. Needed to match generic signature only.

Value

A dataframe

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