

Package ‘r2pmml’

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Type Package

License AGPL-3

Title Convert R Models to PMML

Description R wrapper for the JPMML-R library <<https://github.com/jpmml/jpmml-r>>, which converts R models to Predictive Model Markup Language (PMML).

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URL <https://github.com/jpmml/r2pmml>

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NeedsCompilation no

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Suggests caret, e1071, earth, evtree, glmnet, mlbench, partykit, randomForest, ranger, xgboost

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R topics documented:

as.scorecard	2
decorate	3
decorate.default	3
decorate.earth	4
decorate.elmNN	4
decorate.glmnet	5
decorate.party	5
decorate.randomForest	6
decorate.ranger	7

decorate.svm.formula	7
decorate.train	8
decorate.xgb.Booster	8
genDMatrix	9
genFMap	10
r2pmmml	10
verify	11
verify.default	12
verify.glm	12
verify.train	13
writeFMap	13

Index	14
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as.scorecard	<i>Converts a "glm" object to a "scorecard" object.</i>
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Description

Converts a "glm" object to a "scorecard" object.

Usage

```
as.scorecard(glm, odds = 10, base_points = 500, pdo = 100)
```

Arguments

glm	A "glm" object with binomial family link function.
odds	Odds ratio at base odds.
base_points	Points where odds ratio is defined.
pdo	Points to double the odds.

Value

A "scorecard" object.

decorate	<i>Dispatches execution to the most appropriate model decoration function.</i>
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Description

Dispatches execution to the most appropriate model decoration function.

Usage

```
decorate(x, ...)
```

Arguments

- | | |
|-----|--|
| x | A model object. |
| ... | Arguments to pass on to the selected function. |

decorate.default	<i>Decorates a model object with "preProcess" and "pmml_options" elements.</i>
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Description

Decorates a model object with "preProcess" and "pmml_options" elements.

Usage

```
## Default S3 method:  
decorate(x, preprocess = NULL, pmml_options = NULL,  
...)
```

Arguments

- | | |
|--------------|---|
| x | The model object. |
| preprocess | A "train::preProcess" object. |
| pmml_options | A list of model type-dependent PMML conversion options. |
| ... | Further arguments. |

decorate.earth *Decorates an "earth" object with an "xlevels" element.*

Description

Decorates an "earth" object with an "xlevels" element.

Usage

```
## S3 method for class 'earth'  
decorate(x, data, ...)
```

Arguments

- | | |
|------|--|
| x | An "earth" object. |
| data | The training dataset. |
| ... | Arguments to pass on to the "decorate.default" function. |

decorate.elmNN *Decorates an "elmNN" object with a "model" element.*

Description

Decorates an "elmNN" object with a "model" element.

Usage

```
## S3 method for class 'elmNN'  
decorate(x, data, ...)
```

Arguments

- | | |
|------|--|
| x | An "elmNN" object. |
| data | The training dataset. |
| ... | Arguments to pass on to the "decorate.default" function. |

decorate.glmnet	<i>Decorates a "glmnet" object with a "lambda.s" element.</i>
-----------------	---

Description

Decorates a "glmnet" object with a "lambda.s" element.

Usage

```
## S3 method for class 'glmnet'
decorate(x, lambda.s, ...)
```

Arguments

x	A "glmnet" object.
lambda.s	The best lambda value. Must be one of listed "glmnet\$lambda" values.
...	Arguments to pass on to the "decorate.default" function.

Examples

```
library("glmnet")
library("r2pmml")

data(iris)
iris_x = as.matrix(iris[, -ncol(iris)])
iris_y = iris[, ncol(iris)]
iris.glmnet = glmnet(x = iris_x, y = iris_y, family = "multinomial")
iris.glmnet = decorate(iris.glmnet, lambda.s = iris.glmnet$lambda[49])
r2pmml(iris.glmnet, file.path(tempdir(), "Iris-GLMNet.pmml"))
```

decorate.party	<i>Decorates a "party" object with a "predicted" element.</i>
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Description

Decorates a "party" object with a "predicted" element.

Usage

```
## S3 method for class 'party'
decorate(x, ...)
```

Arguments

- x A "party" object.
- ... Arguments to pass on to the "decorate.default" function.

Examples

```
library("evtree")
library("r2pmml")

data(iris)
iris.party = evtree(Species ~ ., data = iris,
                     control = evtree.control(max_depth = 3))
iris.party = decorate(iris.party)
r2pmml(iris.party, file.path(tempdir(), "Iris-Party.pmml"))
```

decorate.randomForest *Decorates a "randomForest" object with PMML conversion options.*

Description

Decorates a "randomForest" object with PMML conversion options.

Usage

```
## S3 method for class 'randomForest'
decorate(x, compact = FALSE, ...)
```

Arguments

- x A "randomForest" object.
- compact A flag controlling if decision trees should be transformed from binary splits (FALSE) to multi-way splits (TRUE) representation.
- ... Arguments to pass on to the "decorate.default" function.

decorate.ranger *Decorates a "ranger" object with a "variable.levels" element.*

Description

Decorates a "ranger" object with a "variable.levels" element.

Usage

```
## S3 method for class 'ranger'  
decorate(x, data, ...)
```

Arguments

x	A "ranger" object.
data	The training dataset.
...	Arguments to pass on to the "decorate.default" function.

Examples

```
library("ranger")  
library("r2pmml")  
  
data(iris)  
iris.ranger = ranger(Species ~ ., data = iris, num.trees = 17,  
                      write.forest = TRUE, probability = TRUE)  
iris.ranger = decorate(iris.ranger, data = iris)  
r2pmml(iris.ranger, file.path(tempdir(), "Iris-Ranger.pmml"))
```

decorate.svm.formula *Decorates a "svm.formula" object with an "xlevels" element.*

Description

Decorates a "svm.formula" object with an "xlevels" element.

Usage

```
## S3 method for class 'svm.formula'  
decorate(x, data, ...)
```

Arguments

x	A "svm.formula" object.
data	The training dataset.
...	Arguments to pass on to the "decorate.default" function.

decorate.train *Decorates the final model of a "train" object with model type-dependent elements.*

Description

Decorates the final model of a "train" object with model type-dependent elements.

Usage

```
## S3 method for class 'train'
decorate(x, ...)
```

Arguments

- x A "train" object.
- ... Arguments to pass on to the "decorate.default" function.

decorate.xgb.Booster *Decorates an "xgb.Booster" object with "fmap", "schema", "ntreelimit" and "pmml_options" elements.*

Description

Decorates an "xgb.Booster" object with "fmap", "schema", "ntreelimit" and "pmml_options" elements.

Usage

```
## S3 method for class 'xgb.Booster'
decorate(x, fmap, response_name = NULL,
         response_levels = c(), missing = NULL, ntreelimit = NULL,
         compact = FALSE, ...)
```

Arguments

- x An "xgb.Booster" object.
- fmap An XGBoost feature map as a "data.frame" object.
- response_name The name of the target field.
- response_levels A list of category values for a categorical target field.
- missing The string representation of missing input field values.
- ntreelimit The number of decision trees (aka boosting rounds) to convert.
- compact A flag controlling if decision trees should be transformed from binary splits (FALSE) to multi-way splits (TRUE) representation.
- ... Arguments to pass on to the "decorate.default" function.

Examples

```

library("xgboost")
library("r2pmml")

data(iris)
iris_x = iris[, -ncol(iris)]
iris_y = iris[, ncol(iris)]
# Convert from factor to integer[0, num_class]
iris_y = (as.integer(iris_y) - 1)
iris.fmap = genFMap(iris_x)
iris.dmatrix = genDMatrix(iris_y, iris_x)
iris.xgboost = xgboost(data = iris.dmatrix,
                       objective = "multi:softprob", num_class = 3, nrounds = 11)
iris.xgboost = decorate(iris.xgboost, iris.fmap,
                       response_name = "Species", response_levels = c("setosa", "versicolor", "virginica"))
pmmlFile = file.path(tempdir(), "Iris-XGBoost.pmml")
r2pmml(iris.xgboost, pmmlFile, compact = FALSE)
compactPmmlFile = file.path(tempdir(), "Iris-XGBoost-compact.pmml")
r2pmml(iris.xgboost, compactPmmlFile, compact = TRUE)

```

genDMatrix

Generates an XGBoost "DMatrix" object based on label and feature data.

Description

Generates an XGBoost "DMatrix" object based on label and feature data.

Usage

```
genDMatrix(df_y, df_X, file = tempfile(pattern = "DMatrix", fileext =
".libsvm"))
```

Arguments

- | | |
|------|--|
| df_y | A vector with dependent variable values. |
| df_X | A "data.frame" object with independent variable values. |
| file | A filesystem path for storing the temporary LibSVM data format file. |

Value

An "xgb.DMatrix" object.

Examples

```

data(iris)
iris.DMatrix = genDMatrix(as.integer(iris[, 5]) - 1, iris[, 1:4])

```

genFMap	<i>Generates an XGBoost feature map based on feature data.</i>
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Description

Generates an XGBoost feature map based on feature data.

Usage

```
genFMap(df_X)
```

Arguments

df_X A "data.frame" object with independent variables.

Value

A "data.frame" object.

Examples

```
data(iris)
iris.fmap = genFMap(iris[, 1:4])
```

r2pmml	<i>Converts an R model object to PMML.</i>
--------	--

Description

Converts an R model object to PMML.

Usage

```
r2pmml(x, file, converter = NULL, converter_classpath = NULL,
       verbose = FALSE, ...)
```

Arguments

x	An R model object.
file	A filesystem path to the result file.
converter	The name of a custom JPMML-R converter class.
converter_classpath	A list of filesystem paths to library JAR files that provide and support the custom JPMML-R converter class.
verbose	A flag controlling the verbosity of the conversion process.
...	Arguments to be passed on to the "r2pmml::decorate" function.

Examples

```

library("mlbench")
library("randomForest")
library("r2pmml")

data(iris)
iris.rf = randomForest(Species ~ ., data = iris, ntree = 7)
# Convert "randomForest" object to R-style (deep binary splits) MiningModel
pmmlFile = file.path(tempdir(), "Iris-RandomForest.pmml")
r2pmml(iris.rf, pmmlFile)
# Convert "randomForest" object to PMML-style (shallow multi-way splits) MiningModel
compactPmmlFile = file.path(tempdir(), "Iris-RandomForest-compact.pmml")
r2pmml(iris.rf, compactPmmlFile, compact = TRUE)

data(BostonHousing)
housing.glm = glm(medv ~ ., data = BostonHousing, family = "gaussian")
# Convert "glm" object into GeneralRegressionModel
genRegPmmlFile = file.path(tempdir(), "Housing-GLM.pmml")
r2pmml(housing.glm, genRegPmmlFile)
# Convert "glm" object into RegressionModel
regPmmlFile = file.path(tempdir(), "Housing-LM.pmml")
r2pmml(housing.glm, regPmmlFile, converter = "org.jpmml.rexp.LMConverter")

```

verify

Dispatches execution to the most appropriate model verification function.

Description

Dispatches execution to the most appropriate model verification function.

Usage

```
verify(x, newdata, ...)
```

Arguments

- | | |
|----------------------|--|
| <code>x</code> | A model object. |
| <code>newdata</code> | The verification dataset. |
| <code>...</code> | Arguments to pass on to the selected function. |

verify.default *Enhances a model object with verification data.*

Description

Enhances a model object with verification data.

Usage

```
## Default S3 method:  
verify(x, newdata, ...)
```

Arguments

x	A model object.
newdata	The verification dataset.
...	Further arguments.

verify.glm *Enhances a "glm" object with verification data.*

Description

Enhances a "glm" object with verification data.

Usage

```
## S3 method for class 'glm'  
verify(x, newdata, precision = 1e-13,  
      zeroThreshold = 1e-13, ...)
```

Arguments

x	A "glm" object.
newdata	The verification dataset.
precision	Maximal relative error.
zeroThreshold	Maximal absolute error near the zero value.
...	Further arguments.

Examples

```
library("mlbench")
library("r2pmml")

data(BostonHousing)
housing.glm = glm(medv ~ ., data = BostonHousing, family = "gaussian")
housing.glm = verify(housing.glm, newdata = BostonHousing[sample(nrow(BostonHousing), 10), ])
r2pmml(housing.glm, file.path(tempdir(), "Housing-GLM-verified.pmml"))
```

verify.train

Enhances a "train" object with verification data.

Description

Enhances a "train" object with verification data.

Usage

```
## S3 method for class 'train'
verify(x, newdata, precision = 1e-13,
       zeroThreshold = 1e-13, ...)
```

Arguments

- x A "train" object.
- newdata The verification dataset.
- precision Maximal relative error.
- zeroThreshold Maximal absolute error near the zero value.
- ... Arguments to pass on to the "predict.train" method.

writeFMap

Writes XGBoost feature map to a file.

Description

Writes XGBoost feature map to a file.

Usage

```
writeFMap(fmap, file)
```

Arguments

- fmap An XGBoost feature map as a "data.frame" object.
- file A filesystem path to the result file.

Index

as.scorecard, [2](#)
decorate, [3](#)
decorate.default, [3](#)
decorate.earth, [4](#)
decorate.elmNN, [4](#)
decorate.glmnet, [5](#)
decorate.party, [5](#)
decorate.randomForest, [6](#)
decorate.ranger, [7](#)
decorate.svm.formula, [7](#)
decorate.train, [8](#)
decorate.xgb.Booster, [8](#)

genDMatrix, [9](#)
genFMap, [10](#)

r2pmml, [10](#)

verify, [11](#)
verify.default, [12](#)
verify.glm, [12](#)
verify.train, [13](#)

writeFMap, [13](#)