




simChef: High-quality data science simulations in R

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Summary

simChef is an R package that empowers data science practitioners to rapidly plan, carry out, and summarize statistical simulation studies in a flexible, efficient, and low-code manner. Drawing substantially from the Predictability, Computability, and Stability (PCS) framework (Yu & Kumbier, 2020), simChef emphasizes the scientific best practices encompassed by PCS by removing many of the administrative burdens of simulation design through: (1) an intuitive [tidy grammar](#) of data science simulations; (2) powerful abstractions for distributed simulation processing backed by future (Bengtsson, 2021); and (3) automated generation of interactive [R Markdown](#) simulation documentation, situating results next to the workflows needed to reproduce them. Taken together, simChef's capabilities overcome many of the design, computational, and reproducibility hurdles inherent in nearly every data science simulation study.

Statement of need

Data science simulation studies occupy an important role in scientific research as a means to gain insight into new and existing statistical methods. Simulations serve as statistical sandboxes that open a path toward otherwise inaccessible discoveries. For example, they can be used to establish comprehensive benchmarks of existing procedures for a common task; to demonstrate the strengths and weaknesses of novel methodology applied to synthetic and real-world data; or to probe the validity of a theoretical analysis.

Creating high-quality simulation studies typically involves a number of repetitive and error-prone coding tasks: implementing data-generating processes (DGPs) and statistical methods; sampling from these DGPs; parallelizing computation of simulation replicates; summarizing metrics; visualizing, documenting, presenting, and saving results; and so on. While this

administrative overhead is necessary, it is not sufficient for scientific understanding. Data scientists must navigate a number of important judgment calls such as the choice of DGPs, baseline statistical methods, associated parameters, and evaluation metrics for scientific relevancy.

While the scientific context may vary drastically from one study to the next, the simulation scaffolding remains largely similar. Yet simulation code repositories often lack reusability, both for novel settings and when new questions arise in the original context. *simChef* addresses the need for an intuitive, extensible, and reusable framework for data science simulations, allowing data science practitioners to focus their energies on scientific questions by reducing the burdens of parameterization, parallelization, and documentation.

Core abstractions of data science simulations

At its core, *simChef* breaks down a simulation experiment into four modular components (Figure 1), each implemented as an R6 class (Chang, 2022):

- DGP: the data-generating processes from which to *generate* data
- Method: the methods (or models) to *fit* in the experiment
- Evaluator: the evaluation metrics used to *evaluate* the methods' performance
- Visualizer: the visualization functions used to *visualize* outputs from the method fits or evaluation results (can be tables, plots, or even R Markdown snippets to display)

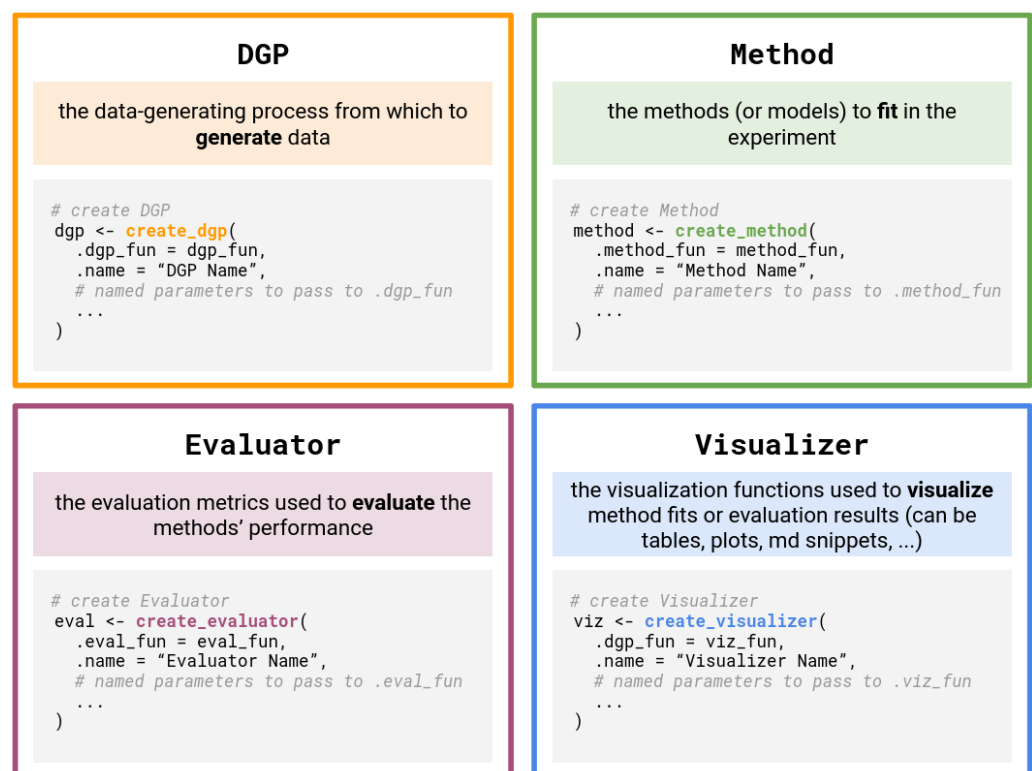


Figure 1: Overview of the four core components in a *simChef* Experiment. *simChef* provides four classes that implement distinct simulation objects in an intuitive and modular manner: DGP, Method, Evaluator, and Visualizer. Using these classes, users can easily build a *simChef* Experiment using reusable, customizable functions (i.e., `dgp_fun`, `method_fun`, `eval_fun`, and `viz_fun`). Optional named parameters can be set in these custom functions via the `...` arguments in the `create_*`() methods.

Using these classes, users can create or reuse custom functions (i.e., `dgp_fun`, `method_fun`,

eval_fun, and viz_fun in Figure 1) aligned with their scientific goals. The custom functions then can be parameterized and encapsulated in one of the corresponding classes via a create_* method, together with optional named parameters (see Figure 1).

A fifth R6 class, Experiment, unites the four components above and serves as a concrete implementation of the user's intent to answer a specific scientific question. Specifically, the Experiment stores references to the DGP(s), Method(s), Evaluator(s), and Visualizer(s) along with the DGP and Method parameters that should be varied and combined during the simulation run.

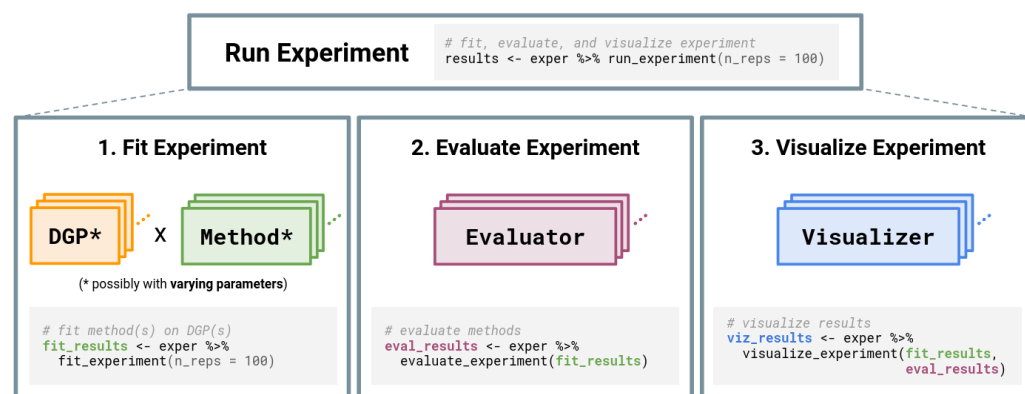


Figure 2: Overview of running a simChef Experiment. The Experiment class handles relationships among the four classes portrayed in Figure 1. Experiments may have multiple DGPs and Methods, which are combined across the Cartesian product of their varying parameters (represented by *). Once computed, each Evaluator and Visualizer takes in the fitted simulation replicates, while Visualizer additionally receives evaluation summaries.

A powerful grammar of data science simulations

Inspired by the tidyverse (Wickham et al., 2019), simChef develops an intuitive grammar for running simulation studies using the aforementioned R6 classes. We provide an illustrative example usage next.

```
library(simChef)

dgp1 <- create_dgp(dgp_fun1, "my_dgp1", sd = 0.5)
dgp2 <- create_dgp(dgp_fun2, "my_dgp2")
method <- create_method(method_fun, "my_method")
eval <- create_evaluator(eval_fun)
viz <- create_visualizer(viz_fun)

exper <- create_experiment(dgp_list = list(dgp1, dgp2)) %>%
  add_method(method) %>%
  add_vary_across(
    list(dgp1, dgp2),
    n = c(1e2, 1e3, 1e4)
  ) %>%
  add_vary_across(
    dgp2,
    sparse = c(FALSE, TRUE)
  ) %>%
  add_vary_across(
    method,
```

```

    scalar_valued_param = c(0.1, 1.0, 10.0),
    vector_valued_param = list(c(1, 2, 3), c(4, 5, 6)),
    list_valued_param = list(list(a1=1, a2=2, a3=3),
                              list(b1=3, b2=2, b3=1))
  ) %>%
add_evaluator(eval) %>%
add_viz(viz)

future::plan(multicore, workers = 64)

results <- exper %>%
  run_experiment(n_reps = 100, save = TRUE)

new_method <- create_method(new_method_fun, 'my_new_method')

exper <- exper %>%
  add_method(new_method)

results <- exper %>%
  run_experiment(n_reps = 100, use_cached = TRUE)

init_docs(exper)
render_docs(exper)

```

In the example usage, DGP(s), Method(s), Evaluator(s), and Visualizer(s) are first created via `create_*`(). These simulation objects can then be combined into an Experiment using either `create_experiment()` and/or `add_*`().

In an Experiment, DGP(s) and Method(s) can also be varied across one or multiple parameters via `add_vary_across()`. For instance, in the example Experiment, there are two DGP instances, both of which are varied across three values of `n` and one of which is additionally varied across two values of `sparse`. This effectively results in nine distinct configurations for data generation (i.e., 3 variations on `dgp1` + 3x2 variations on `dgp2`). For the single Method in the experiment, we use three values of `scalar_valued_param`, two of `vector_valued_param`, and another two of `list_valued_param`, giving 12 distinct configurations. Hence, there are a total of 9x12 = 108 DGP-method-parameter combinations in the Experiment.

Thus far, we have simply instantiated an Experiment object (akin to creating a recipe for an experiment). To compute and run the simulation experiment, we next call `run_experiment` with the desired number of replicates. As summarized in Figure 2, running the experiment will (1) *fit* each Method on each DGP (and for each of the varying parameter configurations), (2) *evaluate* the experiment according to the given Evaluator(s), and (3) *visualize* the experiment according to the given Visualizer(s). Furthermore, the number of replicates per combination of DGP, Method, and parameters specified via `add_vary_across` is determined by the `n_reps` argument to `run_experiment`. Because replication happens at the per-combination level, the effective total number of replicates in the Experiment depends on the number of DGPs, methods, and varied parameters. In the given example, there are 108 DGP-method-parameter combinations, each of which is replicated 100 times. To reduce the computational burden, the Experiment class flexibly handles the computation of simulation replicates in parallel using the `future` package (Bengtsson, 2021). Figure 3 provides a detailed schematic of the `run_experiment` workflow, along with the expected inputs to and outputs from user-defined functions.

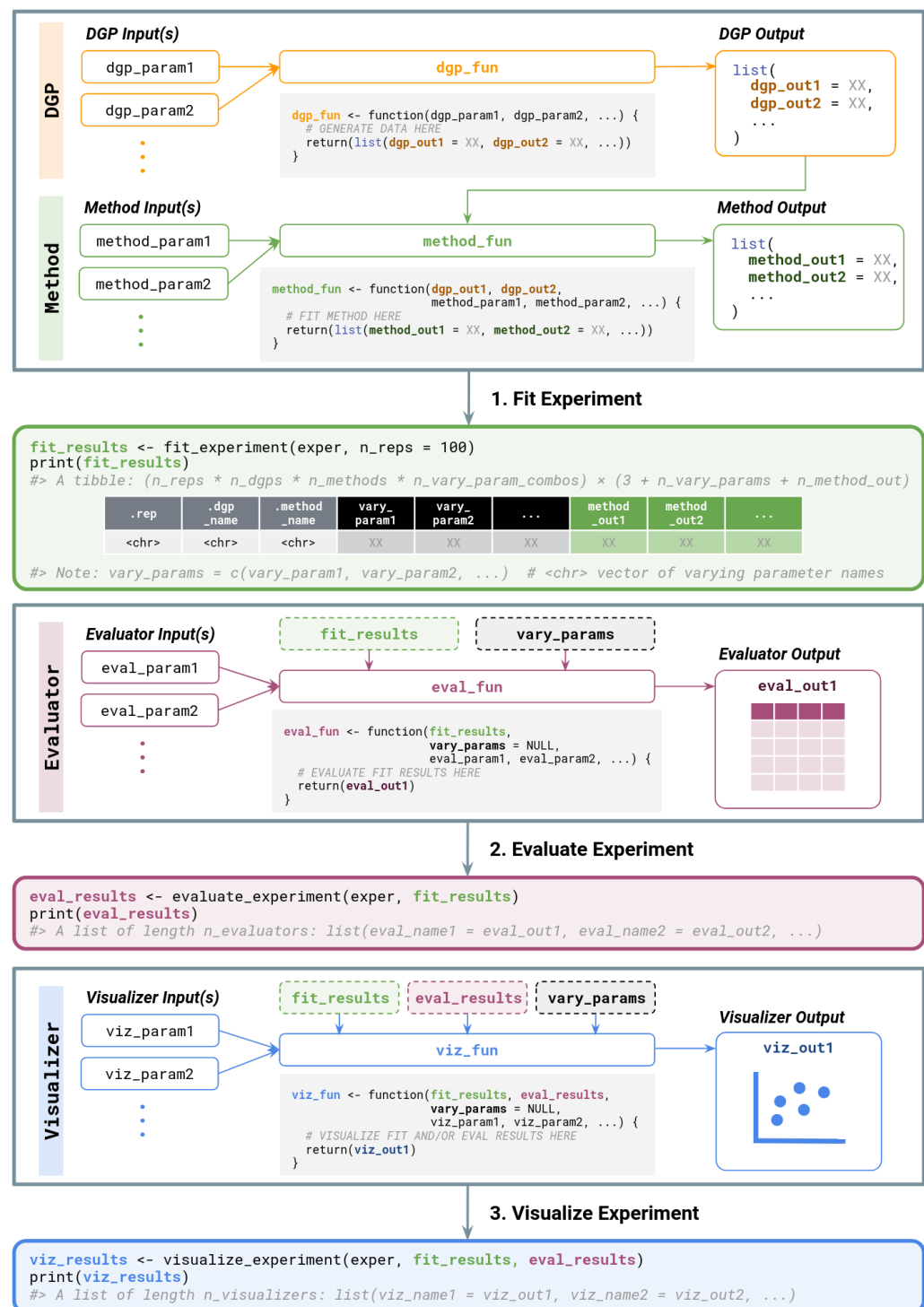


Figure 3: Detailed schematic of the run_experiment workflow using simChef. Expected inputs to and outputs from user-defined functions are also provided.

Additional Features

In addition to the ease of parallelization, simChef enables caching of results to further alleviate the computational burden. Here, users can choose to save the experiment's results to disk by

passing `save = TRUE` to `run_experiment`. Once saved, the user can add new DGP and Method objects to the experiment and compute additional replicates without re-computing existing results via the `use_cached` option. Considering the example above, when we add `new_method` and call `run_experiment` with `use_cached = TRUE`, `simChef` finds that the cached results are missing combinations of `new_method`, existing DGPs, and their associated parameters, giving nine new configurations. Replicates for the new combinations are then appended to the cached results.

`simChef` also provides users with a convenient API to automatically generate an R Markdown document. This documentation gathers the scientific details, summary tables, and visualizations side-by-side with the user's custom source code and parameters for data-generating processes, statistical methods, evaluation metrics, and plots. A call to `init_docs` generates empty markdown files for the user to populate with their overarching simulation objectives and with descriptions of each of the DGP, Method, Evaluator, and Visualizer objects included in the Experiment. Finally, a call to `render_docs` prepares the R Markdown document, either for iterative design and analysis of the simulation or to provide a high-quality overview that can be shared easily. We provide an example of the simulation documentation [here](#). Corresponding R source code is available on [GitHub](#).

Related R packages

A number of existing R packages and projects address needs related `simChef`'s functionality. The `batchtools` package ([Lang et al., 2017](#)) provides abstractions for “problems”, “algorithms”, and “experiments”, similar to `simChef`'s DGP, Method, and Experiment objects, respectively. Additionally, `batchtools` provides a number of utilities for shared-memory and distributed memory computations, including for interacting with high-performance computing cluster schedulers such as Slurm and Torque. `simChef` is able to leverage these utilities for distributed computations via the backends provided by the `future.batchtools` package which is part of the future ecosystem of R packages ([Bengtsson, 2021](#)). Whereas `batchtools` is a general tool for distributed mapping operations, `simChef` specializes in data science simulations and provides additional functionality tailored to that setting including its tidy grammar of simulation experiments, the Evaluator and Visualizer concepts, and automated documentation capabilities discussed above.

Many existing packages aim to simplify the process of creating simulation experiments by reducing coding burden through distributed computing helpers and preset methods for generating, computing, and summarizing simulation replicates. `SimDesign` ([Chalmers, 2020](#)) focuses on Monte Carlo simulation experiments and provides a function `runSimulation` that accepts user-defined `generate`, `analyse`, and `summarise` functions, with support for distributed computation via the `parallel` base R package and `future`. `simulator` ([Bien, 2016](#)) provides a tidy grammar of simulation experiments and highly modular helpers for evaluating and managing simulation outputs, relying on the `parallel` package for distributed computation. Other packages provide a small number of well-tailored helper functions for specific simulation settings or distributed computation, including `simhelpers` ([Joshi & Pustejovsky, 2024](#)), `simTool` ([Scheer, 2020](#)), `parSim` ([Epskamp, 2023](#)), `rsimsum` ([Gasparini, 2018](#)), and `simsalapar` ([Hofert & Mächler, 2016](#)). To our knowledge, no single existing package includes `simChef`'s combination of conceptual modularity, tidy grammar, computational flexibility, simulation workflow management, and automated documentation.

Another category of related packages are those that share conceptual similarities with `simChef` in terms of providing helpful abstractions for the design and analysis of simulation experiments, but at a finer level of detail than `simChef` intends. For example, the package `DeclareDesign` ([Blair et al., 2019](#)) provides various `declare_*` functions for defining and evaluating statistical research questions, with an emphasis on the social sciences. The package `infer` ([Couch et al., 2021](#)) provides a tidy API for statistical inference, providing the ability to specify random variables and their relationships, define a null hypothesis, generate data under that hypothesis,

and calculate distributions of statistics based on that hypothesis. Both of these packages and many of the packages discussed above could be employed in a user's DGP, Method, Evaluator, or Visualizer and deployed via an Experiment to carry out a large-scale simulation with automated documentation in harmony with simChef.

Discussion

While simChef's core functionality focuses on computability (C) – encompassing efficient usage of computational resources, ease of user interaction, reproducibility, and documentation – we emphasize the importance of predictability (P) and stability (S) in data science simulations. The principal goal of simChef is to provide a tool for data scientists to create simulations that incorporate predictability (through fit to real-world data) and stability (through sufficient exploration of uncertainty) in their simulations. In future work, we intend to provide tools that can be flexibly tailored to a user's particular scientific needs and further these goals through automated predictability and stability summaries and documentation.

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