

Package ‘simhelpers’

May 4, 2022

Type Package

Title Helper Functions for Simulation Studies

Version 0.1.2

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Description

Calculates performance criteria measures and associated Monte Carlo standard errors for simulation results. Includes functions to help run simulation studies. Our derivation and explanation of formulas and our general simulation workflow is closely aligned with the approach described by Morris, White, and Crowther (2019) <DOI:10.1002/sim.8086>.

URL <https://meghapsimatrix.github.io/simhelpers/index.html>

BugReports <https://github.com/meghapsimatrix/simhelpers/issues>

Depends R (>= 2.10)

License GPL-3

Encoding UTF-8

LazyData true

RoxygenNote 7.1.2

SystemRequirements RStudio

Imports rlang, stats, dplyr, furrr, magrittr, tibble, rstudioapi,
tidyr, Rdpack

Suggests plyr, purrr, future, knitr, rmarkdown, pkgdown, covr,
testthat, kableExtra, ggplot2, broom

RdMacros Rdpack

VignetteBuilder knitr

NeedsCompilation no

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Repository CRAN

Date/Publication 2022-05-03 22:40:02 UTC

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alpha_res	<i>Cronbach's alpha simulation results</i>
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Description

A dataset containing simulation results from estimating Cronbach's alpha and its variance.

Usage

```
alpha_res
```

Format

A tibble with 1,000 rows and 3 variables:

A estimate of alpha.

Var_A estimate of the variance of alpha.

true_param true alpha used to generate the data.

calc_absolute	<i>Calculate absolute performance criteria and MCSE</i>
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Description

Calculates absolute bias, variance, mean squared error (mse) and root mean squared error (rmse). The function also calculates the associated Monte Carlo standard errors.

Usage

```
calc_absolute(  
  res_dat,  
  estimates,  
  true_param,  
  perfm_criteria = c("bias", "variance", "mse", "rmse")  
)
```

Arguments

`res_dat` data frame or tibble containing the simulation results.
`estimates` name of the column containing the estimates.
`true_param` name of the column containing the true parameters.
`perfm_criteria` character or character vector indicating the performance criteria to be calculated.

Value

A tibble containing the number of simulation iterations, performance criteria estimate(s) and the associated MCSE.

Examples

```
calc_absolute(res_dat = t_res, estimates = est, true_param = true_param)
```

calc_coverage	<i>Calculate confidence interval coverage, width and MCSE</i>
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Description

Calculates confidence interval coverage and width. The function also calculates the associated Monte Carlo standard errors. The confidence interval percentage is based on how you calculated the lower and upper bounds.

Usage

```
calc_coverage(  
  res_dat,  
  lower_bound,  
  upper_bound,  
  true_param,  
  perfm_criteria = c("coverage", "width")  
)
```

Arguments

res_dat	data frame or tibble containing the simulation results.
lower_bound	name of the column containing the lower bound estimates of the confidence intervals.
upper_bound	name of the column containing the upper bound estimates of the confidence intervals.
true_param	name of the column containing the true parameters.
perfm_criteria	character or character vector indicating the performance criteria to be calculated.

Value

A tibble containing the number of simulation iterations, performance criteria estimate(s) and the associated MCSE.

Examples

```
calc_coverage(res_dat = t_res, lower_bound = lower_bound,
              upper_bound = upper_bound, true_param = true_param)
```

calc_rejection	<i>Calculate rejection rate and MCSE</i>
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Description

Calculates rejection rate. The function also calculates the associated Monte Carlo standard error.

Usage

```
calc_rejection(res_dat, p_values, alpha = 0.05)
```

Arguments

res_dat	data frame or tibble containing the simulation results.
p_values	name of the column containing the p-values.
alpha	number indicating the nominal alpha level. Default value is set to the conventional .05.

Value

A tibble containing the number of simulation iterations, performance criteria estimate and the associated MCSE.

Examples

```
calc_rejection(res_dat = t_res, p_values = p_val)
```

calc_relative	<i>Calculate relative performance criteria and MCSE</i>
---------------	---

Description

Calculates relative bias, mean squared error (relative mse), and root mean squared error (relative rmse). The function also calculates the associated Monte Carlo standard errors.

Usage

```
calc_relative(  
  res_dat,  
  estimates,  
  true_param,  
  perfm_criteria = c("relative bias", "relative mse", "relative rmse")  
)
```

Arguments

res_dat	data frame or tibble containing the simulation results.
estimates	name of the column containing the estimates.
true_param	name of the column containing the true parameters.
perfm_criteria	character or character vector indicating the performance criteria to be calculated.

Value

A tibble containing the number of simulation iterations, performance criteria estimate(s) and the associated MCSE.

Examples

```
calc_relative(res_dat = t_res, estimates = est, true_param = true_param)
```

calc_relative_var	<i>Calculate jack-knife Monte Carlo SE for variance estimators</i>
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Description

Calculates relative bias, mean squared error (relative mse), and root mean squared error (relative rmse) of variance estimators. The function also calculates the associated jack-knife Monte Carlo standard errors.

Usage

```
calc_relative_var(  
  res_dat,  
  estimates,  
  var_estimates,  
  perfm_criteria = c("relative bias", "relative mse", "relative rmse")  
)
```

Arguments

res_dat	data frame or tibble containing the simulation results.
estimates	name of the column containing the estimates.
var_estimates	name of the column containing the variance estimates.
perfm_criteria	character or character vector indicating the performance criteria to be calculated.

Value

A tibble containing the number of simulation iterations, performance criteria estimate(s) and the associated MCSE.

Examples

```
calc_relative_var(res_dat = alpha_res, estimates = A, var_estimates = Var_A)
```

create_skeleton	<i>Open a simulation skeleton</i>
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Description

Creates and opens a .R file containing a skeleton for writing a Monte Carlo simulation study.

Usage

```
create_skeleton()
```

Examples

```
## Not run:
create_skeleton()

## End(Not run)
```

evaluate_by_row	<i>Evaluate a simulation function on each row of a data frame or tibble</i>
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Description

Evaluates a simulation function on each row of a data frame or tibble containing parameter values. Returns a single tibble with parameters and simulation results. The function uses `furrr::future_pmap`, which allows for easy parallelization.

Usage

```
evaluate_by_row(
  params,
  sim_function,
  ...,
  .progress = FALSE,
  .options = furrr::furrr_options(),
  system_time = TRUE
)
```

Arguments

<code>params</code>	data frame or tibble containing simulation parameter values. Each row should represent a separate set of parameter values.
<code>sim_function</code>	function to be evaluated, with argument names matching the variable names in <code>params</code> . The function must return a <code>data.frame</code> , <code>tibble</code> , or <code>vector</code> .
<code>...</code>	additional arguments passed to <code>sim_function</code> .
<code>.progress</code>	A single logical. Should a progress bar be displayed? Only works with multisession, multicore, and multiprocess futures. Note that if a multicore/multisession future falls back to sequential, then a progress bar will not be displayed. Warning: The <code>.progress</code> argument will be deprecated and removed in a future version of <code>furrr</code> in favor of using the more robust <code>progressr</code> package.
<code>.options</code>	The future specific options to use with the workers. This must be the result from a call to <code>furrr_options()</code> .
<code>system_time</code>	logical indicating whether to print computation time. TRUE by default.

Value

A tibble containing parameter values and simulation results.

Examples

```
df <- data.frame(  
  n = 3:5,  
  lambda = seq(8, 16, 4)  
)  
  
evaluate_by_row(df, rpois)
```

Tipton_Pusto

Results for Figure 2 of Tipton & Pustejovsky (2015)

Description

A dataset containing simulation results comparing small sample correction methods for cluster robust variance estimation in meta-analysis.

Usage

Tipton_Pusto

Format

A tibble with 15,300 rows and 8 variables:

num_studies the number of studies included in the meta-analysis.

r correlation between outcomes.

Isq measure of heterogeneity of true effects.

contrast type of contrast that was tested.

test small sample method used.

q the number of parameters in the hypothesis test.

rej_rate the Type 1 error rate.

mcse the Monte Carlo standard error for the estimate of the Type 1 error rate.

Source

Tipton E, Pustejovsky JE (2015). "Small-Sample Adjustments for Tests of Moderators and Model Fit Using Robust Variance Estimation in Meta-Regression." *Journal of Educational and Behavioral Statistics*, **40**(6), 604–634. ISSN 1076-9986, 1935-1054, doi: [10.3102/1076998615606099](https://journals.sagepub.com/doi/10.3102/1076998615606099), <https://journals.sagepub.com/doi/10.3102/1076998615606099>.

t_res	<i>t-test simulation results</i>
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Description

A dataset containing simulation results from a study that just runs a t-test.

Usage

t_res

Format

A tibble with 1,000 rows and 5 variables:

est estimate of the mean difference.

p_val p-value from the t-test.

lower_bound lower bound of the confidence interval.

upper_bound upper bound of the confidence interval.

true_param true mean difference used to generate the data.

welch_res	<i>Welch t-test simulation results</i>
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Description

A dataset containing simulation results from a study comparing Welch t-test to the conventional t-test.

Usage

welch_res

Format

A tibble with 16,000 rows and 11 variables:

n1 sample size for Group 1.

n2 sample size for Group 2.

mean_diff true difference in means of two groups used to generate the data.

iterations number of iterations.

seed seed used to generate data.

method indicates whether Welch or conventional t-test was used.

est estimate of the mean difference.

var variance of the estimate.

p_val p-value from the t-test.

lower_bound lower bound of the confidence interval.

upper_bound upper bound of the confidence interval.

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