

# Analysing Results From Monte Carlo Simulation Studies

A Gasparini (@ellesenne) · TP Morris · IR White · MJ Crowther

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

In a simulation study, we:

1. Generate data from a known distribution (so that we know the “truth”);
2. Analyse the data;
3. Compare the analysis results with the truth.

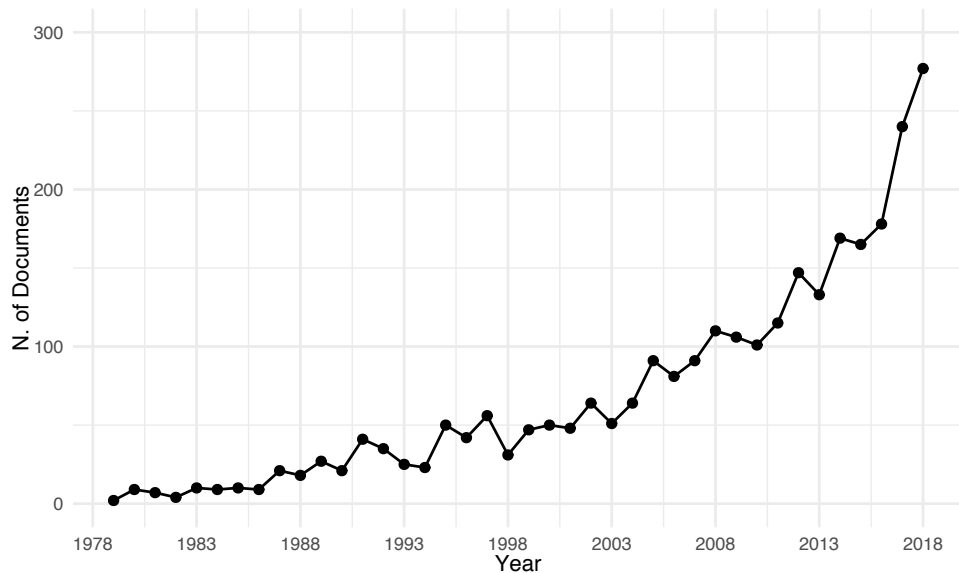
# Monte Carlo simulation studies are useful...

Monte Carlo simulation studies provide an invaluable tool for statistical and biostatistical research.

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.



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).

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

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

Data-generating mechanisms (DGMs):

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## Example: t-test (2)

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Replications: 2,000 per DGM.

## Case study: data

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

```
## Observations: 16,000
```

```
## Variables: 8
```

```
## $ diff <dbl> -3.56987311, -1.37912667, -3.56987311, -1.37912667, -0.45626...
```

```
## $ se <dbl> 1.440554, 1.973592, 1.405570, 2.434108, 1.474513, 2.185057, ...
```

```
## $ df <dbl> 48.00000, 48.00000, 35.55937, 21.15939, 48.00000, 48.00000, ...
```

```
## $ 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)",
```

```
...
```

```
## $ 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...
```

```
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/>

## 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	0.0048 (0.0335)	0.0048 (0.0335)	
## Unequal, Non-Skewed	0.0288 (0.0591)	0.0288 (0.0591)	
## Equal, Skewed	-0.0474 (0.0334)	-0.0474 (0.0334)	
## Unequal, Skewed	0.0219 (0.0583)	0.0219 (0.0583)	

## 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 0.9485 (0.0049) 0.9500 (0.0049)
```

```
## Unequal, Non-Skewed 0.9465 (0.0050) 0.8825 (0.0072)
```

```
##       Equal, Skewed 0.9535 (0.0047) 0.9570 (0.0045)
```

```
##       Unequal, Skewed 0.9200 (0.0061) 0.8775 (0.0073)
```

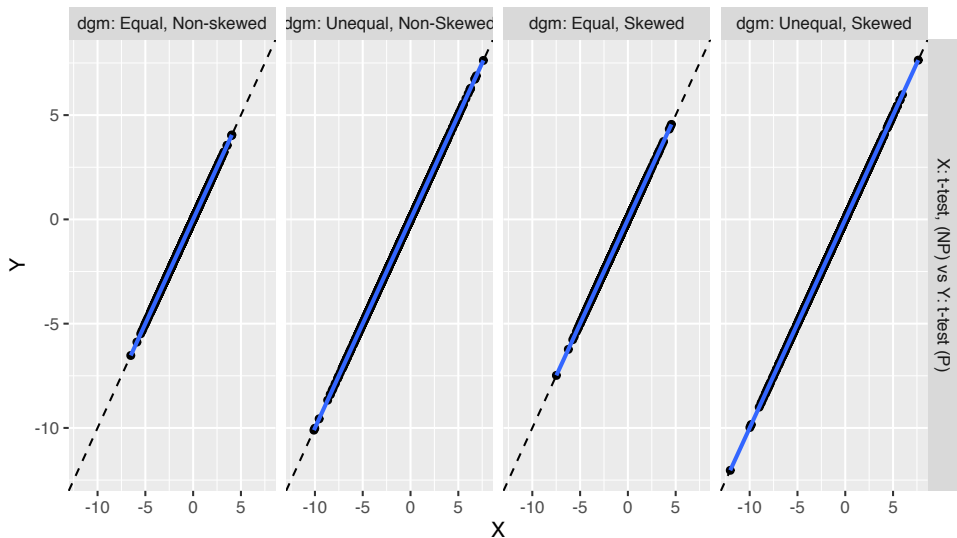
The following performance measures are implemented in `rsimsum`:

- 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

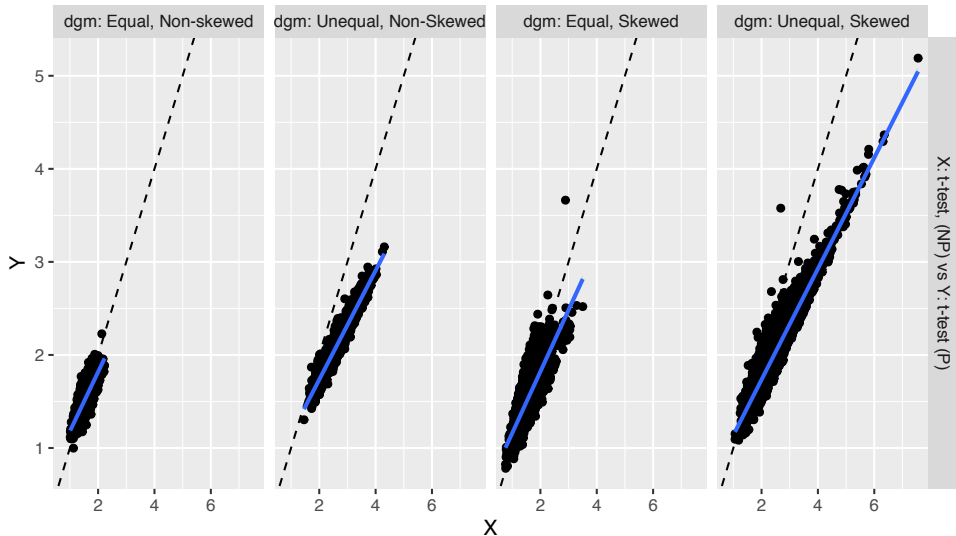
```
autoplot(object = s, type = "est")
```



Comparison of variable 'diff'

# Plotting standard errors

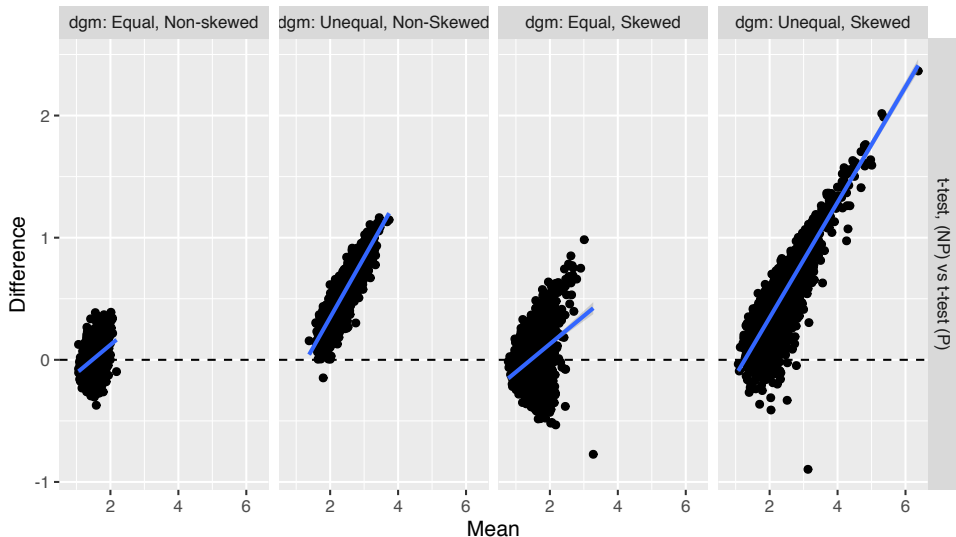
```
autoplot(object = s, type = "se")
```



Comparison of variable 'se'

# Plotting standard errors

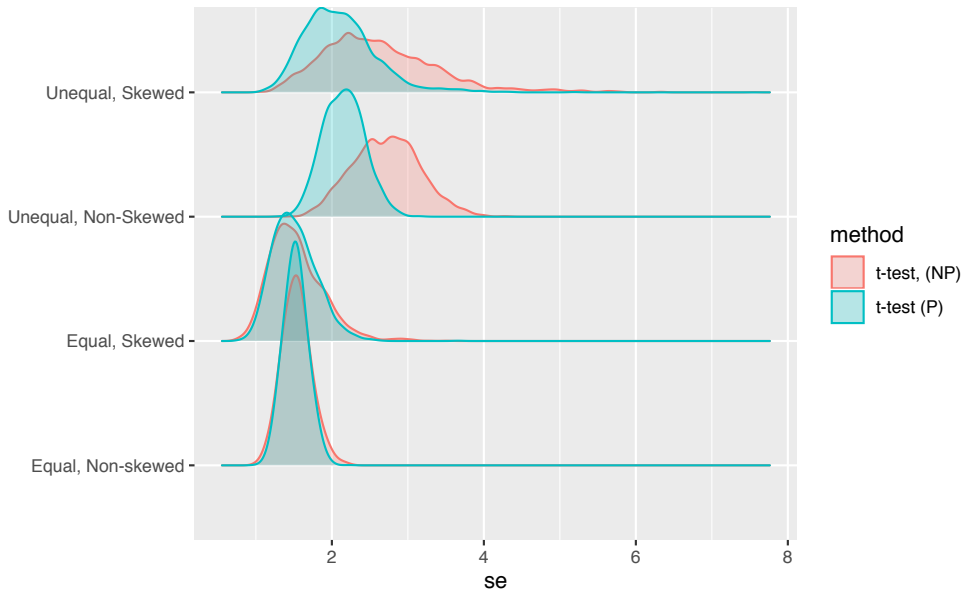
```
autoplot(object = s, type = "se_ba")
```



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

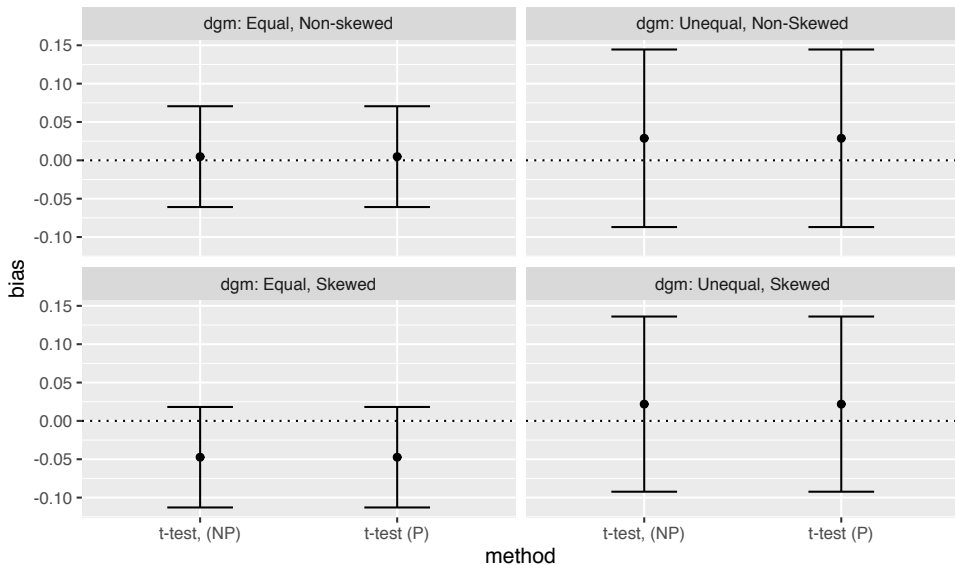
# Plotting standard errors

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



# Plotting results

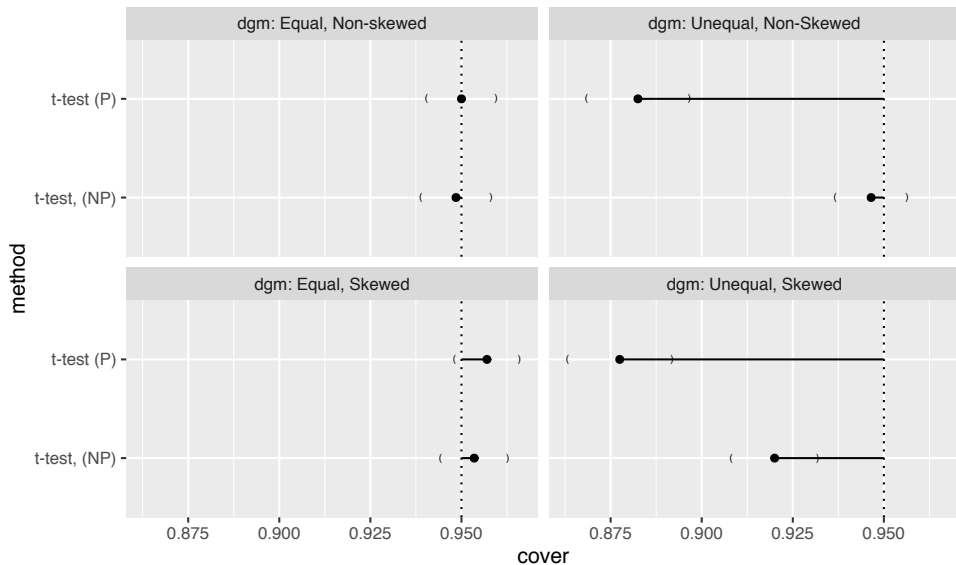
```
autoplot(object = summary(s), type = "forest", stats = "bias")
```





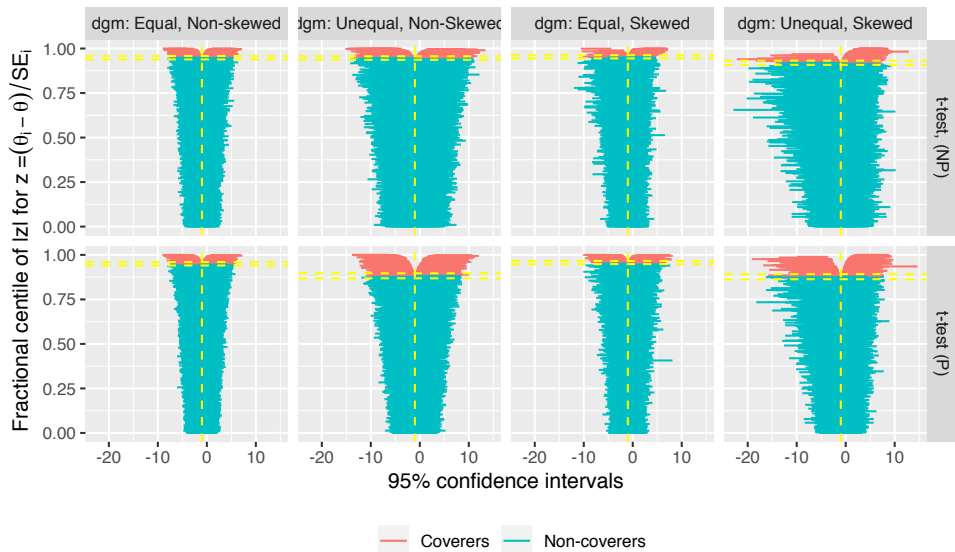
# Plotting results

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



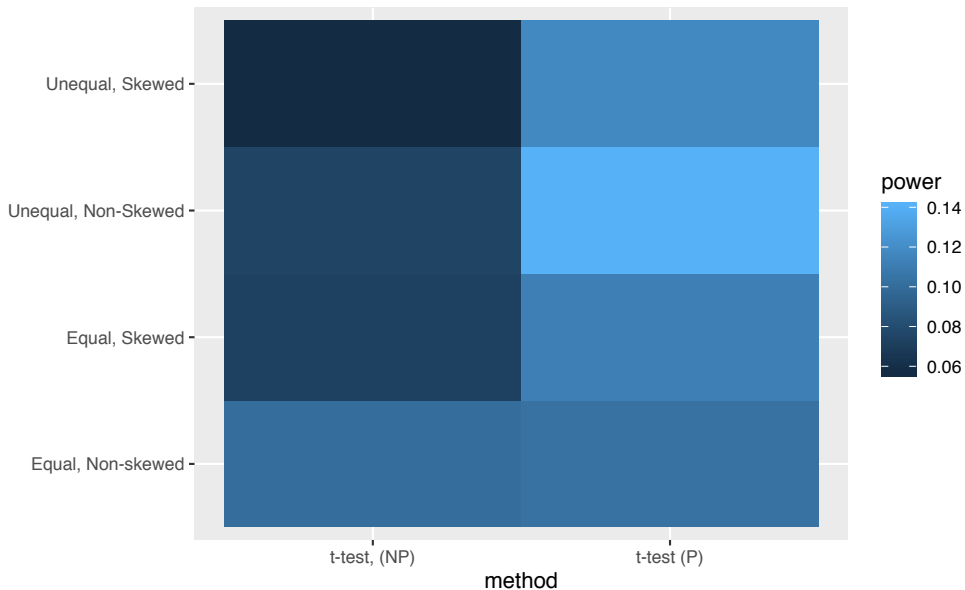
# Plotting results

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



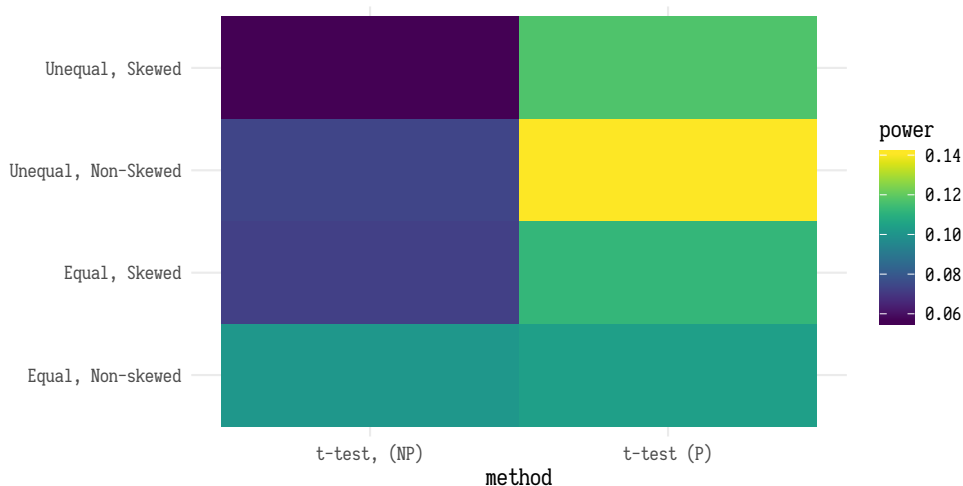
# Plotting results

```
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")
```



```
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
```

# **I**nteractive **T**ool for **E**xploring **R**esults from **S**imulation **sT**udies

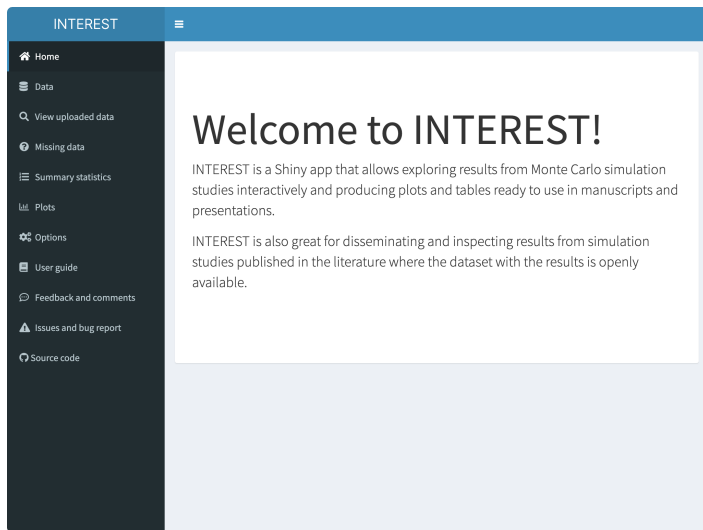
*Why a Shiny app?*

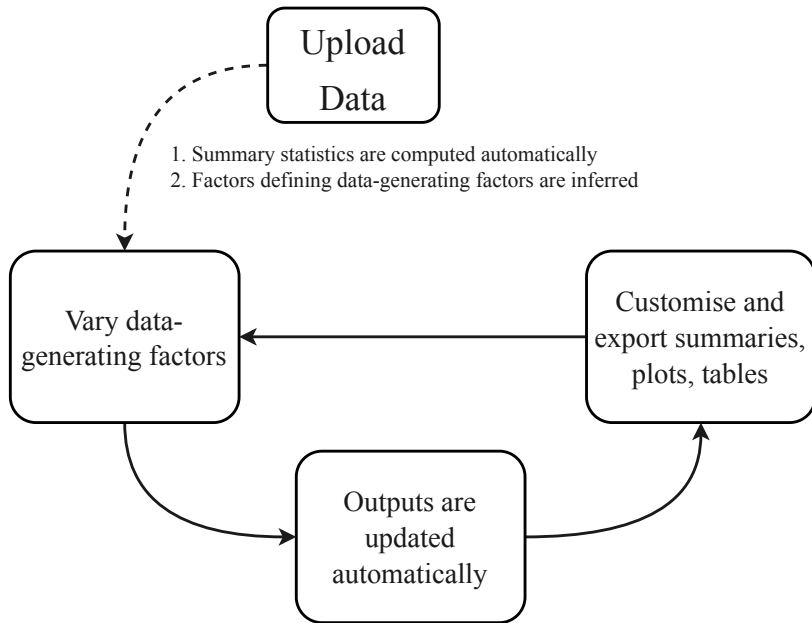
### *Why a Shiny app?*

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



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.





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

## How to get rsumsum and INTEREST?

rsumsum can be installed directly from CRAN:

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

INTEREST is on GitHub:

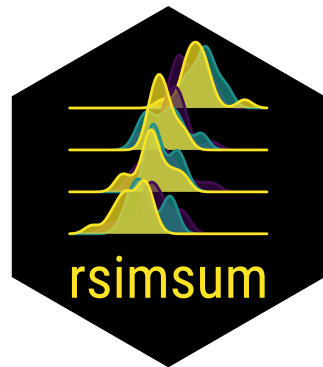
```
# require('remotes')  
remotes::install_github(repo = 'ellessenne/interest')
```

# What is coming next?

- 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)  $\text{\LaTeX}$  tables;
- Additional exporting tools for INTEREST;
- Support for simulation studies with multiple estimands in INTEREST;
- ...

## 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 from [add GitHub \(short\) link here!](#)