

Package ‘DriveML’

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Type Package

Title Self-Drive Machine Learning Projects

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Depends R (>= 3.3.0)

Imports sampling(>= 2.8), rmarkdown(>= 1.9), data.table(>= 1.10.4-3), SmartEDA(>= 0.3.1), caTools, ParamHelpers(>= 1.12), mlr(>= 2.15.0), ggplot2(>= 2.2.1), iml

Description Implementing some of the pillars of an automated machine learning pipeline such as (i) Automated data preparation, (ii) Feature engineering, (iii) Model building in classification context that includes techniques such as (a) Regularised regression [1], (b) Logistic regression [2], (c) Random Forest [3], (d) Decision tree [4] and (e) Extreme Gradient Boosting (xgboost) [5], and finally, (iv) Model explanation (using lift chart and partial dependency plots). Accomplishes the above tasks by running the function instead of writing lengthy R codes. Also provides some additional features such as generating missing at random (MAR) variables and automated exploratory data analysis. Moreover, function exports the model results with the required plots in an HTML vignette report format that follows the best practices of the industry and the academia. [1] Gonzales G B and De Saeger (2018) <doi:10.1038/s41598-018-21851-7>, [2] Sperandei S (2014) <doi:10.11613/BM.2014.003>, [3] Breiman L (2001) <doi:10.1023/A:1010933404324>, [4] Kingford C and Salzberg S (2008) <doi:10.1038/nbt0908-1011>, [5] Chen Tianqi and Guestrin Carlos (2016) <doi:10.1145/2939672.2939785>.

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autoDataprep	<i>Automatic data preparation for ML algorithm</i>
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Description

Final data preparation before ML algorithm. Function provides final data set and highlights of the data preparation

Usage

```
autoDataprep(data, target = NULL, missimpute = "default",
  auto_mar = FALSE, mar_object = NULL, dummyvar = TRUE,
  char_var_limit = 12, aucv = 0.02, corr = 0.99,
  outlier_flag = FALSE, interaction_var = FALSE,
  frequent_var = FALSE, uid = NULL, onlykeep = NULL, drop = NULL,
  verbose = FALSE)
```

Arguments

data	[data.frame Required] dataframe or data.table
target	[integer Required] dependent variable (binary or multiclass)
missimpute	[text Optional] missing value imputation using mlr misimpute function. See more methods in details
auto_mar	[character Optional] identify any missing variable which are completely missing at random or not.(default FALSE). If TRUE this will call autoMAR()
mar_object	[character Optional] object created from autoMAR function
dummyvar	[logical Optional] categorical feature engineering i.e. one hot encoding (default is TRUE)
char_var_limit	[integer Optional] default limit is 12 for a dummy variable preparation. Ex: if gender variable has two different value "M" and "F", then gender has 2 level
aucv	[integer Optional] cut off value for AUC based variable selection
corr	[integer Optional] cut off value for correlation based variable selection
outlier_flag	[logical Optional] to add outlier features (default is False)
interaction_var	[logical Optional] bulk interactions transformer for numerical features
frequent_var	[logical Optional] Frequent transformer for categorical features
uid	[character Optional] unique identifier column if any to keep in the final data set
onlykeep	[character Optional] only consider selected variables for data preparation
drop	[character Optional] exclude variable list from the data preparation
verbose	[logical Optional] display executions steps on console. Default FALSE

Details

Missing imputation using impute function from MLR

MLR package have a appropriate way to impute missing value using multiple methods. default value is listed below #'

- mean value for integer variable
- median value for numeric variable
- mode value for character or factor variable

Optional: You might be interested to impute missing variable using ML method. List of algorithms will be handle missing variables in MLR package `listLearners("classif", check.packages = TRUE, properties = "missings")[c("class", "package")]`

Feature engineering

- Missing not completely at random variable using autoMAR function
- Date transformer like year, month, quarter, week
- Frequent transformer counts each categorical value in the dataset
- Interaction transformer using multiplication

- one hot dummy coding for categorical value
- outlier flag and capping variable for numerical value

Feature reduction

- Zero variance using nearZeroVar caret function
- Pearson's Correlation value
- AUC with target variable

Value

list output contains below objects

`complete_data` Complete data set including new novel features based on the functional understanding of the dataset

`master_data` filtered data set based on the input parameter

`final_var_list` list of master variables

`auc_var` list of auc variables

`cor_var` list of correlation variables

`overall_var` all variables in the dataset

`zerovariance` zero variance variables in the dataset

See Also

[impute](#)

Examples

```
#Auto data prep
traindata <- autoDataprep(heart, target = "target_var", missimpute = "default",
dummyvar = TRUE, aucv = 0.02, corr = 0.98, outlier_flag = TRUE,
interaction_var = TRUE, frequent_var = TRUE)
train <- traindata$master

# Print auto data prep object
printautoDataprep(traindata)
```

autoMAR	<i>Function to identify and generate the Missing at Random features (MAR)</i>
---------	---

Description

this function will automatically identify the missing pattern and flag the variable if they are not missing at random based on AUC method

Usage

```
autoMAR(data, aucv = 0.9, strataname = NULL, stratasize = NULL,
        mar_method = "glm")
```

Arguments

data	dataframe or data.table
aucv	AUC cut-off value for the not missing at random variable selection
strataname	vector of stratification variables
stratasize	vector of stratum sample sizes (in the order in which the strata are given in the input data set).
mar_method	missing at random classification method ("glm", "rf"). Default GLM is used (GLM is run faster for high dimension data)

Value

List output including missing variable summary and number of MAR flag variables

Examples

```
# Create missing at random features
marobj <- autoMAR(heart, aucv = 0.9, strataname = NULL, stratasize = NULL, mar_method = "glm")
```

autoMLmodel	<i>Automated machine learning training of models</i>
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Description

Automated training, tuning and validation of machine learning models. Models are tuned and re-sampling validated on an experiment set and trained on the full set and validated and testing on external sets. Classification models tune the probability threshold automatically and returns the results. Each model contains information of performance, the trained model as well as some plots.

Usage

```
autoMLmodel(train, test = NULL, score = NULL, target = NULL,
  testSplit = 0.2, tuneIters = 200, tuneType = "random",
  models = "all", perMetric = "auc", varImp = 20, liftGroup = 50,
  maxObs = 10000, uid = NULL, pdp = FALSE, positive = 1,
  htmlreport = FALSE, seed = 1991, verbose = FALSE)
```

Arguments

train	[data.frame Required] Training set
test	[data.frame Optional] Optional testing set to validate models on. If none is provided, one will be created internally. Default of NULL
score	[data.frame Optional] Optional score the models on best trained model based on AUC. If none is provided, scorelist will be null. Default of NULL
target	[integer Required] If a target is provided classification or regression models will be trained, if left as NULL unsupervised models will be trained. Default of NULL
testSplit	[numeric Optional] Percentage of data to allocate to the test set. Stratified sampling is done. Default of 0.1
tuneIters	[integer Optional] Number of tuning iterations to search for optimal hyper parameters. Default of 10
tuneType	[character Optional] Tune method applied, list of options are: <ul style="list-style-type: none"> • "random" - random search hyperparameter tuning • "frace" - frace uses iterated f-racing algorithm for the best solution from irace package
models	[character Optional] Which models to train. Default is all. List of strings denoting which algorithms to use for the process: <ul style="list-style-type: none"> • randomForestRandom forests using the randomForest package • rangerRandom forests using the ranger package • xgboostGradient boosting using xgboost • rpartdecision tree classification using rpart • glmnetregularised regression from glmnet • logreglogistic regression from stats
perMetric	[character Optional] Model validation metric. Default "auc" <ul style="list-style-type: none"> • auc - Area under the curve; mlr::auc • accuracy - Accuracy; mlr::acc • balancedAccuracy - Balanced accuracy; mlr::bac • brier - Brier score; mlr::brier • f1 - F1 measure; mlr::f1 • meanPrecRecall - Geometric mean of precision and recall; mlr::gpr • logloss - Logarithmic loss; mlr::logloss
varImp	[integer Optional] Number of important features to plot

liftGroup	[integer Optional] Number of lift value to validate the test model performance
maxObs	[numeric Optional] Number of observations in the experiment training set on which models are trained, tuned and resampled on. Default of 40000. If the training set has less than 40k observations all will be used
uid	[character Optional] unique variable to keep in test output data
pdp	[logical Optional] Partial dependence plot for top important variables
positive	[character Optional] positive class for the target variable
htmlreport	[logical Optional] to view the model outcome in html format
seed	[integer Optional] Random number seed for reproducible results
verbose	[logical Optional] display executions steps on console. Default FALSE

Details

all the models trained using mlr train function, all of the functionality in mlr package can be applied to the autoMLmodel outcome

autoMLmodel provides below information of the machine learning classification models

- trainedModels - Model level list output contains trained model object, hyper parameters, tuned data, test data, performance and Model plots
- results - Summary of all trained model result like AUC, Precision, Recall, F1 score
- modelexp - Model gain chart
- predicted_score - Predicted score
- datasummary - Summary of the input data

Value

List output contains trained models and results

See Also

[mlr train](#) [caret train](#) [makeLearner](#) [tuneParams](#)

Examples

```
# Run only Logistic regression model
mymodel <- autoMLmodel( train = heart, test = NULL, target = 'target_var',
  testSplit = 0.2, tuneIters = 10, tuneType = "random", models = "logreg",
  varImp = 10, liftGroup = 50, maxObs = 4000, uid = NULL, seed = 1991)
```

`autoMLReport`*Display autoMLmodel output in HTML format using Rmarkdown*

Description

This function will generate R markdown report for DriveML model object

Usage

```
autoMLReport(mlobject, mldata = NULL, op_file = NULL, op_dir = NULL)
```

Arguments

<code>mlobject</code>	[autoMLmodel Object Required] autoMLmodel function output
<code>mldata</code>	[autoDataprep Object Optional] autoDataprep function output
<code>op_file</code>	[character Required] output file name (.html)
<code>op_dir</code>	[character Optional] output path. Default path is current working directory

Details

Using this function easily present the model outcome in standard HTML format without writing Rmarkdown scripts

Value

HTML R Markdown output

Examples

```
## Creating HTML report

autoMLReport(heart.model, mldata = NULL, op_file = "sample.html", op_dir = tempdir())
```

`autoPDP`*Generate partial dependence plots*

Description

Partial dependence plots (PDPs) help you to visualize the relationship between a subset of the features and the response while accounting for the average effect of the other predictors in the model. They are particularly effective with black box models like random forests and support vector machines.

Usage

```
autoPDP(train, trainedModel, target, feature, sample = 0.5, modelname,
        seed = 1991)
```

Arguments

train	[data.frame Required] Training sample used to train ML model
trainedModel	[model object Required] The object holding the machine learning model and the data
target	[character Optional] Target variable name. Specify target variable if model object is other than MLR or driveML
feature	[character Optional] The feature name for which to compute the effects
sample	[numeric Optional] Percentage of sample to be considered for training set for faster computation. Default of 0.5
modelName	[character Optional] specify which model to be plotted
seed	[integer Optional] Random seed number. Default 121

Value

List object containing a plot for each feature listed.

See Also

[FeatureEffects](#) [plotPartialDependence](#) [partial](#)

Examples

```
## Example using DriveML model object
mymodel = heart.model
pdp_chol = autoPDP(heart, mymodel, feature = "chol", sample = 0.8, seed = 1234)

# Type one MLR package
mod <- mlr::train(makeLearner("classif.ranger"), iris.task)
cc = autoPDP(iris, mod, feature = c("Sepal.Length", "Sepal.Width", "Petal.Length",
                                   "Petal.Width"), sample = 1, seed = 121)

# Type 2 DriveML object
hearML <- autoMLmodel(heart, target = "target_var", testSplit = 0.2,
                     tuneIters = 10, tuneType = "random",
                     models = "all", varImp = 20, liftGroup = 50, positive = 1, seed = 1991)
cc = autoPDP(heart, hearML, feature = "chol", sample = 0.8, seed = 1234)

cc1 = autoPDP(heart, trainedModel, target = "target_var", feature = "chol",
             sample = 1, modelName = "logreg", seed = 121)

# Type 3 other ML object
library(randomForest)
library(MASS)
```

```
rf = randomForest(medv ~ ., data = Boston, ntree = 50)
cc = autoPDP(Boston, rf, target = "medv", feature = "nox", sample = 1, seed = 121)
```

generateFeature *Automated column transformer*

Description

This function automatically scans through each variable and generate features based on the type listed in the detail

Usage

```
generateFeature(data, varlist, type = "Frequent", method = NULL)
```

Arguments

data	dataframe or data.table
varlist	variable list to generate the additional features
type	variable transformation type 'Dummy', 'Outlier', 'Frequent', 'Interaction'
method	input for variabe transforamtion. For type = 'Frequent' then type should be 'Frequency' or 'Percent'. Other type method list is provided in details

Details

This function is for generating features based on diffenret transformation methods like interaction, outliers, Dummy coding etc.

Interaction type

- multiply - multipliacion
- add - addition
- substract - subtraction
- divide - division

Frequency type

- Frequency - Frequency
- Percent - Percentage

Outlier type

- Flag - Falg outlier values like 1 or 0
- Capping - Impute outlier value by 95th or 5th percentile value

Date type

- Year
- Month
- Quarter
- Week

Value

generated transformed features

Examples

```
# Generate interaction features
generateFeature(heart, varlist = c("cp", "chol", "trestbps"), type = "Interaction",
method = "add")
generateFeature(heart, varlist = c("cp", "chol", "trestbps"), type = "Interaction",
method = "multiply")

# Generate frequency features
generateFeature(heart, varlist = c("cp", "thal"), type = "Frequent", method = "Percent")
generateFeature(heart, varlist = c("cp", "thal"), type = "Frequent", method = "Frequency")
```

heart

Dataset Heart Disease - Classifications

Description

This database contains 76 attributes, but all published experiments refer to using a subset of 14 of them. In particular, the Cleveland database is the only one that has been used by ML researchers to this date. The "goal" field refers to the presence of heart disease in the patient. It is integer valued from 0 (no presence) to 4.

Usage

heart

Format

A data frame with 303 rows and 14 variables:

```
age integer Age
sex integer Sex
cp integer chest pain type (4 values)
trestbps integer resting blood pressure
chol integer serum cholestorol in mg/dl
fbs integer fasting blood sugar > 120 mg/dl
```

restecg integer resting electrocardiographic results (values 0,1,2)
 thalach integer maximum heart rate achieved
 exang integer exercise induced angina
 oldpeak double oldpeak = ST depression induced by exercise relative to rest
 slope integer the slope of the peak exercise ST segment
 ca integer number of major vessels (0-3) colored by flourosopy
 thal integer thal: 3 = normal; 6 = fixed defect; 7 = reversable defect
 target_var integer the presence of heart disease in the patient. It is integer valued from 0 (no presence) to 4

Value

sample data

Source

<https://www.kaggle.com/cdabakoglu/heart-disease-classifications-machine-learning>

Examples

```
## Load heart data
data(heart)
```

heart.model

Heart Classification Drive ML Model.

Description

Contains the task ('heart.model').

Usage

heart.model

Format

An object of class autoMLmodel of length 6.

Value

heart data driveML sample model output

References

See <https://www.kaggle.com/cdabakoglu/heart-disease-classifications-machine-learning>

Examples

```
## Sample model object  
modelobj <- heart.model
```

misspattern

Missing pattern analysis for missing data

Description

this function for summarise the missing variable, missing pattern identification, classifying the columns based on pattern of missing values.

Usage

```
misspattern(data, mfeature, drop = 0.99, print = FALSE)
```

Arguments

data	[data.frame Required] data set with missing values
mfeature	[character Required] only missing variable name
drop	[numeric optional] drop variable percentage. Example, if drop = 0.9, function will automatically drop 90per missing columns from the data set
print	[character optional] default print is FALSE

Value

final variable list, summary of missing data analysis

Examples

```
## Sample iris data  
mdata <- iris  
mobject <- misspattern(mdata, mfeature = c("Sepal.Length", "Petal.Length"), drop = 0.99, print = F)
```

predictAutoMAR *Extract predictions and MAR columns from autoMAR objects*

Description

this function can be used for autoMAR objects to generate the variable for missing variable not completely at random

Usage

```
predictAutoMAR(x, data, mar_var = NULL)
```

Arguments

x	[autoMAR object Required] autoMAR object for which prediction is desired
data	[data.frame Required] prediction data set to prepare the autoMAR outcomes
mar_var	[character list Optional] list of predefined mar variables

Value

flagged variables for missing not completely at random variable

Examples

```
## Missing at random features
train <- heart[1 : 199, ]
test <- heart[200 : 300, ]
marobj <- autoMAR (train, aucv = 0.9, strataname = NULL, stratasize = NULL, mar_method = "glm")

## print summary in console
testobj <- predictAutoMAR(marobj, test)
```

predictDataprep *Extract predictions and generate columns from autoDataprep objects*

Description

this function can be used for autoDataprep objects to generate the same for validation

Usage

```
predictDataprep(x, data)
```

Arguments

x	[autoDataprep object Required] autoDataprep object for which prediction is desired
data	[data.frame Required] prediction data set to prepare the MAR columns

Value

master data set same as train data set

Examples

```
## Sample train data set
train <- heart[1:200, ]
test <- heart[201:350, ]
traindata <- autoDataprep(train, target = "target_var", missimpute = "default",
dummyvar = TRUE, aucv = 0.02, corr = 0.98, outlier_flag = TRUE,
interaction_var = TRUE, frequent_var = TRUE)
train <- traindata$master

## Predict same features for test set
test <- predictDataprep(traindata, test)
```

printautoDataprep *Print Method for the autoDataprep Class*

Description

Print the result of autoDataprep object

Usage

```
printautoDataprep(x)
```

Arguments

x	[Object Required] an object of class autoDataprep
---	---

Value

Print summary autoDataprep function results on console

Examples

```
#Auto data prep
traindata <- autoDataprep(heart, target = "target_var", missimpute = "default",
  dummyvar = TRUE, aucv = 0.02, corr = 0.98, outlier_flag = TRUE,
  interaction_var = TRUE, frequent_var = TRUE)

# Print auto data prep object
printautoDataprep(traindata)
```

printautoMAR	<i>Print Method for the autoMAR Class Print the result of autoMAR object</i>
--------------	--

Description

Print Method for the autoMAR Class Print the result of autoMAR object

Usage

```
printautoMAR(x)
```

Arguments

x [Object | Required] an object of class autoMAR

Value

Print summary of autoMAR output in console

Examples

```
## Missing at random features
marobj <- autoMAR (heart, aucv = 0.9, strataname = NULL, stratasize = NULL, mar_method = "glm")

## print summary in console
printautoMAR(marobj)
```

smartEDA	<i>SmartEDA - functions that automates most of exploratory analyses tasks in modeling</i>
----------	---

Description

SmartEDA includes multiple custom functions to perform initial exploratory analysis on any input data describing the structure and the relationships present in the data. The generated output can be obtained in both summary and graphical form. The graphical form or charts can also be exported as reports.

Usage

```
smartEDA(data, Template = NULL, Target = NULL, label = NULL,
         theme = "Default", op_file = NULL, op_dir = getwd(), sc = NULL,
         sn = NULL, Rc = NULL)
```

Arguments

data	a data frame
Template	R markdown template (.rmd file)
Target	dependent variable. If there is no defined target variable then keep as it is NULL.
label	target variable descriptions, not a mandatory field
theme	customized ggplot theme (default SmartEDA theme) (for Some extra themes use Package: ggthemes)
op_file	output file name (.html)
op_dir	output path
sc	sample number of plots for categorical variable. User can decide how many number of plots to depict in html report.
sn	sample number of plots for numerical variable. User can decide how many number of plots to depict in html report.
Rc	reference category of target variable. If Target is categorical then Pclass value is mandatory and which should not be NULL

Details

SmartEDA has four major functionalities 1. Descriptive statistics

- Numerical variable summary :
- ExpNumStat - Summary statistics for numerical variables [ExpNumStat](#)
- Categorical variable summary :
- ExpCatStat - Function provides summary statistics for all character or categorical columns in the dataframe [ExpCatStat](#)

- ExpCTable - Function to create frequency and custom tables [ExpCTable](#)

2. Data visualization

- Numerical variable plot :
- ExpNumViz - Distributions of numeric variables [ExpNumViz](#)
- Categorical variable plot :
- ExpCatViz - Distributions of categorical variables [ExpCatViz](#)
- Normality testing plot:
- ExpOutQQ - Quantile Quantile Plots [ExpOutQQ](#)
- ExpParcoord - Parallel Co ordinate plots [ExpParcoord](#)

3. Custom tables

- Customized summary statistics :
- ExpCustomStat - Customized summary statistics [ExpCustomStat](#)

4. EDA report

- Function to create HTML EDA report :
- ExpReport - Function to create HTML EDA report [ExpReport](#)

Value

HTML Rmarkdown output file in .html format

Source

Useful links:

- CRAN page <https://CRAN.R-project.org/package=SmartEDA>
- JOSS <https://doi.org/10.21105/joss.01509>

Examples

```
# Generate complete EDA report
smartEDA(iris, op_file="eda_report.html", op_dir = tempdir(), sc = NULL, sn = 2)
```

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