Package 'workboots'

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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.

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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

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 $predict_boots$

Fit and predict from a workflow using many bootstrap resamples.

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

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

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().

summarise_importance

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.

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Examples

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)
```

upper and lower bounds.

Arguments

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 up- per 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

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

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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(gsec ~ 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

Fit and estimate variable importance from a workflow using many bootstrap resamples.

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

workflow

An un-fitted workflow object. An integer for the number of bootstrap resampled models that will be created. training_data A tibble or dataframe of data to be resampled and used for training. verbose A logical. Defaults to FALSE. If set to TRUE, prints progress of training to con-

Additional params passed to rsample::bootstraps().

vi_boots

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)
```

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