Package 'randomForestVIP'

July 19, 2023

Type Package

Title Tune Random Forests Based on Variable Importance & Plot Results **Version** 0.1.3

Description Functions for assessing variable relations and associations prior to modeling with a Random Forest algorithm (although these are

relevant for any predictive model). Metrics such as partial correlations and variance inflation factors are tabulated as well as plotted for the user. A function is available for tuning the main Random Forest hyper-parameter based on model performance and variable importance metrics. This grid-search technique provides tables and plots showing the effect of the main hyper-parameter on each of the assessment metrics. It also returns each of the evaluated models to the user. The package also provides superior variable importance plots for individual models. All of the plots are developed so that the user has the ability to edit and improve further upon the plots. Derivations and methodology are described in Bladen (2022) <https://digitalcommons.usu.edu/etd/8587/>.

License GPL-3

URL https://github.com/KelvynBladen/randomForestVIP

Depends R (>= 4.0.0)

Imports car, dplyr, ggplot2, gridExtra, minerva, randomForest, stats, tidyr

Suggests EZtune, e1071, knitr, MASS, rmarkdown, rpart, testthat (>= 3.0.0)

VignetteBuilder knitr

Config/testthat/edition 3

Encoding UTF-8

LazyData true

RoxygenNote 7.2.3

NeedsCompilation no

Author Kelvyn Bladen [aut, cre],

D. Richard Cutler [aut]

boston

Maintainer Kelvyn Bladen <kelvyn.bladen@usu.edu> Repository CRAN Date/Publication 2023-07-19 11:20:02 UTC

R topics documented:

boston																							2
ggvip															•								3
lichen															•								4
mtry_compare	e .	•		•			•			•			•	•	•			•	•	•	•		6
partial_cor		•		•			•			•			•	•	•			•	•	•	•		7
robust_vifs .															•								7
																							- 9

Index

boston

Housing Values in Suburbs of Boston

Description

The Boston data frame has 506 rows and 14 columns.

Usage

boston

Format

This data frame contains the following columns:

crim per capita crime rate by town.

zn proportion of residential land zoned for lots over 25,000 sq.ft.

indus proportion of non-retail business acres per town.

chas Charles River dummy variable (= 1 if tract bounds river; 0 otherwise).

nox nitrogen oxides concentration (parts per 10 million).

rm average number of rooms per dwelling.

age proportion of owner-occupied units built prior to 1940.

dis weighted mean of distances to five Boston employment centres.

rad index of accessibility to radial highways.

tax full-value property-tax rate per \$10,000.

ptratio pupil-teacher ratio by town.

black $1000(Bk - 0.63)^2$ where Bk is the proportion of blacks by town.

lstat lower status of the population (percent).

medv median value of owner-occupied homes in \$1000s.

ggvip

Source

https://www.stats.ox.ac.uk/pub/MASS4/

ggvip

Variable Importance GGPlot

Description

A ggplot of variable importance as measured by a Random Forest.

Usage

ggvip(x, scale = FALSE, sqrt = TRUE, type = "both", num_var)

Arguments

х	An object of class randomForest.
scale	For permutation based measures such as MSE or Accuracy, should the measures be divided by their "standard errors"? Default is False.
sqrt	Boolean value indicating whether importance metrics should be adjusted via a square root transformation. Default is True.
type	either 1 or 2, specifying the type of importance measure (1=mean decrease in accuracy or node impurity or mean decrease in gini). Default is "both".
num_var	Optional argument for reducing the number of variables to the top 'num_var'. Must be an integer between 1 and the total number of predictor variables in the model.

Value

A ggplot dotchart showing the importance of the variables that were plotted.

Examples

```
rf <- randomForest::randomForest(factor(Species) ~ .,
    importance = TRUE,
    data = iris
)
ggvip(rf, scale = FALSE, sqrt = TRUE, type = "both")
```

lichen

Description

Data were collected between 1993 and 1999 as part of the Lichen Air Quality surveys on public lands in Oregon and southern Washington. Observations were obtained from 1-acre (0.4 ha) plots at Current Vegetation Survey (CVS) sites. Indicator variables denote the presences and absences of 7 lichen species. Data for each sampled plot include the topographic variables elevation, aspect, and slope; bioclimatic predictors including maximum, minimum, daily, and average temperatures, relative humidity precipitation, evapotranspiration, and vapor pressure; and vegetation variables including the average age of the dominant conifer and percent conifer cover. The data in lichenTest were collected from half-acre plots at CVS sites in the same geographical region and contains many of the same variables, including presences and absences for the 7 lichen species. As such, it is a good test dataset for predictive methods applied to the Lichen Air Quality data.

Usage

lichen

Format

A data frame with 840 observations and 40 variables. One variable is a location identifier, 7 (coded as 0 and 1) identify the presence or absence of a type of lichen species, and 32 are characteristics of the survey site where the data were collected.

There were 12 monthly values in the original data for each of the bioclimatic predictors. Principal components analyses suggested that for each of these predictors 2 principal components explained the vast majority (95.0%-99.5%) of the total variability. Based on these analyses, indices were created for each set of bioclimatic predictors. The variables with the suffix Ave in the variable name are the average of 12 monthly variables. The variables with the suffix Diff are contrasts between the sum of the April-September monthly values and the sum of the October-December and January-March monthly values, divided by 12. Roughly speaking, these are summer-to-winter contrasts.

The variables are summarized as follows:

LobaOreg Lobaria oregana (Absent = 0, Present = 1)

EvapoTransAve Average monthly potential evapotranspiration in mm

EvapoTransDiff Summer-to-winter difference in monthly potential evapotranspiration in mm

MoistIndexAve Average monthly moisture index in cm

MoistIndexDiff Summer-to-winter difference in monthly monthly moisture index in cm

PrecipAve Average monthly precipitation in cm

PrecipDiff Summer-to-winter difference in monthly precipitation in cm

RelHumidAve Average monthly relative humidity in percent

RelHumidDiff Summer-to-winter difference in monthly relative humidity in percent

PotGlobRadAve Average monthly potential global radiation in kJ

lichen

PotGlobRadDiff Summer-to-winter difference in monthly potential global radiation in kJ AveTempAve Average monthly average temperature in degrees Celsius AveTempDiff Summer-to-winter difference in monthly average temperature in degrees Celsius **MaxTempAve** Average monthly maximum temperature in degrees Celsius MaxTempDiff Summer-to-winter difference in monthly maximum temperature in degrees Celsius MinTempAve Average monthly minimum temperature in degrees Celsius MinTempDiff Summer-to-winter difference in monthly minimum temperature in degrees Celsius DayTempAve Mean average daytime temperature in degrees Celsius DayTempDiff Summer-to-winter difference in average daytime temperature in degrees Celsius AmbVapPressAve Average monthly average ambient vapor pressure in Pa AmbVapPressDiff Summer-to-winter difference in monthly average ambient vapor pressure in Pa SatVapPressAve Average monthly average saturated vapor pressure in Pa **SatVapPressDiff** Summer-to-winter difference in monthly average saturated vapor pressure in Pa Aspect Aspect in degrees TransAspect Transformed Aspect: TransAspect=(1-cos(Aspect))/2 Elevation Elevation in meters Slope Percent slope **ReserveStatus** Reserve Status (Reserve, Matrix) StandAgeClass Stand Age Class (< 80 years, 80+ years) **ACONIF** Average age of the dominant conifer in years PctVegCov Percent vegetation cover **PctConifCov** Percent conifer cover PctBroadLeafCov Percent broadleaf cover TreeBiomass Live tree (> 1inch DBH) biomass, above ground, dry weight

Source

Cutler, D. Richard., Thomas C. Edwards Jr., Karen H. Beard, Adele Cutler, Kyle T. Hess, Jacob Gibson, and Joshua J. Lawler. 2007. Random Forests for Classification in Ecology. Ecology 88(11): 2783-2792.

https://CRAN.R-project.org/package=EZtune/

mtry_compare

Description

A list of data.frames and useful plots for user evaluations of the randomForest hyperparameter mtry.

Usage

```
mtry_compare(
   formula,
   data = NULL,
   scale = FALSE,
   sqrt = TRUE,
   num_var,
   mvec,
   ...
)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	an optional data frame containing the variables in the model. By default the variables are taken from the environment which randomForest is called from.
scale	For permutation based measures such as MSE or Accuracy, should the measures be divided by their "standard errors"? Default is False.
sqrt	Boolean value indicating whether importance metrics should be adjusted via a square root transformation. Default is True.
num_var	Optional integer argument for reducing the number of variables to the top 'num_var' Should be an integer between 1 and the total number of predictor variables in the model or it should be a positive proportion of variables desired.
mvec	Optional vector argument for defining choices of mtry to have the function con- sider. Should be a vector of integers between 1 and the total number of predictor variables in the model. Or it can be a vector of proportions (strictly less than 1) of the number of predictor variables.
	Other parameters to pass to the randomForest function.

Value

A list of data.frames, useful plots, and forest objects for user evaluations of the randomForest hyperparameter mtry.

Examples

```
m <- mtry_compare(factor(Species) ~ ., data = iris, sqrt = TRUE)
m</pre>
```

partial_cor

Description

A list of data.frames and useful plots for user evaluations of correlations and partial correlations of predictors with a given response.

Usage

partial_cor(formula, data = NULL, model = lm, num_var, ...)

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	a data frame containing the variables in the model. By default the variables are taken from the environment which the model is called from.
model	Model to use for extraction partial correlations. Possible model choices are lm, rpart, randomForest, and svm. Default is lm.
num_var	Optional integer argument for reducing the number of variables to the top 'num_var' Should be an integer between 1 and the total number of predictor variables in the model or it should be a positive proportion of variables desired.
	Additional arguments to be passed to model as needed.

Value

A list of data.frames and useful plots for user evaluations of partial correlations.

Examples

```
pcs <- partial_cor(Petal.Length ~ ., data = iris[-5], model = lm)
pcs$plot_y_part_cors</pre>
```

robust_vifs Non-linear Variance Inflation	on Factors
---	------------

Description

A list of data.frames and useful plots for user evaluations of the randomForest hyperparameter mtry.

Usage

```
robust_vifs(formula, data, model = randomForest, log10 = TRUE, num_var, ...)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	an optional data frame containing the variables in the model. By default the variables are taken from the environment which the model is called from.
model	Model to use for extraction partial correlations. Possible model choices are rpart.
log10	Applies a log10 transformation to VIFs when True. Default is True.
num_var	Optional integer argument for reducing the number of variables to the top 'num_var'. Should be an integer between 1 and the total number of predictor variables in the model or it should be a positive proportion of variables desired.
	Additional arguments to be passed to models as needed.

Value

A list of data.frames and useful plots for user evaluations of VIFs.

Examples

```
rv <- robust_vifs(Petal.Length ~ ., data = iris[-5], model = lm)
rv</pre>
```

Index

* boston boston, 2 * lichen lichen, 4 boston, 2 formula, 6-8 ggvip, 3 lichen, 4 mtry_compare, 6 partial_cor, 7 robust_vifs, 7