

Package ‘simulator’

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Title An Engine for Running Simulations

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Description A framework for performing simulations such as those common in methodological statistics papers. The design principles of this package are described in greater depth in Bien, J. (2016) “The simulator: An Engine to Streamline Simulations,” which is available at [arXiv:1607.00021](https://arxiv.org/abs/1607.00021).

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'model-class.R' 'draws-class.R' 'reference-classes.R'
'simulation-class.R' 'add-to-simulation.R' 'aggregator-class.R'
'utils.R' 'create.R' 'draws.R' 'evaluate.R' 'metric-class.R'
'method-class.R' 'examples.R' 'extended-method-class.R'
'get-from-simulation.R' 'import_from.R' 'load.R' 'manage.R'
'method-extension-class.R' 'methods.R' 'models.R'
'parallel-draws.R' 'parallel-methods.R' 'parallel.R'
'plot_eval.R' 'plot_eval_by.R' 'plot_evals.R' 'tables.R'
'zzz.R'

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R topics documented:

<code>+,ExtendedMethod,MethodExtension-method</code>	3
<code>+,list,MethodExtension-method</code>	4
<code>+,Method,MethodExtension-method</code>	4
<code>add</code>	5
<code>add_bold</code>	6
<code>aggregate_evals</code>	6
<code>Aggregator-class</code>	6
<code>as.data.frame.Evals</code>	7
<code>as.data.frame.listofEvals</code>	7
<code>as.data.frame.listofModels</code>	8
<code>as.data.frame.Model</code>	8
<code>catsim</code>	9
<code>Component-class</code>	9
<code>create</code>	9
<code>describe</code>	10
<code>draws</code>	10
<code>Draws-class</code>	11
<code>DrawsRef-class</code>	12
<code>evals</code>	12
<code>Evals-class</code>	13
<code>EvalsRef-class</code>	14
<code>evaluate</code>	14
<code>evaluate_internal</code>	15
<code>evaluate_single</code>	16
<code>ExtendedMethod-class</code>	16
<code>generate_model</code>	17
<code>get_contents</code>	18
<code>get_files_not_in_simulations</code>	19
<code>get_model_indices</code>	20
<code>get_relative_path</code>	20
<code>get_simulation_with_all_files</code>	21
<code>load,DrawsRef-method</code>	21
<code>load,EvalsRef-method</code>	22
<code>load,list-method</code>	22
<code>load,ModelRef-method</code>	22
<code>load,OutputRef-method</code>	23
<code>load_draws</code>	23
<code>load_evals</code>	24
<code>load_model</code>	25
<code>load_simulation</code>	25
<code>make_my_example_model</code>	26
<code>memory_as_string</code>	27
<code>Method-class</code>	27
<code>MethodExtension-class</code>	28
<code>Metric-class</code>	28
<code>model</code>	29

Model-class	29
ModelRef-class	30
models_as_data.frame	30
model_names	31
my_example_loss	31
my_example_method	32
new_aggregator	32
new_extended_method	33
new_method	33
new_method_extension	34
new_metric	34
new_model	35
new_simulation	35
output	36
Output-class	37
OutputRef-class	38
plot_eval	38
plot_evals	39
plot_eval_by	41
recycle	43
relabel	43
rename	44
run_extendedmethod_single	44
run_method	45
run_method_single	46
save_simulation	46
simulate_from_model	47
simulate_from_model_single	48
simulate_parallel	48
Simulation-class	49
subset_evals	50
subset_models	50
subset_simulation	51
tabulate_eval	51
\$.Model-method	53

Index**54**

+,ExtendedMethod,MethodExtension-method

Create an ExtendedMethod from an ExtendedMethod and MethodExtension

Description

Create an ExtendedMethod from an ExtendedMethod and MethodExtension

Usage

```
## S4 method for signature 'ExtendedMethod,MethodExtension'
e1 + e2
```

Arguments

e1 an object of class [ExtendedMethod](#)
e2 an object of class [MethodExtension](#)

```
+,list,MethodExtension-method
```

Create a list of ExtendedMethod from a list of Methods and a Method-Extension

Description

Create a list of ExtendedMethod from a list of Methods and a MethodExtension

Usage

```
## S4 method for signature 'list,MethodExtension'
e1 + e2
```

Arguments

e1 a list of objects of class [Method](#) or of class [ExtendedMethod](#)
e2 an object of class [MethodExtension](#)

```
+,Method,MethodExtension-method
```

Create an ExtendedMethod from a Method and MethodExtension

Description

Create an ExtendedMethod from a Method and MethodExtension

Usage

```
## S4 method for signature 'Method,MethodExtension'
e1 + e2
```

Arguments

e1 an object of class [Method](#)
e2 an object of class [MethodExtension](#)

add *Add a reference to a simulation*

Description

Adds a ModelRef, DrawsRef, OutputRef, or EvalsRef to a simulation object. To add a DrawsRef, the corresponding ModelRef must already be added. Likewise, to add an OutputRef, the corresponding DrawsRef must already be added. And to add an EvalsRef, the corresponding OutputRef must be added. One can also pass a list of such objects.

Usage

```
add(sim, ref, ...)  
  
## S4 method for signature 'Simulation,ModelRef'  
add(sim, ref, update_saved = TRUE)  
  
## S4 method for signature 'Simulation,DrawsRef'  
add(sim, ref, update_saved = TRUE)  
  
## S4 method for signature 'Simulation,OutputRef'  
add(sim, ref, update_saved = TRUE)  
  
## S4 method for signature 'Simulation,EvalsRef'  
add(sim, ref, update_saved = TRUE)  
  
## S4 method for signature 'Simulation,list'  
add(sim, ref, update_saved = TRUE)
```

Arguments

sim	simulation being added to
ref	the reference object being added
...	not used
update_saved	default is TRUE. Determines whether change to simulation object should be saved to file

Details

The modified simulation object is saved to file if update_saved is TRUE.

<code>add_bold</code>	<i>Make a string bold in a certain format</i>
-----------------------	---

Description

For example, in latex it would take "2" and output "\bf 2"; in html it would output "2".

Usage

```
add_bold(str, output_type)
```

Arguments

<code>str</code>	string or strings (character) to make bold
<code>output_type</code>	output type (see <code>knitr::kable</code> 's format)

<code>aggregate_evals</code>	<i>Apply aggregator to a list of Evals objects</i>
------------------------------	--

Description

Returns a `num_models` by `num_methods` matrix

Usage

```
aggregate_evals(evals_list, aggregator)
```

Arguments

<code>evals_list</code>	a list of Evals objects
<code>aggregator</code>	object of class Aggregator

<code>Aggregator-class</code>	<i>An S4 class for aggregating evaluated metrics</i>
-------------------------------	--

Description

An object of class `Aggregator` consists of a label and a function `aggregate` that has a single argument `ev` that is a list of length equal to the number of draws. This list consists of the evaluated values of all metrics on a single method for a single model.

Slots

<code>label</code>	a human readable label that will be a prefix to the Eval's label
<code>aggregate</code>	a function with argument <code>ev</code> that is a list of length <code>nsim</code> and returns a scalar.

as.data.frame.Evals *Convert an Evals to a data.frame*

Description

This is equivalent to calling `as(x, "data.frame")`

Usage

```
## S3 method for class 'Evals'  
as.data.frame(x, row.names = NULL, optional = FALSE, ...)
```

Arguments

x	object of class Evals
row.names	not used
optional	not used
...	not used

as.data.frame.listofEvals
Convert a list of Evals to a data.frame

Description

When `load` generates a list of Evals, it assigns this to be of (S3) class `listofEvals`, inherited from `list`, so that this function will be invoked instead of `as.data.frame.list`, which is defined in `base`.

Usage

```
## S3 method for class 'listofEvals'  
as.data.frame(x, row.names = NULL, optional = FALSE, ...)
```

Arguments

x	a listofEvals object
row.names	not used
optional	not used
...	not used

```
as.data.frame.listofModels
```

Convert a List of Models to a data.frame

Description

When `load` generates a list of Models, it assigns this to be of (S3) class `listofModels`, inherited from `list`, so that this function will be invoked instead of `as.data.frame.list`, which is defined in `base`.

Usage

```
## S3 method for class 'listofModels'
as.data.frame(x, row.names = NULL, optional = FALSE, ...)
```

Arguments

<code>x</code>	list
<code>row.names</code>	not used
<code>optional</code>	not used
<code>...</code>	not used

```
as.data.frame.Model
```

Convert a Model to a data.frame

Description

Ignores any params that are not length 1 and numeric or character. This is equivalent to calling `as(x, "data.frame")`

Usage

```
## S3 method for class 'Model'
as.data.frame(x, row.names = NULL, optional = FALSE, ...)
```

Arguments

<code>x</code>	object of class <code>Model</code>
<code>row.names</code>	not used
<code>optional</code>	not used
<code>...</code>	not used

catsim	<i>Concatenate and print for the simulator</i>
--------	--

Description

For internal use. This calls `cat` only when `getOption("simulator.verbose")`.

Usage

```
catsim(...)
```

Arguments

... arguments to be passed to `cat`

Component-class	<i>An S4 class representing a component of the simulator.</i>
-----------------	---

Description

This is a virtual class.

Slots

`name` a short name identifier. Must be alphanumeric.

`label` a longer, human readable label that can have other characters such as spaces, hyphens, etc.

create	<i>Create template for a new set of simulations</i>
--------	---

Description

This function is the fastest way to get started. Creates the skeleton of a simulation.

Usage

```
create(dir = "./my_sims")
```

Arguments

`dir` where to create the skeleton of a new set of simulations

Examples

```
## Not run:
  create("./examples")

## End(Not run)
```

describe	<i>Describe the contents of a simulator directory</i>
----------	---

Description

Describe the contents of a simulator directory

Usage

```
describe(dir = ".")
```

Arguments

dir	name of the directory where directory named "files" exists
-----	--

draws	<i>Get one or more draws from a simulation</i>
-------	--

Description

Returns either the draws objects themselves or references to them. See [model](#) function for more information on the `...` and `subset` arguments, which are used to specify a subset of the models.

Usage

```
draws(sim, ..., subset = NULL, index, reference = FALSE)
```

Arguments

sim	a simulation object
...	logical conditions to specify a subset of models. Conditions can only involve params of model that have length 1 and are of class numeric or character.
subset	a vector of integers indexing the models or a vector of model names. To select models based on parameter values, use <code>...</code> . However, using <code>...</code> is slower than using <code>subset</code> .
index	a vector of positive integers specifying which draws objects are desired. If missing, then all draws' outputs are returned.
reference	whether to return the ModelRef or the Model object itself

Examples

```
## Not run:
# suppose previously we had run the following:
sim <- new_simulation(name = "normal-example",
                     label = "Normal Mean Estimation",
                     dir = tempdir()) %>%
  generate_model(make_my_example_model, n = 20) %>%
  simulate_from_model(nsim = 50, index = 1:3)
# then we could get the simulated draws as follows:
d <- draws(sim)
d@draws$r1.1 # first random draw

## End(Not run)
```

 Draws-class

An S4 class representing the random draws from a Model object.

Description

An object of class `Draws` represents the randomly drawn simulated data that is generated when `simulate_from_model` is called on an object of class `Model`. In particular, it contains a named list of `nsim` simulated draws from a model object. The `Model` object's `simulate` function populates this list.

Details

This class inherits from the `Component` class.

Slots

`name` a short name identifier. Must be alphanumeric. Should use the name of the `Model` object that generated it.

`label` a longer, human readable label that indicates what has been randomly drawn.

`draws` a list with `nsim` elements as created by calling the `simulate` function of a `Model` object. This is a named list with each element labeled as `ri.j` where `i` is the index and `j` ranges from 1 to `nsim`. The names are assigned by `simulate_from_model`.

`index` an integer-valued numeric that indicates which block of random draws this is

DrawsRef-class	<i>An S4 class representing a reference to an object of class Draws.</i>
----------------	--

Description

This identifies the necessary information to locate a saved object of class [Draws](#).

Slots

dir directory where the directory `getOption("simulator.files")` is that contains the referenced [Model](#) object
 model_name name of the referenced [Model](#) object
 index the index of the referenced [Draws](#) object. Can alternately be a vector of such indices.
 simulator.files simulator functions will use `getOption("simulator.files")` if simulator.files not provided.

evals	<i>Get one or more evals from a simulation</i>
-------	--

Description

Returns either the Evals object itself or a reference to it.

Usage

```
evals(sim, ..., subset = NULL, index, methods, reference = FALSE)
```

Arguments

sim	a simulation object
...	logical conditions to specify a subset of models. Conditions can only involve params of model that have length 1 and are of class numeric or character.
subset	a vector of integers indexing the models or a vector of model names. To select models based on parameter values, use <code>...</code> . However, using <code>...</code> is slower than using subset.
index	a vector of positive integers specifying which draws' objects are desired. If missing, then all draws' evals are returned.
methods	character vector of method names of interest. If missing, then all methods' evals are returned
reference	whether to return the ModelRef or the Model object itself

See Also

[as.data.frame](#)

Examples

```
## Not run:
# suppose previously we had run the following:
sim <- new_simulation(name = "normal-example",
                    label = "Normal Mean Estimation",
                    dir = tempdir()) %>%
  generate_model(make_my_example_model, n = 20) %>%
  simulate_from_model(nsim = 50, index = 1:3) %>%
  run_method(my_example_method) %>%
  evaluate(my_example_loss)
# then we could get the metric evaluated on the method's output:
e <- evals(sim)
# we can export it as a data.frame
as.data.frame(e)
# or we can get at a particular draw-method-metric triplet
e@evals$`my-method`$r1.1$myloss

## End(Not run)
```

Evals-class

An S4 class representing the evaluation of a metric run by simulator.

Description

An object of class `Evals` consists of information to identify the model, draws, method, and metric objects this output was derived from. It also has a list called `evals`, which is where the output of the metric is stored. Currently, the labels of all these objects are also included so that plot functions can use human-readable labels without requiring re-loading these.

Slots

`model_name` the name of the [Model](#) object this output is derived from.
`model_label` the label of the [Model](#) object this output is derived from.
`index` the index of the [Draws](#) object this output is derived from.
`method_name` the name of the [Method](#) object this output is derived from.
`method_label` the label of the [Method](#) object this output is derived from.
`metric_name` the name of the [Metric](#) object this output is derived from.
`metric_label` the label of the [Metric](#) object this output is derived from.
`evals` a named list with each element labeled by a `method_name` each `evals[[m]]` is itself a named list with each element labeled as `ri.j` where `i` is the index and `j` ranges from 1 to `nsim`. Element `out$ri.j` is output of metric `metric_name` on random draw `ri.j`.

See Also

[evaluate as.data.frame.Evals](#)

EvalsRef-class	<i>An S4 class representing a reference to an object of class Evals</i>
----------------	---

Description

This identifies the necessary information to locate a saved object of class `Evals`. Note that `metric_names` is not needed to identify an `Evals` object since `Evals` objects combine all metrics together into a single file and object.

Slots

`dir` directory where the directory `getOption("simulator.files")` is that contains the referenced `Model` object

`model_name` name of the referenced `Model` object

`index` the index of the referenced `Draws` object.

`method_name` the name of the `Method` object this output is derived from.

`out_loc` a length-1 character vector that gives location (relative to model's path) that method outputs are stored. This can be useful for staying organized when multiple simulations are based on the same `Model` and `Draws` objects.

`simulator.files` simulator functions will use `getOption("simulator.files")` if `simulator.files` not provided.

evaluate	<i>Evaluate outputs of methods according to provided metrics.</i>
----------	---

Description

Given a `Metric` object or list of `Metric` objects, this function evaluates an `Output` object according to these metrics. The computed values of the metrics are saved to file. The "user" time to run the method (as measured by `system.time`) is added to metrics by default unless one of the passed metrics has name "time".

Usage

```
evaluate(object, metrics)
```

Arguments

`object` object of class `OutputRef` as produced by `run_method` (or list of such objects). If `object` is a `Simulation`, then function is applied to the referenced outputs in that simulation and returns the same `Simulation` object but with references added to the new evals created.

`metrics` a list of `Metric` objects or a single `Metric` object.

Details

This function creates objects of class `Evals` and saves each to file (at `dir/model_name/<out_loc>/r<index>_<method_name>`). Since evaluating metrics is usually (in statistical methodological papers) fast, parallel functionality has not been developed for the evaluation component.

See Also

[generate_model](#) [simulate_from_model](#) [run_method](#)

Examples

```
## Not run:
# suppose previously we had run the following:
sim <- new_simulation(name = "normal-example",
                    label = "Normal Mean Estimation",
                    dir = tempdir()) %>%
  generate_model(make_my_example_model, n = 20) %>%
  simulate_from_model(nsim = 50, index = 1:3) %>%
  run_method(my_example_method)
# then we could add
sim <- evaluate(sim, my_example_loss)

## End(Not run)
```

evaluate_internal	<i>Evaluate outputs of methods according to provided metrics.</i>
-------------------	---

Description

Given a `Metric` object or list of `Metric` objects, this function evaluates an `Output` object according to these metrics. The computed values of the metrics are saved to file.

Usage

```
evaluate_internal(
  metrics,
  dir = ".",
  model_name,
  index,
  method_names,
  out_loc = "out"
)
```

Arguments

metrics	a list of <code>Metric</code> objects or a single <code>Metric</code> object
dir	the directory where <code>Model</code> object was saved (by generate_model)

model_name	the Model object's name attribute
index	the index of a computed Draws object. Can alternately be a vector of such indices.
method_names	the Method objects' name attributes as a character vector.
out_loc	(optional) a length-1 character vector that gives location (relative to model's path) that method outputs are stored.

Details

This function creates objects of class [Evals](#) and saves each to file (at dir/model_name/<out_loc>/r<index>_<method_name>_). Since evaluating metrics is usually (in statistical methodological papers) fast, parallel functionality has not been developed for the evaluation component.

evaluate_single *Run one or more metrics on outputs.*

Description

This is an internal function. Users should call the wrapper function [evaluate](#). Here "single" refers to a single output (and thus a single method, though not necessarily a single index). The metrics provided are run and saved together in a file.

Usage

```
evaluate_single(metrics, model, output, draws = NULL)
```

Arguments

metrics	a list of Metric objects
model	a Model object
output	a Output object
draws	(optional) a Draws object or NULL

ExtendedMethod-class *An S4 class representing the extension of a method*

Description

An object of class [ExtendedMethod](#) is like a [Method](#) except it uses the output of another method in addition to the [Model](#) and [Draws](#). We can also form chains of [ExtendedMethod](#)'s, in which one [ExtendedMethod](#) is taken to be the "base_method" of a subsequent [ExtendedMethod](#). This means that the latter [ExtendedMethod](#) would use the output of the former [ExtendedMethod](#).

Details

While one can create an [ExtendedMethod](#) from scratch, typically it will be cleaner to write a [MethodExtension](#) object and then use the addition operator: `my_extended_method = my_base_method + my_method_extension`. For example, if `my_base_method` is the lasso, `my_method_extension` might be cross-validation, and the resulting `my_extended_method` would be the lasso with tuning parameter chosen by cross-validation. The advantage is that if we have several methods, we only have to write the cross-validation [MethodExtension](#) object once.

For an example in which one has a chain of [ExtendedMethod](#)'s, consider the lasso example in which we have a [MethodExtension](#) called, say, `refit`, which takes the nonzeros from the lasso's output and performs least squares on these selected variables. Let `cv` be another [MethodExtension](#). Then, `refitted_lasso = lasso + refit` is an [ExtendedMethod](#) and `refitted_lasso + cv` is as well.

This class inherits from the [Component](#) class.

Slots

`name` a short name identifier. Must be alphanumeric.

`label` a longer, human readable label that can have other characters such as spaces, hyphens, etc.

`base_method` a list of length 1 containing the object of class [Method](#) or [ExtendedMethod](#) that is being extended

`extended_method` a function with arguments "model", "draw", "out", and "base_method".

generate_model

Generate a model.

Description

This function executes the `make_model` function provided by the user and writes to file the resulting [Model](#) object(s). For example, when simulating regression with a fixed design, `X` would be generated in this function and `n`, `p`, `beta`, and `sigma` would also be specified.

Usage

```
generate_model(object = ".", make_model, ..., seed = 123, vary_along = NULL)
```

Arguments

<code>object</code>	the name of the directory where directory named "files" exists (or should be created) to save Model object in. Default is current working directory. Or can be an object of class Simulation , in which case the <code>object@dir</code> is used and a simulation object is returned instead of an object of class ModelRef .
<code>make_model</code>	a function that outputs an object of class Model . Or a list of such functions.
<code>...</code>	optional parameters that may be passed to <code>make_model</code>
<code>seed</code>	an integer seed for the random number generator.
<code>vary_along</code>	character vector with all elements contained in <code>names(...)</code> See description for more details.

Details

When `make_model` has arguments, these can be passed using `...`. These will be passed directly to `make_model` except for any arguments named in `vary_along`. These arguments should be lists and a separate model will be created for each combination of elements in these lists. For example, if `vary_along = c("n", "p")`, then we can pass `n=as.list(c(50, 100, 150))` and `p=as.list(c(10, 100))` and 6 models will be created, one for each pair of `n` and `p`. For each pair (`n,p`), a distinct extension is added to the end of the model name. This extension is generated using a hash function so that different values of the `vary_along` parameters will lead to different model name extensions. This ensures that if one later decides to add more values of the `vary_along` parameters, this will not lead to pre-existing files being overwritten (unless the same values of the `vary_along` combination are used again).

If `object` is a directory name, the function returns a reference or list of references to the model(s) generated. If `object` is a `Simulation`, then function returns the same `Simulation` object but with references added to the new models created. These changes to the `Simulation` object are saved to file.

`make_model` is called generating an object of class `Model`, called `model`, which is saved to `dir/name/model.Rdata` (where `name` is the name attribute of `model`). This file also contains the random number generator state and other information such as the function `make_model` itself and the date when `model` was created.

See Also

[new_model](#) [simulate_from_model](#) [run_method](#)

Examples

```
# initialize a new simulation
sim <- new_simulation(name = "normal-example",
                    label = "Normal Mean Estimation",
                    dir = tempdir())

# generate a model (and add it to the simulation)
sim <- generate_model(sim, make_my_example_model, n = 20)
# generate a sequence of models (and add them to the simulation)
sim <- generate_model(sim, make_my_example_model,
                    n = list(10, 20, 30),
                    vary_along = "n")
```

get_contents

Get the contents of a simulator directory

Description

This function gives detailed information about what is being stored in the "files" directory. In particular, it gives the complete paths for all the draws, outputs, and evals files. This can be useful in situations in which the draws or outputs files are no longer needed and take up a lot of memory. In such a case a user could delete these files with a command such as `system(paste(c("rm", contents$out_files), collapse = " "))`. That said, one must be cautious in deleting these files

since the simulator generally assumes that earlier stages' files will be available and so deleting these may cause errors. However, if one is essentially finished with a simulation and evaluated metrics have been computed and if the methods' raw outputs are taking up a lot of disk space, then one might consider deleting the out_files (and/or the draws_files).

Usage

```
get_contents(dir = ".", out_loc = "out")
```

Arguments

dir	name of the directory where directory named "files" exists
out_loc	a length-1 character vector that gives location (relative to model's path) that method outputs are stored. This can be useful for staying organized when multiple simulations are based on the same Model and Draws objects. Usually this is just "out"

```
get_files_not_in_simulations
```

Find files in simulator directory not referred to by any simulations

Description

Once one has completed all simulation studies, this function can be called to identify any files that may have been created along the way that are no longer being used in any simulations. It would then be safe to delete these files.

Usage

```
get_files_not_in_simulations(dir, out_loc = "out")
```

Arguments

dir	name of the directory where directory named "files" exists
out_loc	a length-1 character vector that gives location (relative to model's path) that method outputs are stored. This can be useful for staying organized when multiple simulations are based on the same Model and Draws objects. Usually this is just "out"

get_model_indices	Returns indices of a specified subset of sim@model_refs
-------------------	---

Description

See [model](#) for information about the various formats of subset.

Usage

```
get_model_indices(sim, subset)
```

Arguments

sim	a simulation object
subset	a vector indicating which models should be returned.

get_relative_path	Get relative path
-------------------	-------------------

Description

Given a base path and a specific path, returns a string str such that file.path(base_path, str) is the same location as path.

Usage

```
get_relative_path(base_path, path)
```

Arguments

base_path	the base path
path	a specific path

`get_simulation_with_all_files`

Returns a simulation object containing references to all files in directory

Description

Returns a simulation object containing references to all files in directory

Usage

```
get_simulation_with_all_files(dir, out_loc = "out")
```

Arguments

<code>dir</code>	name of the directory where directory named "files" exists
<code>out_loc</code>	a length-1 character vector that gives location (relative to model's path) that method outputs are stored. This can be useful for staying organized when multiple simulations are based on the same Model and Draws objects. Usually this is just "out"

`load,DrawsRef-method` *Load a DrawsRef*

Description

Load a DrawsRef

Usage

```
## S4 method for signature 'DrawsRef'  
load(file)
```

Arguments

<code>file</code>	object to load
-------------------	----------------

load,EvalsRef-method *Load an EvalsRef*

Description

Load an EvalsRef

Usage

```
## S4 method for signature 'EvalsRef'  
load(file)
```

Arguments

file object to load

load,list-method *Load a list of reference objects*

Description

Load a list of reference objects

Usage

```
## S4 method for signature 'list'  
load(file)
```

Arguments

file list of objects to load

load,ModelRef-method *Load a ModelRef*

Description

Load a ModelRef

Usage

```
## S4 method for signature 'ModelRef'  
load(file)
```

Arguments

file object to load

load, OutputRef-method *Load an OutputRef*

Description

Load an OutputRef

Usage

```
## S4 method for signature 'OutputRef'
load(file)
```

Arguments

file object to load

load_draws *Load one or more draws objects from file.*

Description

After [simulate_from_model](#) has been called, this function can be used to load one or more of the saved [Draws](#) object(s) (along with RNG information). If multiple indices are provided, these will be combined into a new single [Draws](#) object. If simulation object is available, it is easier to use the function [draws](#) to load it.

Usage

```
load_draws(dir, model_name, index, more_info = FALSE, simulator.files = NULL)
```

Arguments

dir the directory passed to [generate_model](#))

model_name the Model object's name attribute

index a vector of positive integers.

more_info if TRUE, then returns additional information such as state of RNG after calling [generate_model](#)

simulator.files if NULL, then `getOption("simulator.files")` will be used.

See Also

[simulate_from_model](#) [draws](#)

load_evals

Load one or more Evals objects from file.

Description

After `evaluate` has been called, this function can be used to load one or more of the saved `Evals` object(s). If multiple indices are provided, these will be combined by index into a new single `Evals` object. If multiple methods are provided, a list of `Evals` objects will be returned.

Usage

```
load_evals(
  dir,
  model_name,
  index,
  method_names,
  metric_names = NULL,
  out_loc = "out",
  simulator.files = NULL
)

load_evals_from_ref(ref, metric_names = NULL)
```

Arguments

<code>dir</code>	the directory passed to <code>generate_model</code>)
<code>model_name</code>	the <code>Model</code> object's name
<code>index</code>	a vector of positive integers.
<code>method_names</code>	the name of one or more <code>Method</code> objects.
<code>metric_names</code>	(optional) a character vector of which elements of evals should be loaded. If <code>NULL</code> , then all elements are loaded.
<code>out_loc</code>	only needed if it was used in call to
<code>simulator.files</code>	if <code>NULL</code> , then <code>getOption("simulator.files")</code> will be used. <code>run_method</code> .
<code>ref</code>	an object of class <code>EvalsRef</code>

See Also

`load_model` `load_draws` as `data.frame.Evals`

load_model	<i>Load a model from file.</i>
------------	--------------------------------

Description

After [generate_model](#) has been called, this function can be used to load the saved [Model](#) object (along with the RNG state and other information if desired).

Usage

```
load_model(dir, model_name, more_info = FALSE, simulator.files = NULL)
```

Arguments

dir	the directory passed to generate_model)
model_name	the Model object's name attribute
more_info	if TRUE, then returns additional information such as state of RNG after calling generate_model
simulator.files	if NULL, then <code>getOption("simulator.files")</code> will be used.

Details

Depending on `more_info`, either returns [Model](#) object or a list containing [Model](#) object and other information. If simulation object is available, it is easier to use the function [model](#) to load the model.

See Also

[generate_model](#) [model](#)

load_simulation	<i>Load a simulation object</i>
-----------------	---------------------------------

Description

Loads an object of class [Simulation](#). Note that `dir` gives the directory where the Simulation object is stored. Thus, if the working directory is different from the working directory when the Simulation object was created, then `dir` will be different from the one passed to [new_simulation](#).

Usage

```
load_simulation(name, dir = ".")
```

Arguments

name a short name identifier. Must be alphanumeric.
dir directory that contains "files" directory for this simulation

See Also

[new_simulation](#) [save_simulation](#)

Examples

```
sim <- new_simulation(name = "normal-example",  
                      label = "Normal Mean Estimation",  
                      dir = tempdir())  
rm(sim)  
sim <- load_simulation("normal-example", dir = tempdir())
```

make_my_example_model *Make My Example Model*

Description

This function is used in the examples. It returns a [Model](#) object. In particular, it represents n i.i.d. draws from a normal with mean 2 and variance 1.

Usage

```
make_my_example_model(n)
```

Arguments

n number of i.i.d. draws

See Also

[my_example_method](#) [my_example_loss](#)

memory_as_string	<i>Write memory in human readable way</i>
------------------	---

Description

Write memory in human readable way

Usage

```
memory_as_string(memory_in_bytes)
```

Arguments

memory_in_bytes
the amount of memory in Bytes.

Method-class	<i>An S4 class representing a method to be run by simulator.</i>
--------------	--

Description

An object of class Method consists of a name, label, and a function method that takes arguments model and draw. A draw refers to a single element of the list in an object of class [Draws](#).

Details

This class inherits from the [Component](#) class.

Slots

name a short name identifier. Must be alphanumeric.

label a longer, human readable label that can have other characters such as spaces, hyphens, etc.

settings (optional) a list of "settings" for the method (e.g., tuning parameters or related information that might distinguish two otherwise identical methods).

method a function that has arguments "model", "draw" and (optionally) names matching elements within names(settings)

`MethodExtension-class` *An S4 class used to create an extended version of a method*

Description

An object of class `MethodExtension` when added to a `Method` creates a [ExtendedMethod](#).

Details

This class inherits from the [Component](#) class.

Slots

`name` a short name identifier. Must be alphanumeric.

`label` a longer, human readable label that can have other characters such as spaces, hyphens, etc.

`method_extension` a function with arguments "model", "draw", "out", and "base_method". This will become the function `extended_method` in the `ExtendedMethod` object that is created.

`Metric-class` *An S4 class representing an evaluation metric to be used by simulator.*

Description

An object of class `Metric` consists of a name, label, and a function `metric` that takes arguments `model` (of class [Model](#)) and `out` (of class [Output](#)), which is the output of a method.

Details

This class inherits from the [Component](#) class.

Slots

`name` a short name identifier. Must be alphanumeric.

`label` a longer, human readable label that can have other characters such as spaces, hyphens, etc.

`metric` a function with arguments "model" and "out" (and optionally "draw")

model	<i>Get one or more models from a simulation</i>
-------	---

Description

Returns either the models themselves or references to them.

Usage

```
model(sim, ..., subset = NULL, reference = FALSE)
```

Arguments

sim	a simulation object
...	logical conditions to specify a subset of models. Conditions can only involve params of model that have length 1 and are of class numeric or character.
subset	a vector of integers indexing the models or a vector of model names. To select models based on parameter values, use ... However, using ... is slower than using subset.
reference	whether to return the ModelRef or the Model object itself

Details

There are two main ways to specify a subset of the models. (1) The easiest way is by writing a conditional expression involving the parameters and passing it through ... For example, $n > 100$ & $p \leq 20$. Only parameters that are length one and either numeric or character can be used in these expressions. (2) The faster way to retrieve a subset of models is to use the subset argument. This can be either a set of numerical values (specifying which models to load based on the order in which the models are stored in the simulation object. This order can be ascertained by printing the simulation object.) or as a set of a character vector of the model names desired.

While approach (1) is very convenient, it requires loading all models from file. This may be slow in situations in which there are a lot of models and/or the models are large and thus slow to load.

Model-class	<i>An S4 class representing the model component of the simulator.</i>
-------------	---

Description

An object of class Model specifies the statistical model. In particular, all parameters are specified in addition to a function called simulate that allows one to draw random samples from this model.

Details

To get parameters stored in a Model object, a shortcut for `my_model@params$my_parameter` is `my_model$my_parameter`.

This class inherits from the [Component](#) class.

Slots

`name` a short name identifier. Must be alphanumeric (though -, _, and / are allowed as long as they are not at the start or end of name).

`label` a longer, human readable label that can have other characters such as spaces, hyphens, etc.

`params` a list that contains the Model object's parameters

`simulate` a function that has arguments `nsim` and names matching elements within `names(params)`. It returns a list of length `nsim`, where each element of the list represents a random draw from the Model object.

ModelRef-class	<i>An S4 class representing a reference to an object of class Model.</i>
----------------	--

Description

This identifies the necessary information to locate a saved object of class [Model](#).

Slots

`dir` directory where the directory "files" is that contains the referenced [Model](#) object

`name` a short name identifier.

`label` a longer, human readable label that can have other characters

`simulator.files` simulator functions will use `getOption("simulator.files")` if `simulator.files` not provided.

models_as_data.frame	<i>Convert a list of Model objects into a data.frame</i>
----------------------	--

Description

Ignores any params that are not length 1 and numeric or character

Usage

```
models_as_data.frame(m)
```

Arguments

`m` model object

model_names	<i>Get model names in a Simulation</i>
-------------	--

Description

Get model names in a Simulation

Usage

```
model_names(sim)
```

Arguments

sim object of class [Simulation](#)

my_example_loss	<i>My Example Loss</i>
-----------------	------------------------

Description

This [Metric](#) object is used in the examples. It is squared error loss.

Usage

```
my_example_loss
```

Format

An object of class `Metric` of length 1.

See Also

[make_my_example_model](#) [my_example_loss](#)

my_example_method	<i>My Example Method</i>
-------------------	--------------------------

Description

This [Method](#) object is used in the examples. It is the sample mean of the data.

Usage

```
my_example_method
```

Format

An object of class Method of length 1.

See Also

[make_my_example_model](#) [my_example_loss](#)

new_aggregator	<i>Create an Aggregator object</i>
----------------	------------------------------------

Description

Creates a new [Aggregator](#) object.

Usage

```
new_aggregator(label, aggregate)
```

Arguments

label	a human readable label
aggregate	a function with argument <code>ev</code> that is a list of length equal to the number of draws with each element itself being a named list. Each element of this list corresponds to a metric that has been computed. In particular, given an Evals object <code>o</code> , <code>aggregate</code> takes as input <code>o@evals[[method_name]]</code> (which is a list of the kind just described). The function <code>aggregate</code> should return a scalar.

new_extended_method *Create an ExtendedMethod object*

Description

Creates a new [ExtendedMethod](#) object.

Usage

```
new_extended_method(name, label, base_method, extended_method)
```

Arguments

name	a short name identifier. Must be alphanumeric.
label	a longer, human readable label that can have other characters such as spaces, hyphens, etc.
base_method	the object of class Method or of class Method that is being extended
extended_method	a function with arguments "model", "draw", "out", and "base_method".

new_method *Create a Method object*

Description

Creates a new [Method](#) object.

Usage

```
new_method(name, label, method, settings = list())
```

Arguments

name	a short name identifier. Must be alphanumeric.
label	a longer, human readable label that can have other characters such as spaces, hyphens, etc.
method	a function that has arguments "model", "draw" and (optionally) names matching elements within <code>names(settings)</code>
settings	(optional) a list of "settings" for the method (e.g., tuning parameters or related information that might distinguish two otherwise identical methods).

new_method_extension	<i>Create an object that can be used to make an extended version of a method</i>
----------------------	--

Description

Creates an object of class `MethodExtension`, which when added to a `Method` creates an `ExtendedMethod`.

Usage

```
new_method_extension(name, label, method_extension)
```

Arguments

name	a short name identifier. Must be alphanumeric.
label	a longer, human readable label that can have other characters such as spaces, hyphens, etc.
method_extension	a function with arguments "model", "draw", "out", and "base_method". This will become the function <code>extended_method</code> in the <code>ExtendedMethod</code> object that is created.

Details

This class inherits from the `Component` class.

new_metric	<i>Create a Metric object</i>
------------	-------------------------------

Description

Creates a new `Metric` object.

Usage

```
new_metric(name, label, metric)
```

Arguments

name	a short name identifier. Must be alphanumeric.
label	a longer, human readable label that can have other characters such as spaces, hyphens, etc.
metric	a function with arguments "model" and "out" (and optionally "draw")

new_model *Create a Model object*

Description

Creates a new [Model](#) object.

Usage

```
new_model(name, label, params = list(), simulate)
```

Arguments

name	a short name identifier. Must be alphanumeric (though -, _, and / are allowed as long as they are not at the start or end of name).
label	a longer, human readable label that can have other characters such as spaces, hyphens, etc.
params	a list that contains the Model object's parameters
simulate	a function that has arguments <code>nsim</code> and names matching elements within <code>params</code> . It returns a list of length <code>nsim</code> , where each element of the list represents a random draw from the Model object.

Examples

```
make_my_example_model <- function(n) {
  new_model(name = "normal-data",
            label = sprintf("Normal (n = %s)", n),
            params = list(n = n, mu = 2),
            simulate = function(n, mu, nsim) {
              # this function must return a list of length nsim
              x <- matrix(rnorm(n * nsim), n, nsim)
              x <- mu + x # true mean is mu
              return(split(x, col(x))) # make each col its own list element
            })
}
```

new_simulation *Make a new simulation object*

Description

Creates an object of class [Simulation](#). In addition to having a name and label, this object consists of a set of references to objects of class [ModelRef](#), [DrawsRef](#), [OutputRef](#), and [EvalsRef](#).

Usage

```
new_simulation(name, label, dir = ".", refs = list(), save_to_file = TRUE)
```

Arguments

name	a short name identifier. Must be alphanumeric.
label	a longer, human readable label that can have other characters such as spaces, hyphens, etc.
dir	a directory that reference's directories are relative to
refs	a list containing objects of class ModelRef , DrawsRef , OutputRef , and EvalsRef
save_to_file	whether this new simulation should be saved to file. Default is TRUE. If TRUE, then this simulation can be loaded in a new R session using <code>dir</code> and <code>name</code> .

Details

A Simulation object is the basic unit of a simulation study. Roughly, one can think of it as all the files relevant to a single figure. This might be a single plot or a series of related plots/panels. It could also correspond to a single table. Note that a Simulation object is light-weight even for large simulations because it only stores references to the objects not the objects themselves. The functions [model](#), [draws](#), [output](#), [evals](#) can be used to load individual objects of a simulation.

The Simulation object created is saved to a file so that it can be loaded in a new R session. The simulation is saved in `dir/files/name.Rdata`. Note: while "files" is the default, the name of this directory is from `getOption("simulator.files")`, which is the value of `getOption("simulator.files")` when the model was created.

See Also

[load_simulation](#) [save_simulation](#)

Examples

```
sim <- new_simulation(name = "normal-example",
                     label = "Normal Mean Estimation",
                     dir = tempdir())
```

output	<i>Get one or more outputs from a simulation</i>
--------	--

Description

Returns either the output object itself or a reference to it.

Usage

```
output(sim, ..., subset = NULL, index, methods, reference = FALSE)
```

Arguments

sim	a simulation object
...	logical conditions to specify a subset of models. Conditions can only involve params of model that have length 1 and are of class numeric or character.
subset	a vector of integers indexing the models or a vector of model names. To select models based on parameter values, use ... However, using ... is slower than using subset.
index	a vector of positive integers specifying which draws' objects are desired. If missing, then all draws' outputs are returned.
methods	character vector of method names of interest. If missing, then all methods' outputs are returned
reference	whether to return the ModelRef or the Model object itself

Examples

```
## Not run:
# suppose previously we had run the following:
sim <- new_simulation(name = "normal-example",
                    label = "Normal Mean Estimation",
                    dir = tempdir()) %>%
  generate_model(make_my_example_model, n = 20) %>%
  simulate_from_model(nsim = 50, index = 1:3) %>%
  run_method(my_example_method)
# then we could get the method's output as follows:
o <- output(sim)
o@out$r1.1 # first random draw's output

## End(Not run)
```

Output-class

An S4 class representing the output of a method run by simulator.

Description

An object of class Output consists of information to identify the model, draws, and method objects this output was derived from. It also has a list called out, which is where the output of the method is stored.

Slots

model_name the name of the [Model](#) object this output is derived from.

index the index of the [Draws](#) object this output is derived from.

method_name the name of the [Method](#) object this output is derived from.

method_label the label of the [Method](#) object this output is derived from.

out a named list with each element labeled as $r_{i.j}$ where i is the index and j ranges from 1 to nsim. Element $out\$r_{i.j}$ is output of method method_name on random draw $r_{i.j}$.

OutputRef-class	<i>An S4 class representing a reference to an object of class Output.</i>
-----------------	---

Description

This identifies the necessary information to locate a saved object of class `Output`.

Slots

`dir` directory where the directory `getOption("simulator.files")` is that contains the referenced `Model` object

`model_name` name of the referenced `Model` object

`index` the index of the referenced `Draws` object. Can alternately be a vector of such indices.

`method_name` the name of the `Method` object this output is derived from.

`out_loc` a length-1 character vector that gives location (relative to model's path) that method outputs are stored. This can be useful for staying organized when multiple simulations are based on the same `Model` and `Draws` objects.

`simulator.files` simulator functions will use `getOption("simulator.files")` if `simulator.files` not provided.

plot_eval	<i>Plot a metric's value for each method</i>
-----------	--

Description

When the evaluated metric is scalar-valued, this functions makes a boxplot of this metric for each method. When the metric is vector-valued, this function makes a curve with this metric on the y-axis, with one curve for each method (the x-axis is the corresponding entry of that metric's vector). If `evals` is a `listofEvals`, then each model will be its own plot.

Usage

```
plot_eval(
  object,
  metric_name,
  use_ggplot2 = TRUE,
  main,
  facet_mains,
  ylab,
  ylim,
  include_zero = FALSE,
  angle = 0,
  ...
)
```

Arguments

object	an object of class Simulation , Evals , or listofEvals
metric_name	the name of a metric to plot
use_ggplot2	whether to use ggplot2 (requires installation of ggplot2)
main	title of plot. Default is <code>model_label</code> when <code>evals</code> is a single Evals .
facet_mains	only to be used when <code>evals</code> is a listofEvals and should be of the same length. Default will be the <code>model_label</code> for each model.
ylab	the y-axis label (default is <code>metric_label</code>)
ylim	the y-axis limits to use (across all plots)
include_zero	whether <code>ylim</code> should include 0. Ignored if <code>ylim</code> is passed explicitly
angle	angle of labels (only when <code>use_ggplot2 = FALSE</code>)
...	additional arguments to pass to boxplot (only when <code>use_ggplot2 = FALSE</code>).

See Also

[plot_evals](#) [plot_eval_by](#) [tabulate_eval](#)

Examples

```
## Not run:
# suppose previously we had run the following:
sim <- new_simulation(name = "normal-example",
                    label = "Normal Mean Estimation",
                    dir = tempdir()) %>%
  generate_model(make_my_example_model, n = 20) %>%
  simulate_from_model(nsim = 50, index = 1:3) %>%
  run_method(my_example_method) %>%
  evaluate(my_example_loss)
# then we could plot this
plot_eval(sim, "myloss") # "myloss" is my_example_loss@name

## End(Not run)
```

plot_evals

Plot one metric versus another for each method

Description

This function is used when both evaluated metrics are vector-valued, so a curve is plotted, parametrized by the two metrics. To plot a single metric that is vector-valued, pass `NULL` for `metric_name_x`. This behaves similarly to `plot(runif(5))`, in which the x-axis variable is simply `1:5`. If `evals` is a [listofEvals](#), then each model will be its own plot.

Usage

```

plot_evals(
  object,
  metric_name_x,
  metric_name_y,
  use_ggplot2 = TRUE,
  main,
  facet_mains,
  xlab,
  ylab,
  xlim,
  ylim,
  include_zero = FALSE,
  legend_location = "topright",
  method_col = seq(num_methods),
  method_lty = rep(1, num_methods),
  method_lwd = rep(1, num_methods),
  method_pch = rep(NA, num_methods),
  ...
)

```

Arguments

object	an object of class Simulation , Evals , or listofEvals
metric_name_x	the name of metric to plot on x axis (or NULL)
metric_name_y	the name of metric to plot on y axis
use_ggplot2	whether to use ggplot2 (requires installation of ggplot2)
main	title of plot. Default is <code>model_label</code> when <code>evals</code> is a single Evals .
facet_mains	only to be used when <code>evals</code> is a listofEvals and should be of the same length. Default will be the <code>model_label</code> for each model.
xlab	the x-axis label (default is <code>metric_label_x</code>)
ylab	the y-axis label (default is <code>metric_label_y</code>)
xlim	the limits of the x-axis
ylim	the limits of the y-axis
include_zero	whether <code>ylim</code> should include 0. Ignored if <code>ylim</code> is passed explicitly
legend_location	location of legend. Set to NULL to remove legend.
method_col	color to use for each method
method_lty	line style to use for each method
method_lwd	line thickness to use for each method
method_pch	point style to use for each method (default is that no points, only lines are drawn)
...	additional arguments to pass to boxplot (only when <code>use_ggplot2 = FALSE</code>).

plot_eval_by

Plot a metric across multiple values of a model parameter

Description

This function is to be used on simulations in which `generate_model` was called using the `vary_along` parameter. When this is a single (scalar) numeric parameter, a single plot is created in which the x-axis is this parameter. Eventually, this function should handle one or two categorical variables (in which facets are used) and one categorical combined with one continuous variable.

Usage

```
plot_eval_by(
  sim,
  metric_name,
  varying,
  type = c("aggregated", "raw"),
  center_aggregator = NULL,
  spread_aggregator = NULL,
  use_ggplot2 = TRUE,
  main,
  xlab,
  ylab,
  xlim,
  ylim,
  include_zero = FALSE,
  legend_location = "topright",
  method_col = seq(num_methods),
  method_lty = rep(1, num_methods),
  method_lwd = rep(1, num_methods),
  method_pch = rep(1, num_methods),
  ...
)
```

Arguments

<code>sim</code>	an object of class <code>Simulation</code>
<code>metric_name</code>	the name of a metric to plot (ignored if custom aggregator is provided)
<code>varying</code>	character vector giving the name of a parameter that is varied across the models in evals. For now, this parameter must be numeric and there cannot be multiple models having the same value of this parameter.
<code>type</code>	if "aggregated" then shows line with error bars (line represents <code>center_aggregator</code> and error bars represent <code>spread_aggregator</code> ; by default these are sample mean and estimated standard error); if <code>type</code> is "raw" then shows the raw data as points (with smoother overlaid)

center_aggregator	ignored if type is "raw". When NULL (which is default), the sample mean aggregator is used. User can write specialized aggregators (see definition of class Aggregator) as necessary, for example, when the evaluated metric is not scalar-valued.
spread_aggregator	ignored if type is "raw". When NULL (which is default), the sample mean aggregator is used. User can write specialized aggregators (see definition of class Aggregator) as necessary, for example, when the evaluated metric is not scalar-valued. Set spread_aggregator to NA to hide error bars.
use_ggplot2	whether to use ggplot2 (requires installation of ggplot2)
main	title of plot.
xlab	the x-axis label (default is varying)
ylab	the y-axis label (default is metric_label)
xlim	the x-axis limits to use
ylim	the y-axis limits to use
include_zero	whether ylim should include 0. Ignored if ylim is passed explicitly
legend_location	location of legend. Set to NULL to remove legend.
method_col	color to use for each method
method_lty	line style to use for each method
method_lwd	line thickness to use for each method
method_pch	point style to use for each method (default is that no points, only lines are drawn)
...	additional arguments to pass to plot (only when use_ggplot2 = FALSE).

Details

When type is "raw", the individual evals are shown (one point per model-draw-method triplet) along with a loess smooth. When type is "aggregated", then center_aggregator and spread_aggregator are used. center_aggregator is used to draw a single line per method in which the individual evals computed for each draw has been aggregated in some way. By default, the mean_aggregator is used, which simply averages the evals computed across all draws. When spread_aggregator is non-NULL, "error bars" are drawn with (half)widths computed using spread_aggregator. By default, the se_aggregator is used, which gives an estimate of the standard error of the sample mean.

The arguments method_col, method_lty, method_lwd, method_pch only apply when use_ggplot2 is FALSE.

Examples

```
## Not run:
# suppose previously we had run the following:
sim <- new_simulation(name = "normal-example",
                     label = "Normal Mean Estimation",
                     dir = tempdir()) %>%
```

```

generate_model(make_my_example_model,
               n = list(10, 20, 30),
               vary_along = "n") %>%
simulate_from_model(nsim = 50, index = 1:3) %>%
run_method(my_example_method) %>%
evaluate(my_example_loss)
# then we could plot this
plot_eval_by(sim, "myloss", varying = "n", include_zero = TRUE)

## End(Not run)

```

recycle	<i>Recycles elements to create vector of desired length</i>
---------	---

Description

Recycles elements to create vector of desired length

Usage

```
recycle(x, length)
```

Arguments

x	vector to be expanded to proper length
length	desired length

relabel	<i>Give simulation a new label</i>
---------	------------------------------------

Description

Note that [save_simulation](#) needs to be called for this change to be saved to file.

Usage

```
relabel(sim, label)
```

Arguments

sim	object of class Simulation
label	a longer, human readable label that can have other characters such as spaces, hyphens, etc.

See Also

[rename](#)

rename	<i>Give simulation a new name</i>
--------	-----------------------------------

Description

Note that [save_simulation](#) needs to be called for this change to be saved to file.

Usage

```
rename(sim, name)
```

Arguments

sim	object of class Simulation
name	a short name identifier. Must be an alphanumeric (but can also have - or _ within

See Also

[relabel](#)

run_extendedmethod_single	<i>Run a single extended method on a single index of simulated data.</i>
---------------------------	--

Description

This is an internal function. Users should call the wrapper function, [run_method](#). Here "single" refers to a single index-ExtendedMethod pair.

Usage

```
run_extendedmethod_single(extmethod, model, draws, base_output_list)
```

Arguments

extmethod	a ExtendedMethod object
model	a Model object
draws	a Draws object generated by model
base_output_list	the result of loading a Output object with more_info = TRUE so that it includes RNG endstate.

run_method	<i>Run one or more methods on simulated data.</i>
------------	---

Description

Given a [Method](#) object or list of [Method](#) objects, this function runs the method(s) on the draws passed through object. The output of each method is saved to file.

Usage

```
run_method(object, methods, out_loc = "out", parallel = NULL)
```

Arguments

object	an object of class DrawsRef (or a list of such objects) as returned by <code>link{simulate_from_model}</code> . If object is a Simulation , then function is applied to the referenced draws in that simulation and returns the same Simulation object but with references added to the new outputs created.
methods	a list of Method and/or ExtendedMethod objects or a single Method or object ExtendedMethod
out_loc	(optional) a length-1 character vector that gives location (relative to model's path) that method outputs are stored. This can be useful for staying organized when multiple simulations are based on the same Model and Draws objects.
parallel	either <code>NULL</code> or a list containing <code>socket_names</code> and (optionally) <code>libraries</code> and <code>save_locally</code> (see Details for more information)

Details

This function creates objects of class [Output](#) and saves each to file (at `dir/model_name/<out_loc>/r<index>_<method_name>`). If `parallel` is not `NULL`, then it must be a list containing `socket_names`, which can either be a positive integer specifying the number of copies to run on localhost or else a character vector of machine names (e.g., "mycluster-0-0"). The list `parallel` can also contain `libraries`, a character vector of R packages that will be needed on the slaves and `save_locally`, a logical that indicates whether the files generated should be saved on the slaves (i.e., locally) or on the master.

Before running each method on index `i`, the RNG state is restored to what it was at the end of calling `simulate_from_model` on this index. This is only relevant for randomized methods. The choice to do this ensures that one will get identical results regardless of the order in which methods and indices are run in. When [ExtendedMethod](#) objects are passed, these are run after all [Method](#) objects have been run. This is because each [ExtendedMethod](#) object depends on the output of its base method. Furthermore, before an [ExtendedMethod](#) is called, the RNG state is restored to what it was after the base method had been called.

See Also

[generate_model](#) [simulate_from_model](#)

Examples

```
## Not run:
# suppose previously we had run the following:
sim <- new_simulation(name = "normal-example",
                    label = "Normal Mean Estimation",
                    dir = tempdir()) %>%
  generate_model(make_my_example_model, n = 20) %>%
  simulate_from_model(nsim = 50, index = 1:3)
# then we could add
sim <- run_method(sim, my_example_method)

## End(Not run)
```

run_method_single	<i>Run a single method on a single index of simulated data.</i>
-------------------	---

Description

This is an internal function. Users should call the wrapper function. [run_method](#). Here "single" refers to a single index-method pair.

Usage

```
run_method_single(method, model, draws_list)
```

Arguments

method	a Method object
model	a Model object
draws_list	the result of loading a Draws object with <code>more_info = TRUE</code> so that it includes RNG endstate.

save_simulation	<i>Save a simulation object</i>
-----------------	---------------------------------

Description

Saves an object of class [Simulation](#) to `sim@dir/files/sim@name.Rdata`. Note: while "files" is the default, the name of this directory is from `getOption("simulator.files")`, which is the value of `getOption("simulator.files")` when the model was created.

Usage

```
save_simulation(sim)
```

Arguments

sim an object of class [Simulation](#)

Details

This function overwrites any pre-existing file in that location without apology.

See Also

[new_simulation](#) [load_simulation](#)

simulate_from_model *Simulate from a model.*

Description

Given a reference to a [Model](#) object, this function calls the model's `simulate` function on its `params`. It repeats this `nsim` times. For example, when simulating regression with a fixed design, this function would generate `nsim` response vectors `y`.

Usage

```
simulate_from_model(object, nsim, index = 1, parallel = NULL)
```

Arguments

`object` an object of class [ModelRef](#) as returned by `link{generate_model}`. Or a list of such objects. If `object` is a [Simulation](#), then function is applied to the referenced models in that simulation and returns the same [Simulation](#) object but with references added to the new draws created.

`nsim` number of simulations to be conducted. If a scalar, then value repeated for each index. Otherwise can be a vector of length `length(index)`

`index` a vector of positive integer indices. Allows simulations to be carried out in chunks. Each chunk gets a separate RNG stream, meaning that the results will be identical whether we run these in parallel or sequentially.

`parallel` either `NULL` or a list containing `socket_names` and (optionally) `libraries` and `save_locally` (see [Details](#) for more information)

Details

This function creates objects of class [Draws](#) and saves each to file (at `dir/files/model_name/r<index>.Rdata`). Note: while "files" is the default, the name of this directory is from `getOption("simulator.files")`, which is the value of `getOption("simulator.files")` when the model was created.

If `parallel` is not `NULL`, then it must be a list containing `socket_names`, which can either be a positive integer specifying the number of copies to run on localhost or else a character vector of machine names (e.g., "mycluster-0-0"). The list `parallel` can also contain `libraries`, a character vector of R packages that will be needed on the slaves and `save_locally`, a logical that indicates whether the files generated should be saved on the slaves (i.e., locally) or on the master.

See Also

[load_draws](#) [generate_model](#) [run_method](#)

Examples

```
## Not run:
sim <- new_simulation(name = "normal-example",
                    label = "Normal Mean Estimation",
                    dir = tempdir()) %>%
  generate_model(make_my_example_model, n = 20) %>%
  simulate_from_model(nsim = 50, index = 1:3,
                    parallel = list(socket_names = 3))

## End(Not run)
```

```
simulate_from_model_single
    Simulate from a model.
```

Description

This is an internal function. Users should call the wrapper function [simulate_from_model](#).

Usage

```
simulate_from_model_single(model, nsim, index, seed)
```

Arguments

model	a Model object
nsim	number of simulations to be conducted.
index	a positive integer index.
seed	this is the 7 digit seed used by L'Ecuyer RNG

```
simulate_parallel    Simulate from a model in parallel.
```

Description

This is an internal function. Draws are done in chunks labeled by indices and of size determined by nsim. Users should call the wrapper function [simulate_from_model](#).

Usage

```
simulate_parallel(
  model_ref,
  nsim,
  index,
  seeds,
  socket_names,
  libraries,
  save_locally = TRUE
)
```

Arguments

model_ref	object of class ModelRef
nsim	number of simulations to be conducted on each chunk. Vector of same length as index
index	a vector of positive integer indices. Allows simulations to be carried out in chunks. Each chunk gets a separate RNG stream, meaning that the results will be identical whether we run these in parallel or sequentially.
seeds	a list of length(index) L'Ecuyer-CMRG seed vectors. Each should be from a separate stream. In particular, starting from the seed used to generate the model object, seeds[i] should be the result of calling nextRNGStream index[i] times.
socket_names	(quoting from makePSOCKcluster "either a character vector of host names on which to run the worker copies of R, or a positive integer (in which case that number of copies is run on localhost)."
libraries	character vector of R packages that will be needed on the slaves.
save_locally	if TRUE, then files will be saved on slaves. If FALSE, they will be saved on master.

Simulation-class	<i>An S4 class representing a simulation.</i>
------------------	---

Description

A simulation is a set of references to simulator objects that have been saved to file. The [DrawsRef](#), [OutputRef](#), and [EvalsRef](#) objects are organized by model into separate lists.

Details

When a reference ref is added to a simulation sim, ref@dir is changed so that the referenced file is located at file.path(sim@dir, ref@dir).

Slots

name a short name identifier. Must be an alphanumeric (but can also have - or _ within)
label a longer, human readable label that can have other characters such as spaces, hyphens, etc.
dir name of the directory where directory named "files" exists.
model_refs a list of [ModelRef](#) objects
draws_refs a list of lists of [DrawsRef](#) objects
output_refs a list of lists of [OutputRef](#) objects
evals_refs a list of lists of [EvalsRef](#) objects

subset_evals	<i>Reduce an Evals object to a subset of methods and/or metrics</i>
--------------	---

Description

If `method_names` is NULL, then subsetting is not done over methods. Likewise for `metric_names`.

Usage

```
subset_evals(evals, method_names = NULL, metric_names = NULL)
```

Arguments

<code>evals</code>	an object of class Evals or <code>listofEvals</code> .
<code>method_names</code>	a character vector of method names
<code>metric_names</code>	a character vector of metric names

subset_models	<i>Subset Models</i>
---------------	----------------------

Description

Given a list of [Model](#) objects, returns model names which meet conditions. Uses [subset](#)

Usage

```
subset_models(m, ...)
```

Arguments

<code>m</code>	list of Model objects
<code>...</code>	logical expression involving parameters of Models. For now, can only be parameters that are of length 1 and either of class numeric or character

subset_simulation	<i>Create a simulation that is a subset of a preexisting simulation object</i>
-------------------	--

Description

Given a simulation, creates a new simulation that is a subset of the preexisting simulation. Does not save this new one to file. To do so, first change the name (and, potentially, label) of the simulation and then use `save_simulation`. If you call `save_simulation` before changing the name, you will overwrite the preexisting simulation. Use `rename` and `relabel`.

Usage

```
subset_simulation(sim, ..., subset = NULL, index, methods)
```

Arguments

sim	a simulation object
...	logical conditions to specify a subset of models. Conditions can only involve params of model that have length 1 and are of class numeric or character.
subset	a vector of integers indexing the models or a vector of model names. To select models based on parameter values, use <code>...</code> . However, using <code>...</code> is slower than using <code>subset</code> .
index	a vector of positive integers specifying which draws' objects are desired. If missing, then all draws' evals are returned.
methods	character vector of method names of interest. If missing, then all methods' evals are returned

tabulate_eval	<i>Make a table of a metric for each pair of models and methods</i>
---------------	---

Description

Each row of the table corresponds to a different model and each column to a different method. The metric must be a scalar. The way in which standard error is shown (or not shown) is controlled by `se_format`.

Usage

```
tabulate_eval(
  object,
  metric_name,
  method_names = NULL,
  caption = NULL,
  center_aggregator = NULL,
```

```

spread_aggregator = NULL,
se_format = c("Paren", "PlusMinus", "None"),
output_type = "latex",
format_args = list(nsmall = 0, digits = NULL, scientific = FALSE),
na_string = "--",
bold = c("None", "Smallest", "Largest")
)

```

Arguments

object	an object of class Simulation , Evals , or <code>listofEvals</code> . Each evals object should just differ by <code>model_name</code> .
metric_name	the name of a metric to tabulate. Must be scalar valued.
method_names	character vector indicating methods to include in table. If <code>NULL</code> , then will include all methods found in object's evals.
caption	caption of plot. If <code>NULL</code> , then default caption used; if <code>FALSE</code> then no caption (and returns tabular without table).
center_aggregator	When <code>NULL</code> (which is default), the sample mean aggregator is used. User can write specialized aggregators (see definition of class Aggregator) as necessary, for example, when the evaluated metric is not scalar-valued.
spread_aggregator	When <code>NULL</code> (which is default), the standard error of the sample mean is used. User can write specialized aggregators (see definition of class Aggregator) as necessary, for example, when the evaluated metric is not scalar-valued. Set <code>spread_aggregator</code> to <code>NA</code> to hide error bars.
se_format	format of the standard error
output_type	see kable 's argument <code>format</code> for options. Default is "latex" but other options include "html" and "markdown"
format_args	arguments to pass to the function format
na_string	what to write in table in place of NA
bold	puts in bold the value that is smallest/largest for each model

Details

Uses `knitr`'s function `kable` to put table in various formats, including latex, html, markdown, etc.

Examples

```

## Not run:
# suppose previously we had run the following:
sim <- new_simulation(name = "normal-example",
                    label = "Normal Mean Estimation",
                    dir = tempdir()) %>%
  generate_model(make_my_example_model,
                n = list(10, 20, 30),
                vary_along = "n") %>%

```

```
simulate_from_model(nsim = 50, index = 1:3) %>%  
run_method(my_example_method) %>%  
evaluate(my_example_loss)  
# then we could plot this  
tabulate_eval(sim, "myloss")  
  
## End(Not run)
```

\$.Model-method *Get element of [Model](#)'s params list*

Description

Get element of [Model](#)'s params list

Usage

```
## S4 method for signature 'Model'  
x$name
```

Arguments

x	object of class Model
name	name of an element appearing in x@params

Index

- * **datasets**
 - my_example_loss, 31
 - my_example_method, 32
- +, ExtendedMethod, MethodExtension-method, 3
- +, Method, MethodExtension-method, 4
- +, list, MethodExtension-method, 4
- \$, Model-method, 53

- add, 5
- add, Simulation, DrawsRef-method (add), 5
- add, Simulation, EvalsRef-method (add), 5
- add, Simulation, list-method (add), 5
- add, Simulation, ModelRef-method (add), 5
- add, Simulation, OutputRef-method (add), 5
- add_bold, 6
- aggregate_evals, 6
- Aggregator, 32, 42, 52
- Aggregator-class, 6
- as.data.frame, 12
- as.data.frame.Evals, 7, 13, 24
- as.data.frame.listofEvals, 7
- as.data.frame.listofModels, 8
- as.data.frame.Model, 8

- cat, 9
- catsim, 9
- Component, 11, 17, 27–29, 34
- Component-class, 9
- create, 9

- describe, 10
- Draws, 12–14, 16, 23, 27, 37, 38, 44, 46, 47
- draws, 10, 23, 36
- Draws-class, 11
- DrawsRef, 35, 36, 45, 50
- DrawsRef-class, 12

- Evals, 7, 14–16, 24, 32, 39, 40, 50, 52
- evals, 12, 36

- Evals-class, 13
- EvalsRef, 24, 35, 36, 50
- EvalsRef-class, 14
- evaluate, 13, 14, 16, 24
- evaluate_internal, 15
- evaluate_single, 16
- ExtendedMethod, 4, 17, 28, 33, 34, 44, 45
- ExtendedMethod-class, 16

- format, 52

- generate_model, 15, 17, 23–25, 41, 45, 48
- get_contents, 18
- get_files_not_in_simulations, 19
- get_model_indices, 20
- get_relative_path, 20
- get_simulation_with_all_files, 21

- kable, 52

- load, 7, 8
- load, DrawsRef-method, 21
- load, EvalsRef-method, 22
- load, list-method, 22
- load, ModelRef-method, 22
- load, OutputRef-method, 23
- load_draws, 23, 24, 48
- load_evals, 24
- load_evals_from_ref (load_evals), 24
- load_model, 24, 25
- load_simulation, 25, 36, 47

- make_my_example_model, 26, 31, 32
- makePSOCKcluster, 49
- memory_as_string, 27
- Method, 4, 13, 14, 16, 17, 24, 32, 33, 37, 38, 45, 46
- Method-class, 27
- MethodExtension, 4
- MethodExtension-class, 28
- Metric, 13–16, 31, 34

- Metric-class, 28
- Model, 8, 12–18, 24–26, 28, 30, 35, 37, 38, 44, 46, 47, 50, 53
- model, 10, 20, 25, 29, 36
- Model-class, 29
- model_names, 31
- ModelRef, 17, 35, 36, 47, 49, 50
- ModelRef-class, 30
- models_as_data.frame, 30
- my_example_loss, 26, 31, 31, 32
- my_example_method, 26, 32

- new_aggregator, 32
- new_extended_method, 33
- new_method, 33
- new_method_extension, 34
- new_metric, 34
- new_model, 18, 35
- new_simulation, 25, 26, 35, 47
- nextRNGStream, 49

- Output, 14–16, 28, 38, 44, 45
- output, 36, 36
- Output-class, 37
- OutputRef, 14, 35, 36, 50
- OutputRef-class, 38

- plot_eval, 38
- plot_eval_by, 39, 41
- plot_evals, 39, 39

- recycle, 43
- relabel, 43, 44, 51
- rename, 43, 44, 51
- run_extendedmethod_single, 44
- run_method, 14, 15, 18, 24, 44, 45, 46, 48
- run_method_single, 46

- save_simulation, 26, 36, 43, 44, 46, 51
- simulate_from_model, 11, 15, 18, 23, 45, 47, 48
- simulate_from_model_single, 48
- simulate_parallel, 48
- Simulation, 14, 17, 25, 31, 35, 39–41, 43–47, 52
- Simulation-class, 49
- subset, 50
- subset_evals, 50
- subset_models, 50

- subset_simulation, 51
- system.time, 14
- tabulate_eval, 39, 51