

# Package ‘workboots’

October 12, 2022

**Title** Generate Bootstrap Prediction Intervals from a 'tidymodels' Workflow

**Version** 0.2.0

**Description** Provides functions for generating bootstrap prediction intervals from a 'tidymodels' workflow. 'tidymodels' <<https://www.tidymodels.org/>> is a collection of packages for modeling and machine learning using 'tidyverse' <<https://www.tidyverse.org/>> principles. This package is not affiliated with or maintained by 'RStudio' or the 'tidymodels' maintainers.

**License** MIT + file LICENSE

**URL** <https://github.com/markjrieke/workboots>,  
<https://markjrieke.github.io/workboots/>

**BugReports** <https://github.com/markjrieke/workboots/issues>

**Imports** assertthat, dplyr, generics, lifecycle, Metrics, purrr, rlang, rsample, stats, tibble, tidyr, vip, workflows

**Encoding** UTF-8

**RoxygenNote** 7.1.2

**Suggests** forcats, ggplot2, knitr, readr, recipes, rmarkdown, scales, testthat (>= 3.0.0), tidymodels, tune, xgboost

**VignetteBuilder** knitr

**Config/testthat/edition** 3

**Depends** R (>= 2.10)

**NeedsCompilation** no

**Author** Mark Rieke [aut, cre]

**Maintainer** Mark Rieke <markjrieke@gmail.com>

**Repository** CRAN

**Date/Publication** 2022-05-16 06:40:02 UTC

## R topics documented:

predict_boots . . . . .	2
summarise_importance . . . . .	3
summarise_predictions . . . . .	4
vi_boots . . . . .	5

<b>Index</b>	<b>7</b>
--------------	----------

---

predict_boots	<i>Fit and predict from a workflow using many bootstrap resamples.</i>
---------------	--

---

### Description

Generate a prediction interval from arbitrary model types using bootstrap resampling. `predict_boots()` generates `n` bootstrap resamples, fits a model to each resample (creating `n` models), then creates `n` predictions for each observation in `new_data`.

### Usage

```
predict_boots(
  workflow,
  n = 2000,
  training_data,
  new_data,
  interval = c("prediction", "confidence"),
  verbose = FALSE,
  ...
)
```

### Arguments

<code>workflow</code>	An un-fitted workflow object.
<code>n</code>	An integer for the number of bootstrap resampled models that will be created.
<code>training_data</code>	A tibble or dataframe of data to be resampled and used for training.
<code>new_data</code>	A tibble or dataframe used to make predictions.
<code>interval</code>	One of prediction, confidence. Specifies the interval type to be generated.
<code>verbose</code>	A logical. Defaults to FALSE. If set to TRUE, prints progress of training to console.
<code>...</code>	Additional params passed to <code>rsample::bootstraps()</code> .

### Details

Since `predict_boots()` fits a new model to each resample, the argument `workflow` must not yet be fit. Any tuned hyperparameters must be finalized prior to calling `predict_boots()`.

**Value**

A tibble with a column indicating the row index of each observation in `new_data` and a nested list of the model predictions for each observation.

**Examples**

```
## Not run:
library(tidymodels)

# setup a workflow without fitting
wf <-
  workflow() %>%
  add_recipe(recipe(qsec ~ wt, data = mtcars)) %>%
  add_model(linear_reg())

# fit and predict 2000 bootstrap resampled models to mtcars
set.seed(123)
wf %>%
  predict_boots(n = 2000, training_data = mtcars, new_data = mtcars)

## End(Not run)
```

---

`summarise_importance` *Append a tibble of variable importances returned by `vi_boots()` with upper and lower bounds.*

---

**Description**

Append a tibble of variable importances returned by `vi_boots()` with upper and lower bounds.

**Usage**

```
summarise_importance(.data, interval_width = 0.95, conf = NULL)
summarize_importance(.data, interval_width = 0.95, conf = NULL)
```

**Arguments**

`.data` a tibble of variable importances returned by `vi_boots()`.  
`interval_width` a value between (0, 1) specifying the interval range.  
`conf` deprecated - please use `interval_width` instead.

**Details**

Generates a summary of variable importances with an upper and lower interval range. Uses the `vi()` function from the `{vip}` package to compute variable importances (not all model types are supported by `vip::vi()`; please refer to `{vip}` package documentation for supported model types). Presently, the `quantile()` function from the `{stats}` package is used to determine the lower, 50th percentile, and upper interval ranges.

## Examples

```
## Not run:
library(tidymodels)

# setup a workflow without fitting
wf <-
  workflow() %>%
  add_recipe(recipe(qsec ~ wt, data = mtcars)) %>%
  add_model(linear_reg())

# evaluate variable importance from 2000 models fit to mtcars
set.seed(123)
importances <-
  wf %>%
  vi_boots(n = 2000, training_data = mtcars, new_data = mtcars)

# append with lower and upper bound importance summary columns
importances %>%
  summarise_importance(interval_width = 0.95)

## End(Not run)
```

---

`summarise_predictions` *Append a tibble of predictions returned by `predict_boots()` with upper and lower bounds.*

---

## Description

Append a tibble of predictions returned by `predict_boots()` with upper and lower bounds.

## Usage

```
summarise_predictions(.data, interval_width = 0.95, conf = NULL)
```

```
summarize_predictions(.data, interval_width = 0.95, conf = NULL)
```

## Arguments

<code>.data</code>	a tibble of predictions returned by <code>predict_boots()</code> .
<code>interval_width</code>	a value between (0, 1) specifying the interval range.
<code>conf</code>	deprecated - please use <code>interval_width</code> instead.

## Details

Generates a summary of predictions with a upper and lower interval range. Presently, the `quantile()` function from the `{stats}` package is used to determine the lower, 50th percentile, and upper interval ranges.

**Value**

Appends the tibble of predictions returned by `predict_boots()` with three new columns: `.pred_lower`, `.pred`, and `.pred_upper`.

**Examples**

```
## Not run:
library(tidymodels)

# setup a workflow without fitting
wf <-
  workflow() %>%
  add_recipe(recipe(qsec ~ wt, data = mtcars)) %>%
  add_model(linear_reg())

# fit and predict 2000 bootstrap resampled models to mtcars
set.seed(123)
preds <-
  wf %>%
  predict_boots(n = 2000, training_data = mtcars, new_data = mtcars)

# append with prediction interval summary columns
preds %>%
  summarise_predictions(conf = 0.95)

## End(Not run)
```

---

vi_boots	<i>Fit and estimate variable importance from a workflow using many bootstrap resamples.</i>
----------	---

---

**Description**

Generate variable importances from a tidymodel workflow using bootstrap resampling. `vi_boots()` generates `n` bootstrap resamples, fits a model to each (creating `n` models), then creates `n` estimates of variable importance for each variable in the model.

**Usage**

```
vi_boots(workflow, n = 2000, training_data, verbose = FALSE, ...)
```

**Arguments**

<code>workflow</code>	An un-fitted workflow object.
<code>n</code>	An integer for the number of bootstrap resampled models that will be created.
<code>training_data</code>	A tibble or dataframe of data to be resampled and used for training.
<code>verbose</code>	A logical. Defaults to FALSE. If set to TRUE, prints progress of training to console.
<code>...</code>	Additional params passed to <code>rsample::bootstraps()</code> .

## Details

Since `vi_boots()` fits a new model to each resample, the argument `workflow` must not yet be fit. Any tuned hyperparameters must be finalized prior to calling `vi_boots()`.

## Value

A tibble with a column indicating each variable in the model and a nested list of variable importances for each variable. The shape of the list may vary by model type. For example, linear models return two nested columns: the absolute value of each variable's importance and the sign (POS/NEG), whereas tree-based models return a single nested column of variable importance. Similarly, the number of nested rows may vary by model type as some models may not utilize every possible predictor.

## Examples

```
## Not run:
library(tidymodels)

# setup a workflow without fitting
wf <-
  workflow() %>%
  add_recipe(recipe(qsec ~ wt, data = mtcars)) %>%
  add_model(linear_reg())

# fit and estimate variable importance from 125 bootstrap resampled models
set.seed(123)
wf %>%
  vi_boots(n = 2000, training_data = mtcars)

## End(Not run)
```

# Index

`predict_boots`, [2](#)

`summarise_importance`, [3](#)

`summarise_predictions`, [4](#)

`summarize_importance`

    (`summarise_importance`), [3](#)

`summarize_predictions`

    (`summarise_predictions`), [4](#)

`vi_boots`, [5](#)