Package 'TidyML'

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Title Machine Learning Modelling For Everyone

Version 0.0.0.9000

Description
TidyML is a minimal library focused on providing all the essential tools for the workflow of a machine learning modelling process. The whole process is divided into 5 steps: preprocessing() -> build_model() -> fine_tuning() -> show_results() -> sensitivity_analysis()

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Encoding UTF-8

Roxygen list(markdown = TRUE)

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Depends R (>= 2.10),
tidyverse

Imports broom,
dials,
parsnip,
recipes,
rsample,
tune,
workflows,
yardstick,
R6,
magrittr,
vip,
glue,
fmsb,
tidyr,
ggpubr,
innsight,
torch,
shapr,
DiagrammeR

Suggests testthat (>= 3.0.0)

Config/testthat/edition 3

URL https://github.com/JMartinezGarcia/TidyML

BugReports https://github.com/JMartinezGarcia/TidyML/issues

LazyData true

Contents

`build_model` *Create ML Model*

Description

Create ML Model

Usage

```
build_model(analysis_object, model_name, hyperparameters = NULL)
```

Arguments

`analysis_object`	analysis_object created from preprocessing function.
`hyperparameters`	Hyperparameters of the ML model. List containing the name of the hyperparameter and its value or range of values.
`model_names`	Name of the ML Model. A string of the model name: "Neural Network", "Random Forest", "SVM" or "XGBOOST".

Value

Updated analysis_object

Hyperparameters

Neural Network:

Parsnip model using **brulee** engine. Hyperparameters:

- **hidden_units**: Number of Hidden Neurons. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(5, 20).
- **activation**: Activation Function. A vector with any of ("relu", "sigmoid", "tanh") or NULL for default values c("relu", "sigmoid", "tanh").
- **learn_rate**: Learning Rate. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(-3, -1) in log10 scale.

Random Forest:

Parsnip model using **ranger** engine. Hyperparameters:

- **trees**: Number of Trees. A single value, a vector with range values `c(min_val, max_val)`. Default range c(100, 300).

- **mtry**: Number of variables randomly selected as candidates at each split. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(3, 8).
- **min_n**: Minimum Number of samples to split at each node. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(2, 25).

XGBOOST:

Parsnip model using **xgboost** engine. Hyperparameters:

- **trees**: Number of Trees. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(100, 300).
- **mtry**: Number of variables randomly selected as candidates at each split. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(3, 8).
- **min_n**: Minimum Number of samples to split at each node. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(5, 25).
- **tree_depth**: Maximum tree depth. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(3, 10).
- **learn_rate**: Learning Rate. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(-4, -1) in log10 scale.
- **loss_reduction**: Minimum loss reduction required to make a further partition on a leaf node. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(-5, 1.5) in log10 scale.

SVM:

Parsnip model using **kernlab** engine. Hyperparameters:

- **cost**: Penalty parameter that regulates model complexity and misclassification tolerance. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(-3, 3) in log10 scale.
- **margin**: Distance between the separating hyperplane and the nearest data points. A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(0, 0.2).
- **type**: Kernel to be used. A single value from ("linear", "rbf", "polynomial")
- **rbf_sigma**: A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(-5, 0) in log10 scale.
- **degree**: Polynomial Degree (polynomial kernel only). A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(1, 3).
- **scale_factor**: Scaling coefficient applied to inputs. (polynomial kernel only) A single value, a vector with range values `c(min_val, max_val)` or NULL for default range c(-5, -1) in log10 scale.

`fine_tuning` *Fine Tune ML Model*

Description

Fine Tune ML Model

Usage

```
fine_tuning(analysis_object, tuner, metrics, plot_results = F, verbose = FALSE)
```

Arguments

analysis_object	analysis_object created from build_model function.
tuner	Name of the Hyperparameter Tuner. A string of the tuner name: "Bayesian Optimization" or "Grid Search CV".
metrics	Metric used for Model Selection. A string of the name of metric (see Metrics).
plot_results	Whether to plot the tuning results. Boolean TRUE or FALSE (default).
verbose	Whether to show tuning process. Boolean TRUE or FALSE (default).

Value

Updated analysis_object

Tuners

Bayesian Optimization:

- Initial data points: 20
- Maximum number of iterations: 25
- Convergence after 5 iterations without improvement
- Train / Validation / Test : 0.6 / 0.2 / 0.2

Grid Search CV:

- Number of Folds: 5
- Max grid size per hyperparameter: 10
- Train / Test : 0.75 / 0.25

Metrics

Regression Metrics:

- rmse
- mae
- mpe
- mape
- ccc
- smape
- rpiq
- rsq

Classification Metrics:

- accuracy
- bal_accuracy
- recall
- sensitivity
- specificity
- kap
- f_meas
- mcc
- j_index

- detection_prevelance
- roc_auc
- pr_auc
- gain_capture
- brier_class
- roc_aunp

`preprocessing` *Preprocessing Data Matrix*

Description

Preprocessing Data Matrix

Usage

```
preprocessing(
  df,
  formula,
  task = "regression",
  num_vars = NULL,
  cat_vars = NULL,
  norm_num_vars = "all",
  encode_cat_vars = "all",
  y_levels = NULL
)
```

Arguments

`df`	Input DataFrame. Either a data.frame or tibble.
`formula`	Modelling Formula. A string of characters or formula.
`task`	Modelling Task. Either "regression" or "classification".
`num_vars`	Optional vector of names of the numerical features.
`cat_vars`	Optional vector of names of the categorical features.
`norm_num_vars`	Normalize numeric features as z-scores. Either vector of names of numerical features to be normalized or "all" (default).
`encode_cat_vars`	One Hot Encode Categorical Features. Either vector of names of categorical features to be encoded or "all" (default).
`y_levels`	Optional ordered vector with names of the target variable levels (Classification task only).

Value

An analysis_object

sensitivity_analysis *Perform Sensitivity Analysis and Interpretable ML methods*

Description

Perform Sensitivity Analysis and Interpretable ML methods

Usage

```
sensitivity_analysis(analysis_object, methods = c("PFI"), metric = NULL)
```

Arguments

analysis_object	analysis_object created from fine_tuning function.
metric	Metric used for "PFI" method (Permutation Feature Importance). A string of the name of metric (see Metrics).
type	Type of method used. A string of the method name: "PFI" (Permutation Feature Importance), "SHAP" (SHapley Additive exPlanations), "Integrated Gradients" (Neural Network only) or "Olden" (Neural Network only).

Value

Updated analysis_object

show_results *Showcase Summary Results and Plots*

Description

Showcase Summary Results and Plots

Usage

```
show_results(
  analysis_object,
  summary = FALSE,
  roc_curve = FALSE,
  pr_curve = FALSE,
  gain_curve = FALSE,
  lift_curve = FALSE,
  dist_by_class = FALSE,
  reliability_plot = FALSE,
  confusion_matrix = FALSE,
  scatter_residuals = FALSE,
  scatter_predictions = FALSE,
  residuals_dist = FALSE,
  new_data = "test"
)
```

Arguments

analysis_object	analysis_object created from fine_tuning function.
summary	Whether to plot summary results table. Boolean (FALSE by default).
roc_curve	Whether to plot ROC Curve (Classification task only). Boolean (FALSE by default).
pr_curve	Whether to plot ROC Curve (Classification task only). Boolean (FALSE by default).
gain_curve	Whether to plot ROC Curve (Classification task only). Boolean (FALSE by default).
lift_curve	Whether to plot ROC Curve (Classification task only). Boolean (FALSE by default).
dist_by_class	Whether to plot distribution of output probability by class (Classification task only). Boolean (FALSE by default).
reliability_plot	Whether to plot Reliability Plot (Binary Classification task only). Boolean (FALSE by default).
confusion_matrix	Whether to Confusion Matrix (Classification task only). Boolean (FALSE by default).
scatter_residuals	Whether to plot Residuals vs Predictions (Regression task only). Boolean (FALSE by default).
scatter_predictions	Whether to plot Predictions vs Observed (Regression task only). Boolean (FALSE by default).
residuals_dist	Whether to plot Residuals Distribution (Regression task only). Boolean (FALSE by default).
new_data	Data to be used for Confusion Matrix, Reliability Plot, Distribution by Class Plot, Residuals vs Predictions Plot, Predictions vs Observed Plot and Residuals Distribution Plot. A string with the name of the data_set: "train", "validation", "test" (default) or "all".

Value

Updated analysis_object

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sim_data *Example Data Set*

Description

This dataset contains simulated data of a psychometric trial.

Usage

```
sim_data
```

Format

A data frame with 1000 rows and 10 columns:

psych_well Psychological Wellbeing Indicator. Continous with (0,100)

psych_well_bin Psychological Wellbeing Binary Indicator. Factor with ("Low", "High")

psych_well_pol Psychological Wellbeing Polytomic Indicator. Factor with ("Low", "Somewhat", "Quite a bit", "Very Much")

gender Patient Gender. Factor ("Female", "Male")

age Patient Age. Continous (18, 85)

socioec_status Socioeconomial Status Indicator. Factor ("Low", "Medium", "High")

emot_intel Emotional Intelligence Indicator. Continous (24, 120)

resilience Resilience Indicator. Continous (4, 20)

depression Depression Indicator. Continous (0, 63)

life_sat Life Satisfaction Indicator. Continous (5, 35)

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