

# Package ‘workflowsets’

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**Title** Create a Collection of 'tidymodels' Workflows

**Version** 0.1.0

**Description** A workflow is a combination of a model and preprocessors (e.g, a formula, recipe, etc.) (Kuhn and Silge (2021) <<https://www.tmwr.org/>>). In order to try different combinations of these, an object can be created that contains many workflows. There are functions to create workflows en masse as well as training them and visualizing the results.

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**URL** <https://github.com/tidymodels/workflowsets>,  
<https://workflowsets.tidymodels.org>

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as_workflow_set	<i>Save results from tuning or resampling functions as a workflow set</i>
-----------------	---

---

### Description

If results have been generated directly from functions like `tune::tune_grid()`, they can be combined into a workflow set using this function.

### Usage

```
as_workflow_set(...)
```

### Arguments

... One or more named objects. Names should be unique and the objects should have at least one of the following classes: `iteration_results`, `tune_results`, `resample_results`, or `tune_race`. Each element should also contain the original workflow (accomplished using the `save_workflow` option in the control function).

### Value

A workflow set. Note that the `option` column will not reflect the options that were used to create each object.

---

autoplot.workflow\_set *Plot the results of a workflow set*

---

### Description

This autoplot() method can performance metrics that have been ranked using a metric. It can also run autoplot() on the individual results (per wflow\_id).

### Usage

```
## S3 method for class 'workflow_set'
autoplot(
  object,
  rank_metric = NULL,
  metric = NULL,
  id = "workflow_set",
  select_best = FALSE,
  std_errs = qnorm(0.95),
  ...
)
```

### Arguments

object	A workflow_set whose elements have results.
rank_metric	A character string for which metric should be used to rank the results. If none is given, the first metric in the metric set is used (after filtering by the metric option).
metric	A character vector for which metrics (apart from rank_metric) to be included in the visualization.
id	A character string for what to plot. If a value of "workflow_set" is used, the results of each model (and sub-model) are ordered and plotted. Alternatively, a value of the workflow set's wflow_id can be given and the autoplot() method is executed on that workflow's results.
select_best	A logical; should the results only contain the numerically best submodel per workflow?
std_errs	The number of standard errors to plot (if the standard error exists).
...	Other options to pass to autoplot().

### Details

This function is intended to produce a default plot to visualize helpful information across all possible applications of a workflow set. A more appropriate plot for your specific analysis can be created by calling [rank\\_results\(\)](#) and using standard ggplot2 code for plotting.

The x-axis is the workflow rank in the set (a value of one being the best) versus the performance metric(s) on the y-axis. With multiple metrics, there will be facets for each metric.

If multiple resamples are used, confidence bounds are shown for each result (90% confidence, by default).

### Value

A ggplot object.

### Examples

```
autoplot(two_class_res)
autoplot(two_class_res, select_best = TRUE)
autoplot(two_class_res, id = "yj_trans_cart", metric = "roc_auc")
```

---

```
collect_metrics.workflow_set
```

*Obtain and format results produced by tuning functions for workflow sets*

---

### Description

Return a tibble of performance metrics for all models or submodels.

### Usage

```
## S3 method for class 'workflow_set'
collect_metrics(x, summarize = TRUE, ...)

## S3 method for class 'workflow_set'
collect_predictions(
  x,
  summarize = TRUE,
  parameters = NULL,
  select_best = FALSE,
  metric = NULL,
  ...
)
```

### Arguments

x	A workflow_set object where all workflows have been evaluated.
summarize	A logical for whether the performance estimates should be summarized via the mean (over resamples) or the raw performance values (per resample) should be returned along with the resampling identifiers. When collecting predictions, these are averaged if multiple assessment sets contain the same row.
...	Not currently used.
parameters	An optional tibble of tuning parameter values that can be used to filter the predicted values before processing. This tibble should only have columns for each tuning parameter identifier (e.g. "my_param" if tune("my_param") was used).

select_best	A single logical for whether the numerically best results are retained. If TRUE, the parameters argument is ignored.
metric	A character string for the metric that is used for select_best.

### Details

When applied to a workflow set, the metrics and predictions that are returned do not contain the actual tuning parameter columns and values (unlike when these collect functions are run on other objects). The reason is that workflow sets can contain different types of models or models with different tuning parameters.

If the columns are needed, there are two options. First, the `.config` column can be used to merge the tuning parameter columns into an appropriate object. Alternatively, the `map()` function can be used to get the metrics from the original objects (see the example below).

### Value

A tibble.

### See Also

[tune::collect\\_metrics\(\)](#), [rank\\_results\(\)](#)

### Examples

```
library(dplyr)
library(purrr)
library(tidyr)

# -----

collect_metrics(two_class_res)

# Alternatively, if the tuning parameter values are needed:
two_class_res %>%
  dplyr::filter(grepl("cart", wflow_id)) %>%
  mutate(metrics = map(result, collect_metrics)) %>%
  dplyr::select(wflow_id, metrics) %>%
  tidyr::unnest(cols = metrics)

collect_metrics(two_class_res, summarize = FALSE)
```

---

comment_add	<i>Add annotations and comments for workflows</i>
-------------	---

---

### Description

comment\_add() can be used to log important information about the workflow or its results as you work. Comments can be appended or removed.

### Usage

```
comment_add(x, id, ..., append = TRUE, collapse = "\n")
```

```
comment_get(x, id)
```

```
comment_reset(x, id)
```

```
comment_print(x, id = NULL, ...)
```

### Arguments

x	A workflow set
id	A single character string for a value in the wflow_id column. For comment_print(), id can be a vector or NULL (and this indicates that all comments are printed).
...	One or more character strings.
append	A logical value to determine if the new comment should be added to the existing values.
collapse	A character string that separates the comments.

### Value

comment\_add() and comment\_reset() return an updated workflow set. comment\_get() returns a character string. comment\_print() returns NULL invisibly.

### Examples

```
two_class_set %>% comment_get("none_cart")

new_set <-
  two_class_set %>%
  comment_add("none_cart", "What does 'cart' stand for\u2753") %>%
  comment_add("none_cart", "Classification And Regression Trees.")

comment_print(new_set)

new_set %>% comment_get("none_cart")

new_set %>% comment_reset("none_cart") %>% comment_get("none_cart")
```

---

`extract_workflow_set_result`*Extract elements of workflow sets*

---

## Description

These functions extract various elements from a workflow set object. If they do not exist yet, an error is thrown.

- `extract_preprocessor()` returns the formula, recipe, or variable expressions used for preprocessing.
- `extract_spec_parsnip()` returns the parsnip model specification.
- `extract_fit_parsnip()` returns the parsnip model fit object.
- `extract_fit_engine()` returns the engine specific fit embedded within a parsnip model fit. For example, when using `parsnip::linear_reg()` with the "lm" engine, this returns the underlying lm object.
- `extract_mold()` returns the preprocessed "mold" object returned from `hardhat::mold()`. It contains information about the preprocessing, including either the prepped recipe, the formula terms object, or variable selectors.
- `extract_recipe()` returns the recipe. The `estimated` argument specifies whether the fitted or original recipe is returned.
- `extract_workflow_set_result()` returns the results of `workflow_map()` for a particular workflow.
- `extract_workflow()` returns the workflow object. The workflow will not have been estimated.

## Usage

```
extract_workflow_set_result(x, id, ...)  
  
## S3 method for class 'workflow_set'  
extract_workflow(x, id, ...)  
  
## S3 method for class 'workflow_set'  
extract_spec_parsnip(x, id, ...)  
  
## S3 method for class 'workflow_set'  
extract_recipe(x, id, ..., estimated = TRUE)  
  
## S3 method for class 'workflow_set'  
extract_fit_parsnip(x, id, ...)  
  
## S3 method for class 'workflow_set'  
extract_fit_engine(x, id, ...)
```

```
## S3 method for class 'workflow_set'
extract_mold(x, id, ...)
```

```
## S3 method for class 'workflow_set'
extract_preprocessor(x, id, ...)
```

### Arguments

x	A workflow set.
id	A single character string for a workflow ID.
...	Other options (not currently used).
estimated	A logical for whether the original (unfit) recipe or the fitted recipe should be returned.

### Details

These functions supersede the `pull_*()` functions (e.g., `extract_workflow_set_result()`).

### Value

The extracted value from the object, x, as described in the description section.

### Examples

```
library(tune)

extract_workflow_set_result(two_class_res, "none_cart")

extract_workflow(two_class_res, "none_cart")
```

---

```
leave_var_out_formulas
```

*Create formulas without each predictor*

---

### Description

From an initial model formula, create a list of formulas that exclude each predictor.

### Usage

```
leave_var_out_formulas(formula, data, full_model = TRUE, ...)
```

### Arguments

formula	A model formula that contains at least two predictors.
data	A data frame.
full_model	A logical; should the list include the original formula?
...	Options to pass to <code>stats::model.frame()</code>



**Details**

The new formulas obey the hierarchy rule so that interactions without main effects are not included (unless the original formula contains such terms).

Factor predictors are left as-is (i.e., no indicator variables are created).

**Value**

A named list of formulas

**See Also**

[workflow\\_set\(\)](#)

**Examples**

```
data(penguins, package = "modeldata")

leave_var_out_formulas(
  bill_length_mm ~ .,
  data = penguins
)

leave_var_out_formulas(
  bill_length_mm ~ (island + sex)^2 + flipper_length_mm,
  data = penguins
)

leave_var_out_formulas(
  bill_length_mm ~ (island + sex)^2 + flipper_length_mm +
    I(flipper_length_mm^2),
  data = penguins
)
```

---

option\_add

*Add and edit options saved in a workflow set*

---

**Description**

These functions are helpful for manipulating the information in the option column.

**Usage**

```
option_add(x, ..., id = NULL, strict = FALSE)
```

```
option_remove(x, ...)
```

```
option_add_parameters(x, id = NULL, strict = FALSE)
```

**Arguments**

x	A workflow set
...	A list of named options. For <code>option_remove()</code> this can be a series of unquoted option names.
id	A character string of one or more values from the <code>wflow_id</code> column that indicates which options to update. By default, all workflows are updated.
strict	A logical; show execution stop if existing options are being replaced?

**Details**

`option_add()` is used to update all of the options in a workflow set.

`option_remove()` will eliminate specific options across rows.

`option_add_parameters()` adds a parameter object to the `option` column (if parameters are being tuned).

Note that executing a function on the workflow set, such as `tune_grid()`, will add any options given to that function to the `option` column.

**Value**

An updated workflow set.

**Examples**

```
two_class_set %>%
  option_add(a = 1)

two_class_set %>%
  option_add(a = 1) %>%
  option_add(b = 2, id = "none_cart")

library(tune)
two_class_set %>%
  option_add_parameters()
```

---

option_list	<i>Make a classed list of options</i>
-------------	---------------------------------------

---

**Description**

This function returns a named list with an extra class of "workflow\_set\_options" that has corresponding formatting methods for printing inside of tibbles.

**Usage**

```
option_list(...)
```

**Arguments**

... A set of named options (or nothing)

**Value**

A classed list.

**Examples**

```
option_list(a = 1, b = 2)
option_list()
```

---

pull\_workflow\_set\_result

*Extract elements from a workflow set*

---

**Description**

**[Soft-deprecated]**

**Usage**

```
pull_workflow_set_result(x, id)
```

```
pull_workflow(x, id)
```

**Arguments**

x A workflow set.

id A single character string for a workflow ID.

**Details**

pull\_workflow\_set\_result() retrieves the results of [workflow\\_map\(\)](#) for a particular workflow while pull\_workflow() extracts the unfitted workflow from the info column.

The [extract\\_workflow\\_set\\_result\(\)](#) and [extract\\_workflow\(\)](#) functions should be used instead of these functions.

**Value**

pull\_workflow\_set\_result() produces a tune\_result or resample\_results object. pull\_workflow() returns an unfit workflow object.

**Examples**

```
library(tune)

pull_workflow_set_result(two_class_res, "none_cart")

pull_workflow(two_class_res, "none_cart")
```

---

rank_results	<i>Rank the results by a metric</i>
--------------	-------------------------------------

---

**Description**

This function sorts the results by a specific performance metric.

**Usage**

```
rank_results(x, rank_metric = NULL, select_best = FALSE)
```

**Arguments**

x	A workflow set that has all results.
rank_metric	A character string for a metric.
select_best	A logical; should the results only contain the numerically best submodel per workflow.

**Details**

If some models have the exact same performance, `rank(value, ties.method = "random")` is used (with a reproducible seed) so that all ranks are integers.

No columns are returned for the tuning parameters since they are likely to be different (or not exist) for some models. The `wflow_id` and `.config` columns can be used to determine the corresponding parameter values.

**Value**

A tibble with columns: `wflow_id`, `.config`, `.metric`, `mean`, `std_err`, `n`, `preprocessor`, `model`, and `rank`.

**Examples**

```
rank_results(chi_features_res)
rank_results(chi_features_res, select_best = TRUE)
rank_results(chi_features_res, rank_metric = "rsq")
```

---

two_class_set	<i>Example Data Sets</i>
---------------	--------------------------

---

### Description

Example Data Sets

### Details

Example workflow sets and associated model fits.

two\_class\_set and two\_class\_res were generated using the data in the package file example-data/two-class-set.R  
chi\_features\_set and chi\_features\_res were generated using the data in the package file example-data/chi-features-res.R. It is meant to approximate the sequence of models built in Section 1.3 of Kuhn and Johnson (2019).

### Value

Workflow sets.

### References

Max Kuhn and Kjell Johnson (2019) *Feature Engineering and Selection*, <https://bookdown.org/max/FES/a-more-complex-example.html>

### Examples

```
data(two_class_set)
two_class_set
```

---

workflow_map	<i>Process a series of workflows</i>
--------------	--------------------------------------

---

### Description

workflow\_map() will execute the same function across the workflows in the set. The various tune\_\*() functions can be used as well as `tune::fit_resamples()`.

### Usage

```
workflow_map(  
  object,  
  fn = "tune_grid",  
  verbose = FALSE,  
  seed = sample.int(10^4, 1),  
  ...  
)
```

**Arguments**

object	A workflow set.
fn	The function to run. Acceptable values are: <code>tune::tune_grid()</code> , <code>tune::tune_bayes()</code> , <code>tune::fit_resamples()</code> , <code>finetune::tune_race_anova()</code> , <code>finetune::tune_race_win_loss()</code> , or <code>finetune::tune_sim_anneal()</code> .
verbose	A logical for logging progress.
seed	A single integer that is set prior to each function execution.
...	Options to pass to the modeling function. See details below.

**Details**

When passing options, anything passed in the ... will be combined with any values in the option column. The values in ... will override that column's values and the new options are added to the options column.

Any failures in execution result in the corresponding row of results to contain a try-error object.

In cases where a model has no tuning parameters is mapped to one of the tuning functions, `tune::fit_resamples()` will be used instead and a warning is issued if `verbose = TRUE`.

If a workflow required packages that are not installed, a message is printed and `workflow_map()` continues with the next workflow (if any).

**Value**

An updated workflow set. The option column will be updated with any options for the tune package functions given to `workflow_map()`. Also, the results will be added to the result column. If the computations for a workflow fail, a try-catch object will be saved in place of the results (without stopping execution).

**See Also**

`workflow_set()`, `as_workflow_set()`, `extract_workflow_set_result()`

**Examples**

```
# An example of processed results
chi_features_res

# Code examples at
if (interactive()) {
  system.file("example-data", package = "workflowsets")
}
```

---

workflow_set	<i>Generate a set of workflow objects from preprocessing and model objects</i>
--------------	--

---

### Description

Generate a set of workflow objects from preprocessing and model objects

### Usage

```
workflow_set(preproc, models, cross = TRUE)
```

### Arguments

preproc	A list (preferably named) with preprocessing objects: formulas, recipes, or <a href="#">workflows::workflow_variables()</a> .
models	A list (preferably named) of parsnip model specifications.
cross	A logical: should all combinations of the preprocessors and models be used to create the workflows? If FALSE, the length of preproc and models should be equal.

### Details

The preprocessors that can be combined with the model objects can be one or more of:

- A traditional R formula.
- A recipe definition (un-prepared) via [recipes::recipe\(\)](#).
- A selectors object created by [workflows::workflow\\_variables\(\)](#).

Since preproc is a named list column, any combination of these can be used in that argument (i.e., preproc can be mixed types).

### Value

A tibble with extra class 'workflow\_set'. A new set includes four columns (but others can be added):

- `wflow_id` contains character strings for the preprocessor/workflow combination. These can be changed but must be unique.
- `info` is a list column with tibbles containing more specific information, including any comments added using [comment\\_add\(\)](#). This tibble also contains the workflow object (which can be easily retrieved using [extract\\_workflow\(\)](#)).
- `option` is a list column that will include a list of optional arguments passed to the functions from the tune package. They can be added manually via [option\\_add\(\)](#) or automatically when options are passed to [workflow\\_map\(\)](#).
- `result` is a list column that will contain any objects produced when [workflow\\_map\(\)](#) is used.

**See Also**

[workflow\\_map\(\)](#), [comment\\_add\(\)](#), [option\\_add\(\)](#), [as\\_workflow\\_set\(\)](#)

**Examples**

```

library(workflows)
library(modeldata)
library(recipes)
library(parsnip)
library(dplyr)
library(rsample)
library(tune)
library(yardstick)

# -----

data(cells)
cells <- cells %>% dplyr::select(-case)

set.seed(1)
val_set <- validation_split(cells)

# -----

basic_recipe <-
  recipe(class ~ ., data = cells) %>%
  step_YeoJohnson(all_predictors()) %>%
  step_normalize(all_predictors())

pca_recipe <-
  basic_recipe %>%
  step_pca(all_predictors(), num_comp = tune())

ss_recipe <-
  basic_recipe %>%
  step_spatialsign(all_predictors())

# -----

knn_mod <-
  nearest_neighbor(neighbors = tune(), weight_func = tune()) %>%
  set_engine("kknn") %>%
  set_mode("classification")

lr_mod <-
  logistic_reg() %>%
  set_engine("glm")

# -----

preproc <- list(none = basic_recipe, pca = pca_recipe, sp_sign = ss_recipe)

```



```
models <- list(knn = knn_mod, logistic = lr_mod)

cell_set <- workflow_set(preproc, models, cross = TRUE)
cell_set

# -----
# Using variables and formulas

# Select predictors by their names
channels <- paste0("ch_", 1:4)
preproc <- purrr::map(channels, ~ workflow_variables(class, c(contains(!.x))))
names(preproc) <- channels
preproc$everything <- class ~ .
preproc

cell_set_by_group <- workflow_set(preproc, models["logistic"])
cell_set_by_group
```

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