Package 'fastml'

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Title Fast Machine Learning Model Training and Evaluation				
Version 0.3.0				
Description Streamlines the training, evaluation, and comparison of multiple machine learning models with minimal code by providing comprehensive data preprocessing and support for a wide range of algorithms with hyperparam ter tuning. It offers performance metrics and visualization tools to facilitate efficient and effective machine learning workflows.				
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evaluate_models

Evaluate Models Function

Description

Evaluates the trained models on the test data and computes performance metrics.

Usage

```
evaluate_models(models, train_data, test_data, label, task, metric = NULL)
```

Arguments

models A list of trained model objects.

train_data Preprocessed training data frame.

test_data Preprocessed test data frame.

label Name of the target variable.

task Type of task: "classification" or "regression".

metric The performance metric to optimize (e.g., "accuracy", "rmse").

Value

A list with two elements:

performance A named list of performance metric tibbles for each model.

predictions A named list of data frames with columns including truth, predictions, and probabilities per model.

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explain

Explain the fastml_model (DALEX + SHAP + Permutation-based VI)

Description

Provides model explainability using DALEX. This function: - Creates a DALEX explainer. - Computes permutation-based variable importance with boxplots showing variability, displays the table and plot. - Computes partial dependency-like model profiles if 'features' are provided. - Computes Shapley values (SHAP) for a sample of the training observations, displays the SHAP table, and plots a summary bar chart of mean(ISHAP valuel) per feature. For classification, it shows separate bars for each class.

Usage

```
explain(
  object,
 method = "dalex",
  features = NULL,
  grid_size = 20,
  shap\_sample = 5,
  vi_iterations = 10,
  colormap = "viridis",
  top_features = NULL,
  seed = 123,
)
```

Arguments

object

method	Currently only "dalex" is supported.
features	Character vector of feature names for partial dependence (model profiles). Default NULL.
grid_size	Number of grid points for partial dependence. Default 20.
shap_sample	Integer number of observations from processed training data to compute SHAP values for. Default 5.
vi_iterations	Integer. Number of permutations for variable importance. Default 10.
colormap	Character. Name of a color palette to use (e.g., "viridis"). Default "viridis".
ton features	Integer I imit the SHAP summary plot to top N features by mean als SHAP

Integer. Limit the SHAP summary plot to top N features by mean abs SHAP. top_features

Default NULL (no limit).

A fastml_model object.

Integer. A value specifying the random seed. seed

Additional arguments (not currently used).

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Details

1. Custom number of permutations for VI (vi_iterations): You can now specify how many permutations (B) to use for permutation-based variable importance. More permutations yield more stable estimates but take longer.

- 2. Custom color palette (colormap): A 'colormap' parameter allows you to select a color palette (e.g., "viridis") for SHAP summary plots and variable importance plots. This improves aesthetics over default palettes.
- 3. Top Features in SHAP Summary (top_features): You can limit the SHAP summary plot to the top N features by mean absolute SHAP value. This helps focus on the most influential features.
- 4. Support for calibration plot if probably is available (calibration): If 'calibration = TRUE' and 'probably' is installed, it attempts to produce a model-based calibration plot (e.g., 'cal_plot_logistic'). This provides a smoothed, nonparametric view of model calibration.
- 5. Better error messages and checks: Improved checks and messages if certain packages or conditions are not met.

Value

Prints DALEX explanations: variable importance table & plot, model profiles (if any), SHAP table & summary plot, and optionally a calibration plot.

fastml

Fast Machine Learning Function

Description

Trains and evaluates multiple classification or regression models automatically detecting the task based on the target variable type.

```
fastml(
  data,
  label,
  algorithms = "all",
  test_size = 0.2,
  resampling_method = "cv",
  folds = ifelse(grepl("cv", resampling_method), 10, 25),
  repeats = ifelse(resampling_method == "repeatedcv", 1, NA),
  tune_params = NULL,
  metric = NULL,
  n_cores = 1,
  stratify = NULL,
  impute_method = "error",
  encode_categoricals = TRUE,
  scaling_methods = c("center", "scale"),
```

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```
summaryFunction = NULL,
use_default_tuning = FALSE,
tuning_strategy = "grid",
tuning_iterations = 10,
early_stopping = FALSE,
adaptive = FALSE,
seed = 123,
recipe = NULL
)
```

Arguments

data A data frame containing the features and target variable.

label A string specifying the name of the target variable.

algorithms A vector of algorithm names to use. Default is "all" to run all supported algo-

rithms.

test_size A numeric value between 0 and 1 indicating the proportion of the data to use for

testing. Default is 0.2.

resampling_method

A string specifying the resampling method for model evaluation. Default is "cv" (cross-validation). Other options include "none", "boot", "repeatedcy", etc.

folds An integer specifying the number of folds for cross-validation. Default is 10 for

methods containing "cv" and 25 otherwise.

repeats Number of times to repeat cross-validation (only applicable for methods like

"repeatedcv").

tune_params A list specifying hyperparameter tuning ranges. Default is NULL.

metric The performance metric to optimize during training. Default depends on the

task.

n_cores An integer specifying the number of CPU cores to use for parallel processing.

Default is 1.

stratify Logical indicating whether to use stratified sampling when splitting the data.

Default is TRUE for classification and FALSE for regression.

impute_method Method for handling missing values. Options include:

"medianImpute" Impute missing values using median imputation.

"knnImpute" Impute missing values using k-nearest neighbors.

"bagImpute" Impute missing values using bagging.

"remove" Remove rows with missing values from the data.

"error" Do not perform imputation; if missing values are detected after preprocessing, stop execution with an error.

NULL Equivalent to "error". No imputation is performed, and the function will stop if missing values are present.

Default is "error".

encode_categoricals

Logical indicating whether to encode categorical variables. Default is TRUE.

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scaling_methods

Vector of scaling methods to apply. Default is c("center", "scale").

summaryFunction

A custom summary function for model evaluation. Default is NULL.

use_default_tuning

Logical indicating whether to use default tuning grids when tune_params is NULL. Default is FALSE.

tuning_strategy

A string specifying the tuning strategy. Options might include "grid", "bayes", or "none". Default is "grid".

tuning_iterations

Number of tuning iterations (applicable for Bayesian or other iterative search methods). Default is 10.

early_stopping Logical indicating whether to use early stopping in Bayesian tuning methods (if

supported). Default is FALSE.

adaptive Logical indicating whether to use adaptive/racing methods for tuning. Default

is FALSE.

seed An integer value specifying the random seed for reproducibility.

recipe A user-defined recipe object for custom preprocessing. If provided, internal

recipe steps (imputation, encoding, scaling) are skipped.

Details

Fast Machine Learning Function

Trains and evaluates multiple classification or regression models. The function automatically detects the task based on the target variable type and can perform advanced hyperparameter tuning using various tuning strategies.

Value

An object of class fastml_model containing the best model, performance metrics, and other information.

Examples

```
# Example 1: Using the iris dataset for binary classification (excluding 'setosa')
data(iris)
iris <- iris[iris$Species != "setosa", ] # Binary classification
iris$Species <- factor(iris$Species)

# Train models with Bayesian optimization
model <- fastml(
   data = iris,
   label = "Species",
   algorithms = c("random_forest", "xgboost", "svm_radial")
)
# View model summary</pre>
```

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```
summary(model)

# Example 2: Using the mtcars dataset for regression
data(mtcars)

# Train models
model <- fastml(
   data = mtcars,
   label = "mpg",
   algorithms = c("random_forest", "xgboost", "svm_radial")
)

# View model summary
summary(model)</pre>
```

load_model

Load Model Function

Description

Loads a trained model object from a file.

Usage

```
load_model(filepath)
```

Arguments

filepath

A string specifying the file path to load the model from.

Value

An object of class fastml_model.

plot.fastml_model

Plot Function for fastml_model

Description

Generates plots to compare the performance of different models.

```
## S3 method for class 'fastml_model'
plot(x, ...)
```

predict.fastml_model

Arguments

x An object of class fastml_model.

... Additional arguments (not used).

Value

Displays comparison plots of model performances.

Description

Makes predictions on new data using the trained model.

Usage

```
## S3 method for class 'fastml_model'
predict(object, newdata, type = "auto", ...)
```

Arguments

object An object of class fastml_model.

newdata A data frame containing new data for prediction.

type Type of prediction. Default is "auto", which returns class labels for classifica-

tion and numeric predictions for regression. Other options include "prob" for

class probabilities (classification only).

... Additional arguments (not used).

Value

A vector or data frame of predictions.

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save_model

Save Model Function

Description

Saves the trained model object to a file.

Usage

```
save_model(model, filepath)
```

Arguments

model An o

An object of class fastml_model.

filepath

A string specifying the file path to save the model.

Value

No return value, called for its side effect of saving the model object to a file.

```
summary.fastml_model Summary Function for fastml_model (Using yardstick for ROC Curves)
```

Description

Provides a concise, user-friendly summary of model performances. For classification: - Shows Accuracy, F1 Score, Kappa, Precision, ROC AUC, Sensitivity, Specificity. - Produces a bar plot of these metrics. - Shows ROC curves for binary classification using yardstick::roc_curve(). - Displays a confusion matrix and a calibration plot if probabilities are available.

```
## $3 method for class 'fastml_model'
summary(
   object,
   sort_metric = NULL,
   plot = TRUE,
   combined_roc = TRUE,
   notes = "",
   ...
)
```

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Arguments

object An object of class fastml_model.

sort_metric The metric to sort by. Default uses optimized metric.

plot Logical. If TRUE, produce bar plot, yardstick-based ROC curves (for binary

classification), confusion matrix (classification), smooth calibration plot (if prob-

abilities), and residual plots (regression).

combined_roc Logical. If TRUE, combined ROC plot; else separate ROC plots.

notes User-defined commentary.
... Additional arguments.

Details

For regression: - Shows RMSE, R-squared, and MAE. - Produces a bar plot of these metrics. - Displays residual diagnostics (truth vs predicted, residual distribution).

Value

Prints summary and plots if requested.

train_models

Train Specified Machine Learning Algorithms on the Training Data

Description

Trains specified machine learning algorithms on the preprocessed training data.

```
train_models(
  train_data,
 label.
  task,
  algorithms,
  resampling_method,
  folds,
  repeats,
  tune_params,
 metric,
  summaryFunction = NULL,
  seed = 123,
  recipe,
  use_default_tuning = FALSE,
  tuning_strategy = "grid",
  tuning_iterations = 10,
  early_stopping = FALSE,
  adaptive = FALSE
)
```

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Arguments

train_data Preprocessed training data frame.

label Name of the target variable.

task Type of task: "classification" or "regression".

algorithms Vector of algorithm names to train.

resampling_method

Resampling method for cross-validation (e.g., "cv", "repeatedcv", "boot", "none").

folds Number of folds for cross-validation.

repeats Number of times to repeat cross-validation (only applicable for methods like

"repeatedcv").

tune_params List of hyperparameter tuning ranges.

metric The performance metric to optimize.

summaryFunction

A custom summary function for model evaluation. Default is NULL.

seed An integer value specifying the random seed for reproducibility.

recipe A recipe object for preprocessing.

use_default_tuning

Logical indicating whether to use default tuning grids when tune_params is

NULL.

tuning_strategy

A string specifying the tuning strategy ("grid", "bayes", or "none"), possibly

with adaptive methods.

tuning_iterations

Number of iterations for iterative tuning methods.

early_stopping Logical for early stopping in Bayesian tuning.

adaptive Logical indicating whether to use adaptive/racing methods.

Value

A list of trained model objects.

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