UNIVERSITY OF LEICESTER

Analysing Results From Monte Carlo Simulation Studies

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useR! 2019

Monte Carlo simulation studies

Monte Carlo simulation studies are computer experiments that involve generating data by pseudo-random sampling.

The Process

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3. Compare the analysis results with the truth.

Monte Carlo simulation studies are useful...

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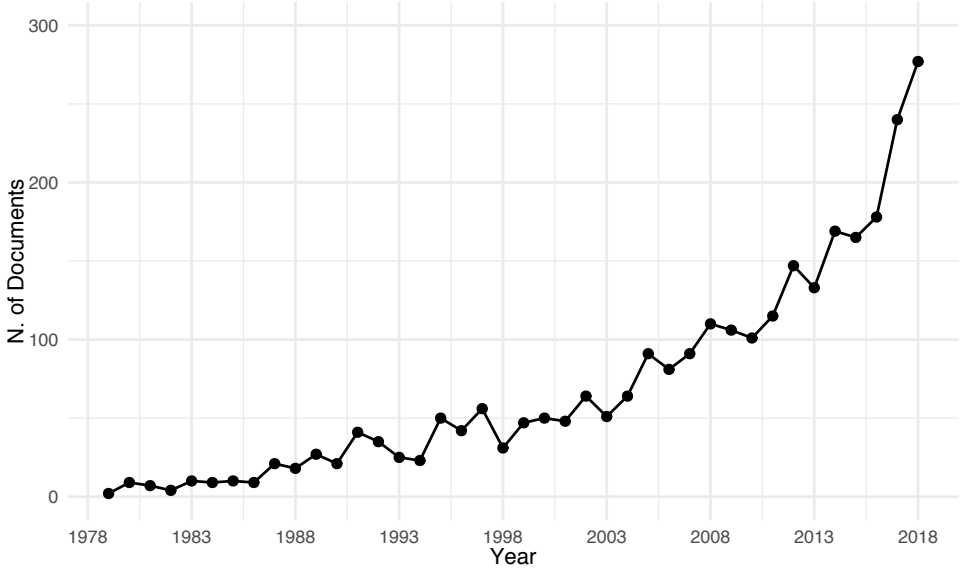
They can also help to answer questions such as:

- Is an estimator biased in a finite sample?
- Do confidence intervals for a given parameter achieve the desired nominal level of coverage?
- How does a newly developed method compare to an established one?
- What is the power to detect a desired effect size under complex experimental settings and analysis methods?
- You name it!

...and increasingly popular!



Scopus search key: `TITLE-ABS-KEY ("Monte Carlo simulation study")`

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2. Reproducibility of results;
3. Dissemination of results.

Disclaimer!

I will not cover how to plan, design, and run a Monte Carlo simulation study.

An full example on modelling survival data is included with `rsimsum`:

```
vignette("relhaz", package = "rsimsum")
```

Also, check out the tutorial paper by Morris *et al*. (2019).

Why yet *another R package?*

Why yet *another R package?*

- There is a similar package in Stata, but nothing comparable in R (that I know);
- Several performance measures are supported - no need to do tedious (and error-prone) calculations by hand;
- Monte Carlo standard errors are computed and displayed by default;
- Several quick plotting options for fast iteration/exploration.

rsimsum's main function: simsum

```
args(rsimsum::simsum)
```

```
## function (data, estvarname, true, se, methodvar = NULL, ref = NULL,
##     by = NULL, ci.limits = NULL, dropbig = FALSE, x = FALSE,
##     control = list())
## NULL
```

Documentation: https://ellessenne.github.io/rsimsum/

Example: t-test (1)

Aim: investigate the performance of the two-sample t-test when

1. Data is skewed;
2. Variances are unequal in the two groups.

Example: t-test (2)

Data-generating mechanisms (DGMs):

1. Simulating 60 observations, with 2:1 groups ratio;
2. Fully fractional design, varying equal/unequal variance, and skewed/non-skewed data.

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Replications: 2,000 per DGM.

Case study: data

```
dplyr::glimpse(data)
```

```
## Observations: 16,000
## Variables: 8
## $ diff <dbl> -2.802464229, -0.079683569, -2.802464229, -0.079683569, -1.7...
## $ se <dbl> 1.279933, 2.218313, 1.197805, 2.947133, 1.363249, 1.723423, ...
## $ df <dbl> 58.00000, 58.00000, 46.14587, 22.18708, 58.00000, 58.00000, ...
## $ i <int> 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 2, 3, 3, 3, 3, ...
## $ dgm <fct> "Equal, Non-skewed", "Unequal, Non-Skewed", "Equal, Non-skew...
## $ method <fct> t-test (P), t-test (P), t-test (NP), t-test (NP), t-test
(P)...
## $ dist <fct> N, N, N, N, Gamma, Gamma, Gamma, Gamma, N, N, N, N, Gamma, G...
## $ var <fct> Equal, Unequal, Equal, Unequal, Equal, Unequal, Equal, Unequ...
```

Summarising a simulation study (1)

```
s <- rsimsum::simsum(
  data = data, estvarname = "diff", se = "se", true = -1, methodvar = "method",
  by = "dgm", ref = "t-test (NP)", x = TRUE
)
s
```

```
## Summary of a simulation study with a single estimand.
##
## Method variable: method
##  Unique methods: t-test (NP), t-test (P)
##  Reference method: t-test (NP)
##
## By factors: dgm
##
## Monte Carlo standard errors were computed.
```

Summarising a simulation study (2)

```
summary(s, stats = "bias")
```

```
## Values are:
##  Point Estimate (Monte Carlo Standard Error)
##
## Bias in point estimate:
##                   dgm      t-test (NP)       t-test (P)
##    Equal, Non-skewed  0.0054 (0.0302)  0.0054 (0.0302)
##  Unequal, Non-Skewed -0.0461 (0.0547) -0.0461 (0.0547)
##        Equal, Skewed -0.0610 (0.0300) -0.0610 (0.0300)
##      Unequal, Skewed -0.0258 (0.0533) -0.0258 (0.0533)
```

Summarising a simulation study (3)

```
summary(s, stats = "cover")

## Values are:
##  Point Estimate (Monte Carlo Standard Error)
##
## Coverage of nominal 95% confidence interval:
##                   dgm     t-test (NP)      t-test (P)
##    Equal, Non-skewed 0.9590 (0.0044) 0.9600 (0.0044)
##  Unequal, Non-Skewed 0.9350 (0.0055) 0.8725 (0.0075)
##        Equal, Skewed 0.9495 (0.0049) 0.9525 (0.0048)
##      Unequal, Skewed 0.9290 (0.0057) 0.8770 (0.0073)
```

Performance measures implemented in `rsimsum`

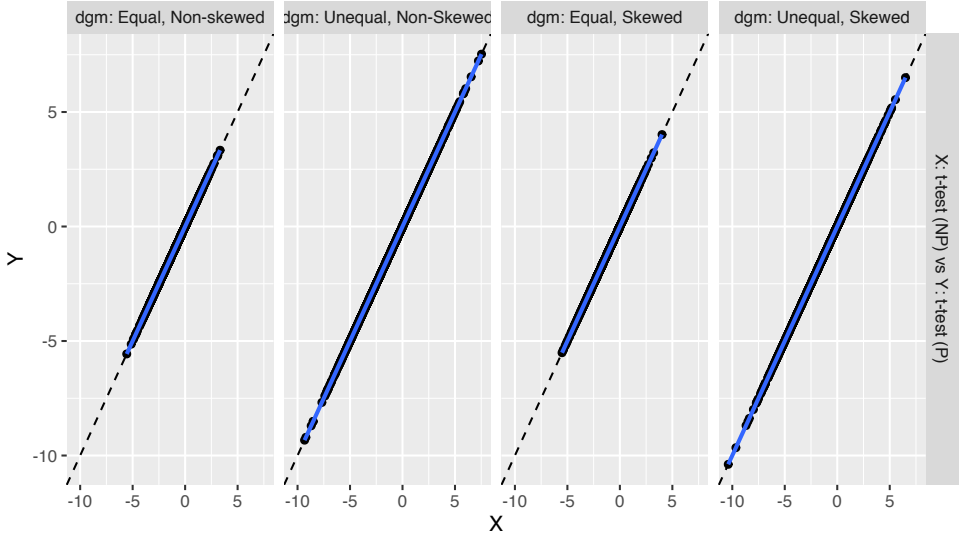
The following performance measures are implemented in `rsimsum`:

- Bias;
- Empirical SE, relative % increase in precision, model-based SE, and relative % error in model-based SE;
- Mean squared error (MSE);
- Coverage probability and bias-corrected coverage probability;
- Power of type I error.

Each performance measure is described in more detail elsewhere (Morris *et al.*, 2019).

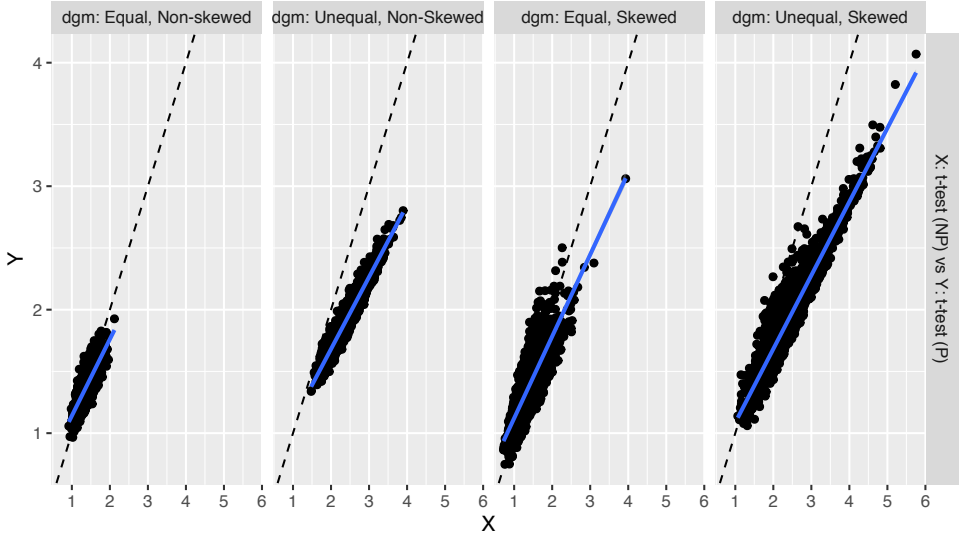
Plotting point estimates

```
autoplot(object = s, type = "est")
```



Plotting standard errors

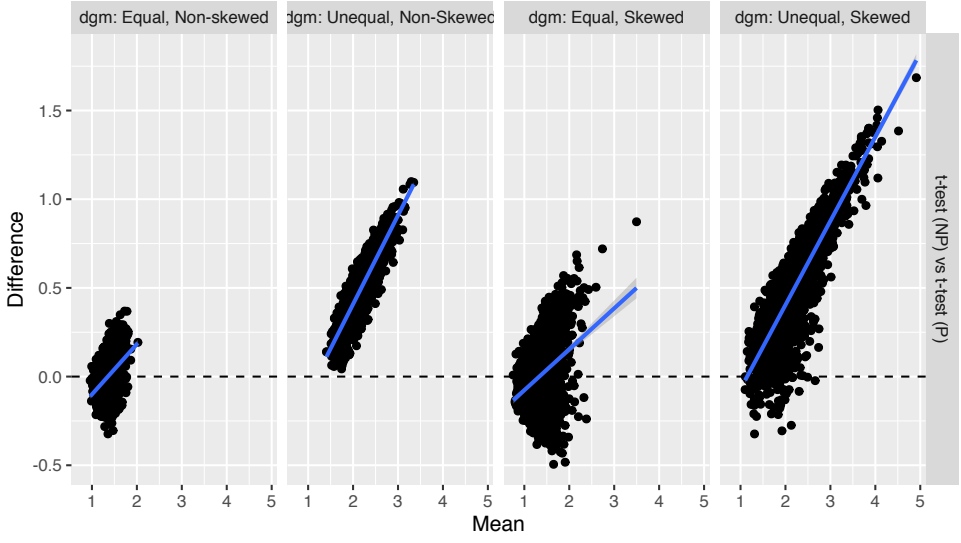
```
autoplot(object = s, type = "se")
```



Comparison of variable 'se'

Plotting standard errors

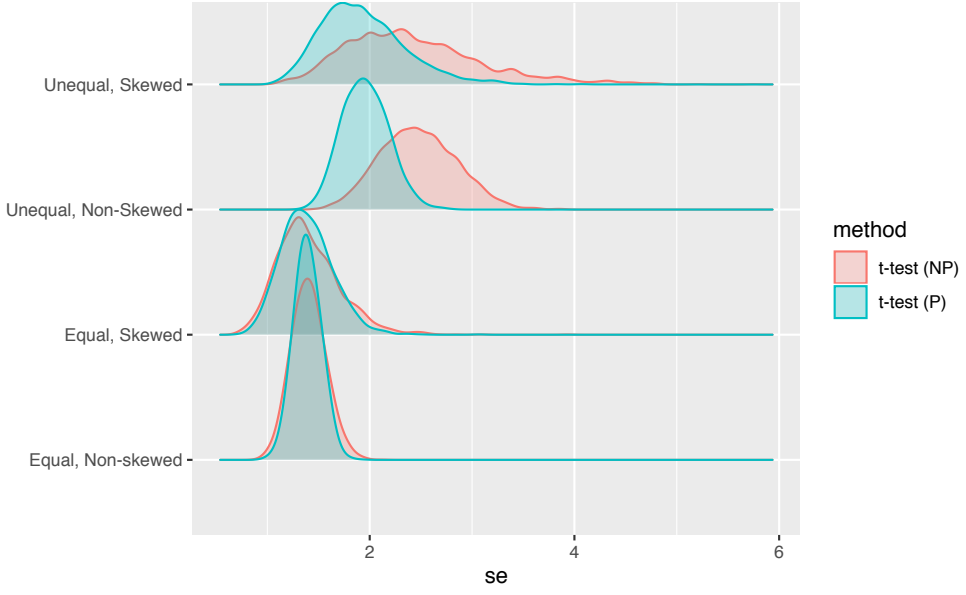
```
autoplot(object = s, type = "se_ba")
```



Comparison of variable 'se'; Bland-Altman type plot

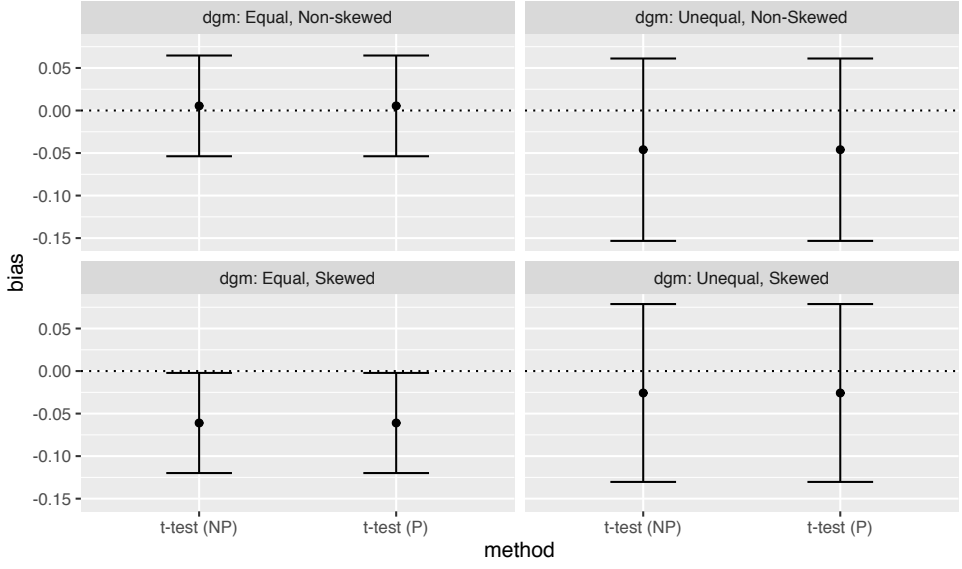
Plotting standard errors

```
autoplot(object = s, type = "se_ridge")
```



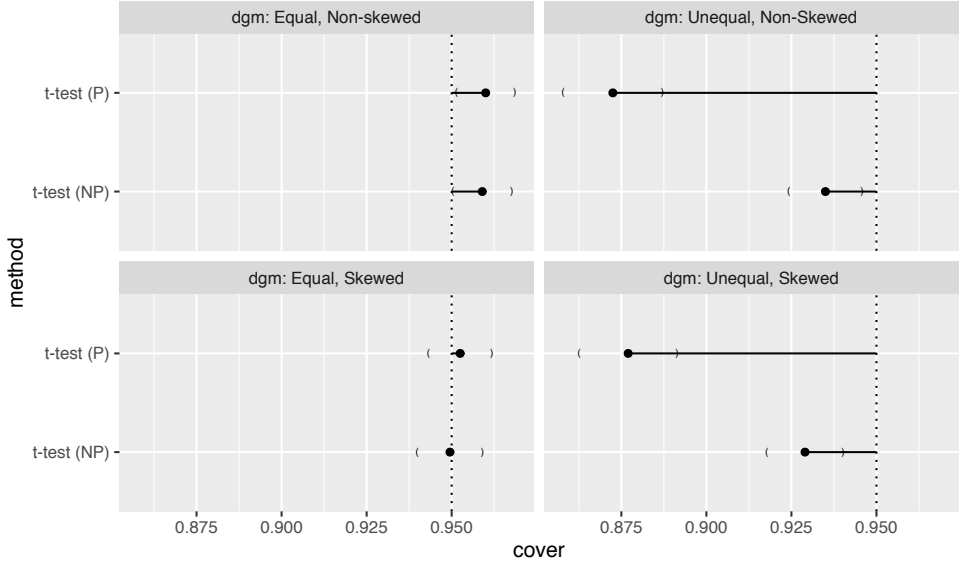
Plotting results

```
autoplot(object = summary(s), type = "forest", stats = "bias")
```



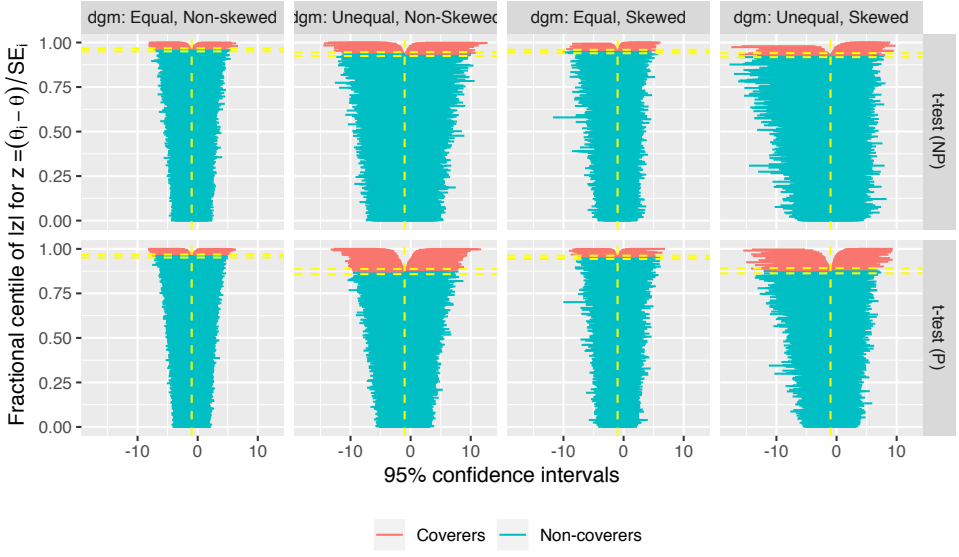
Plotting results

```
autoplot(object = summary(s), type = "lolly", stats = "cover")
```



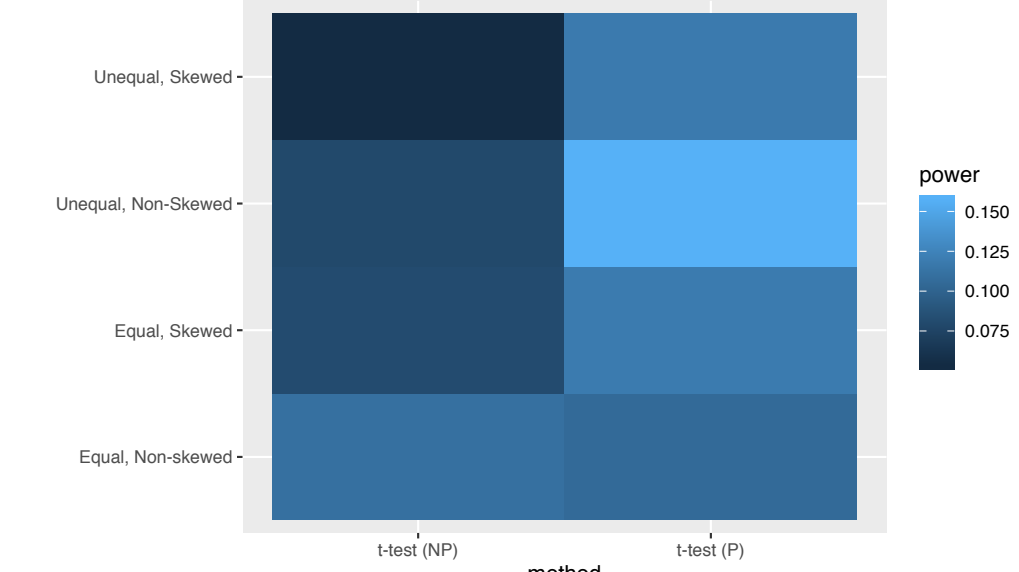
Plotting results

```
autoplot(object = s, type = "zip")
```



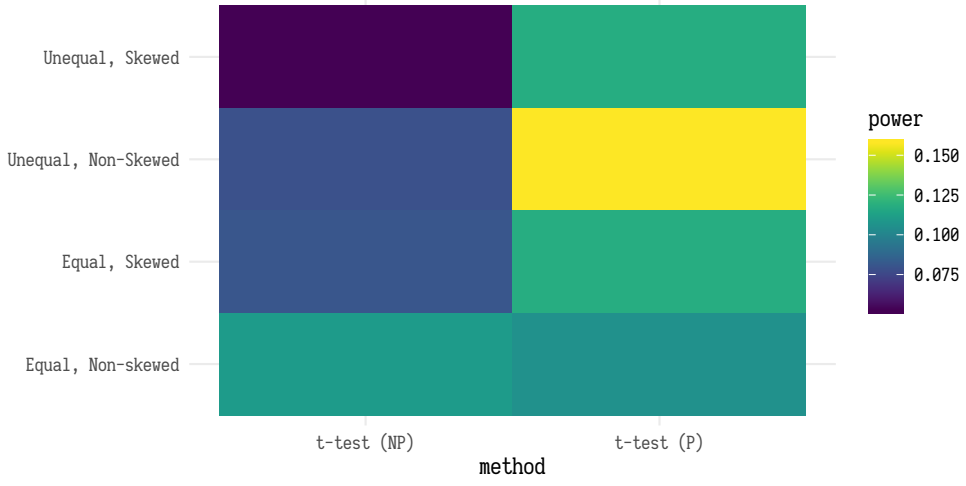
Plotting results

```
autoplot(object = s, type = "heat", stats = "power")
```



Plotting results

```
autoplot(object = s, type = "heat", stats = "power") +
  viridis::scale_fill_viridis() +
  ggplot2::theme_minimal(base_family = "Iosevka Slab")
```



Multiple estimands

```
args(rsimsum::multisimsum)

## function (data, par, estvarname, true, se, methodvar = NULL,
##     ref = NULL, by = NULL, ci.limits = NULL, dropbig = FALSE,
##     x = FALSE, control = list())
## NULL
```

INteractive **T**ool for **E**xploring
REsults from **S**imulation s**T**udies

Why a Shiny app?

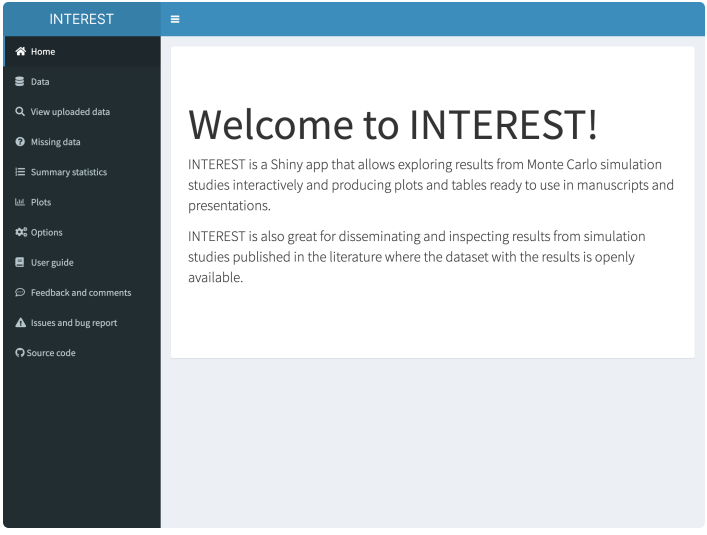
Why a Shiny app?

- Dissemination of results and open science;
- Fast(er) iteration and exploration of results;
- Supporting devices where R does not run natively (smartphones, Chromebooks, ...).

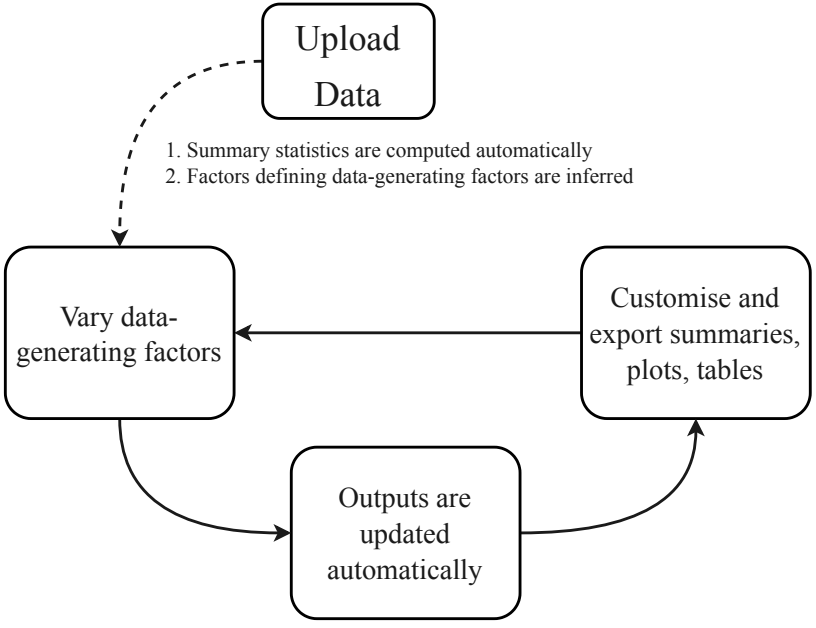
Dissemination is key

1. Can drive practitioners and applied statisticians to methods that have been shown to perform well in their practical settings;
2. Can guide researchers to develop new methods in promising directions;
3. Can provide insights into less established methods.

INTEREST: Homepage



INTEREST: Workflow



http://interest.shinyapps.io/interest/

How to get rsimsum and INTEREST?

rsimsum can be installed directly from CRAN:

```
install.packages("rsimsum")
# Development version on GitHub:
# require("remotes")
# remotes::install_github(repo = "ellessenne/rsimsum")
```

INTEREST is on GitHub:

```
# require("remotes")
remotes::install_github(repo = "ellessenne/interest")
```

What is coming next?

`rsimsum`:

- Nested loop plot for simulation studies with several DGMs (Rücker and Schwarzer, 2014);
- Methods to easily reproduce plots generated by `autoplot`;
- Methods to directly export (pretty) LaTeX tables;
- ...

INTEREST:

- Additional exporting tools for INTEREST;
- Support for simulation studies with multiple estimands in INTEREST;
- ...

Thank you!

References:

- *Using simulation studies to evaluate statistical methods*. Morris TP, White IR, and Crowther MJ (2019). Statistics in Medicine 38(11):2074–2102, DOI: 10.1002/sim.8086
- *rsimsum: Summarise results from Monte Carlo simulation studies*. Gasparini A (2018). Journal of Open Source Software 3(26):739, DOI: 10.21105/joss.00739

- `rsimsum`'s website: `https://ellessenne.github.io/rsimsum/`

Slides available online: `https://tinyurl.com/useR-2019`