

# Package ‘ODRF’

December 18, 2022

**Type** Package

**Title** Oblique Decision Random Forest for Classification and Regression

**Version** 1.0-1

**Date** 2022-09-21

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**Description** The oblique decision tree (ODT), which uses linear combinations of predictors as partitioning variables. Oblique Decision Random Forest (ODRF) is the assembly of multiple ODTs, which can effectively reduce the overfitting of individual ODTs and improve the accuracy of classification and regression.

**BugReports** <https://github.com/liuyu-star/ODRF/issues>

**License** GPL (>= 2)

**LazyData** yes

**NeedsCompilation** yes

**Depends** R (>= 3.1.0), partykit

**Imports** Rcpp (>= 0.11.0), stats, doParallel, foreach, Pursuit, grid, nnet

**LinkingTo** Rcpp, RcppArmadillo

**RoxygenNote** 7.2.0

**Suggests** knitr

**VignetteBuilder** knitr

## R topics documented:

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as.party.ODT	<i>ODT as party</i>
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**Description**

Functions coercing ODT object to objects of class party.

**Usage**

```
## S3 method for class 'ODT'  
as.party(ppTree, data, ...)
```

**Arguments**

- ppTree            an object of class [ODT](#).
- data             Training data of class data.frame is used to convert the object of class ODRF. and data must be the training data data in [ODT](#).
- ...              arguments to be passed to methods

**References**

Lee, EK(2017) PPtreeViz: An R Package for Visualizing Projection Pursuit Classification Trees, Journal of Statistical Software <doi:10.18637/jss.v083.i08>

**See Also**

[ODT party](#)

## Examples

```
data(iris)
tree <- ODT(Species~.,data = iris)
tree
party.tree<- as.party(tree,data = iris)
party.tree
plot(party.tree)
```

---

best.cut.node	<i>Find best split variable and node.</i>
---------------	---

---

## Description

Three criterion functions for splitting variables.

## Usage

```
best.cut.node(
  X,
  y,
  type = "i-classification",
  weights = 1,
  MinLeaf = ifelse(type == "regression", 5, 1),
  numLabels = ifelse(type == "regression", 0, length(unique(y)))
)
```

## Arguments

X	An n by d numeric matrix (preferable) or data frame.
y	a n vector.
type	The criterion used for splitting the variable. 'i-classification': information gain (classification, default), 'g-classification': gini impurity index (classification) or 'regression': mean square error (regression).
weights	a vector of values which weigh the samples when considering a split.
MinLeaf	the minimum amount of samples in a leaf.
numLabels	the number of categories.

## Value

a list which contains:

- BestCutVar: the best split variable.
- BestCutVal: the best split point for the best split variable.
- BestIndex: Each variable corresponds to the min gini impurity index(method='g-classification'), the max information gain(method='i-classification') or the min squared error(method='regression').

### Examples

```
### Find the best split variable ###
data(iris)
X=as.matrix(iris[, 1:4])
y=iris[[5L]]
bestcut=best.cut.node(X,y,type='i-classification')
print(bestcut)
```

---

body\_fat

*Body Fat Prediction Dataset*

---

### Description

Lists estimates of the percentage of body fat determined by underwater weighing and various body circumference measurements for 252 men. Accurate measurement of body fat is inconvenient/costly and it is desirable to have easy methods of estimating body fat that are not inconvenient/costly.

### Format

A data frame with 252 rows and 15 covariate variables and 1 response variable

### Details

The variables listed below, from left to right, are:

- Density determined from underwater weighing
- Age (years)
- Weight (lbs)
- Height (inches)
- Neck circumference (cm)
- Chest circumference (cm)
- Abdomen 2 circumference (cm)
- Hip circumference (cm)
- Thigh circumference (cm)
- Knee circumference (cm)
- Ankle circumference (cm)
- Biceps (extended) circumference (cm)
- Forearm circumference (cm)
- Wrist circumference (cm)

### Source

<https://www.kaggle.com/datasets/fedesoriano/body-fat-prediction-dataset>

## References

- Bailey, Covert (1994). Smart Exercise: Burning Fat, Getting Fit, Houghton-Mifflin Co., Boston, pp. 179-186.
- Behnke, A.R. and Wilmore, J.H. (1974). Evaluation and Regulation of Body Build and Composition, Prentice-Hall, Englewood Cliffs, N.J.
- Siri, W.E. (1956), "Gross composition of the body", in Advances in Biological and Medical Physics, vol. IV, edited by J.H. Lawrence and C.A. Tobias, Academic Press, Inc., New York.
- Katch, Frank and McArdle, William (1977). Nutrition, Weight Control, and Exercise, Houghton Mifflin Co., Boston.
- Wilmore, Jack (1976). Athletic Training and Physical Fitness: Physiological Principles of the Conditioning Process, Allyn and Bacon, Inc., Boston.

## Examples

```
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])

rf = ODRF(Density~.,train_data,type='regression')
pred <- predict(rf,test_data[,-1],weight = FALSE)$prediction
#estimation error
mean((pred-test_data[,1])^2)

tree = ODT(Density~.,train_data,type='regression')
pred <- predict(tree,test_data[,-1])
#estimation error
mean((pred-test_data[,1])^2)
```

---

breast\_cancer

*Breast Cancer Dataset*


---

## Description

Breast cancer is the most common cancer amongst women in the world. It accounts for 25% of all cancer deaths. It starts when cells in the breast begin to grow out of control. These cells usually form tumors that can be seen via X-ray or felt as lumps in the breast area. The key challenges against its detection is how to classify tumors into malignant (cancerous) or benign (non cancerous).

## Format

A data frame with 569 rows and 30 covariate variables and 1 response variable

## Details

The actual linear program used to obtain the separating plane in the 3-dimensional space is that described in:

- ID number
- Diagnosis (M = malignant, B = benign)

- Ten real-valued features are computed for each cell nucleus:
  - radius (mean of distances from center to points on the perimeter)
  - texture (standard deviation of gray-scale values)
  - perimeter
  - area
  - smoothness (local variation in radius lengths)
  - compactness ( $\text{perimeter}^2 / \text{area} - 1.0$ )
  - concavity (severity of concave portions of the contour)
  - concave points (number of concave portions of the contour)
  - symmetry
  - fractal dimension ("coastline approximation" - 1)

### Source

<https://www.kaggle.com/datasets/yasserh/breast-cancer-dataset?select=breast-cancer.csv> and [https://archive.ics.uci.edu/ml/datasets/breast+cancer+wisconsin+\(diagnostic\)](https://archive.ics.uci.edu/ml/datasets/breast+cancer+wisconsin+(diagnostic))

### References

Wolberg WH, Street WN, Mangasarian OL. Machine learning techniques to diagnose breast cancer from image-processed nuclear features of fine needle aspirates. *Cancer Lett.* 1994 Mar 15;77(2-3):163-71.

O.L. Mangasarian, W.N. Street and W.H. Wolberg. Breast cancer diagnosis and prognosis via linear programming. *Operations Research*, 43(4), pages 570-577, July-August 1995.

### Examples

```
data(breast_cancer)

set.seed(221212)
train = sample(1:569,200)
train_data = data.frame(breast_cancer[train,-1])
test_data = data.frame(breast_cancer[-train,-1])

rf = ODRF(diagnosis~.,train_data,type='i-classification')
pred <- predict(rf,test_data[,-1],weight = FALSE)$prediction
#estimation error
(mean(pred!=test_data[,1]))

tree = ODT(diagnosis~.,train_data,type='i-classification')
pred <- predict(tree,test_data[,-1])
#estimation error
(mean(pred!=test_data[,1]))
```

---

ODRF

*Classification and Regression with Oblique Decision Random Forest*

---

### Description

ODRF is the assembly of multiple ODTs, which can effectively reduce the overfitting of individual ODTs and improve the accuracy of classification and regression.

**Usage**

```

ODRF(
  formula,
  data = NULL,
  type = NULL,
  NodeRotateFun = "RotMatPPO",
  FunDir = getwd(),
  paramList = NULL,
  ntrees = 100,
  storeOOB = TRUE,
  replacement = TRUE,
  stratify = TRUE,
  numOOB = 1/3,
  parallel = TRUE,
  numCores = Inf,
  seed = 220924,
  MaxDepth = Inf,
  numNode = Inf,
  MinLeaf = 5,
  subset = NULL,
  weights = NULL,
  na.action = na.fail,
  catLabel = NULL,
  Xcat = 0,
  Xscale = "Min-max",
  TreeRandRotate = FALSE,
  ...
)

```

**Arguments**

formula	Object of class formula with a response but no interaction terms describing the model to fit. If this is a data frame, it is taken as the model frame. (see <a href="#">model.frame</a> )
data	Training data of class data.frame in which to interpret the variables named in the formula. If data is missing it is obtained from the current environment by formula.
type	The criterion used for splitting the nodes. 'g-classification': gini impurity index(default) and 'i-classification': information gain for classification; 'regression': mean square error for regression. y is a factor then is a classification.
NodeRotateFun	Name of the function of class character that implements a linear combination of predictor variables in the split node. Default is "RotMatPPO" with model="PPR". (see <a href="#">RotMatPPO</a> ) Users can define this function, for details see <a href="#">RotMatMake</a> .
FunDir	The path to the function of the user-defined NodeRotateFun. (default current Workspace)
paramList	Parameters in a named list to be used by NodeRotateFun. If left unchanged, default values will be populated, for details see <a href="#">defaults</a> .
ntrees	The number of trees in the forest. (default 100)
storeOOB	if TRUE then the samples omitted during the creation of a tree are stored as part of the tree.

replacement	if TRUE then n samples are chosen, with replacement, from training data. (default TRUE)
stratify	if TRUE then class sample proportions are maintained during the random sampling. Ignored if replacement = FALSE. (default TRUE)
numOOB	Ratio of 'out-of-bag'.
parallel	Parallel computing or not. (default TRUE)
numCores	Number of cores to be used for parallel computing. (default Inf)
seed	Random seeds in order to reproduce results.
MaxDepth	The maximum depth of the tree (default Inf).
numNode	The number of nodes used by the tree (default Inf).
MinLeaf	Minimal node size. Default 1 for classification, 5 for regression.
subset	An index vector indicating which rows should be used. (NOTE: If given, this argument must be named.)
weights	A vector of length same as data that are positive weights.(default NULL)
na.action	A function to specify the action to be taken if NAs are found. (NOTE: If given, this argument must be named.)
catLabel	A category labels of class list in prediction variables. (for details see Details and Examples)
Xcat	A class vector is used to indicate which variables are class variables. The default Xcat=0 means that no special treatment is given to category variables. When Xcat=NULL, the variable x that satisfies the condition $(\text{length}(\text{unique}(x)) < 10) \& (n > 20)$ is judged to be a category variable # ' @param Xscale Predictor variable standardization methods." Min-max", "Quantile", "No" denote Min-max transformation, Quantile transformation and No transformation (default "Min-max"), respectively.
TreeRandRotate	If or not to randomly rotate the Training data before building the tree. (default FALSE)
...	optional parameters to be passed to the low level function.

## Value

An object of class ODT, which is a list with the following components:

- call: The original call to ODT.
- terms: An object of class `c("terms", "formula")` (see [terms.object](#)) summarizing the formula. Used by various methods, but typically not of direct relevance to users.
- ppTrees: Each tree used to build the forest.
  - oobErr: 'out-of-bag' error for tree, classification error rate for classification or mean square error for regression.
  - oobIndex: Which training data to use as 'out-of-bag'?
  - oobPred: Predicted value for 'out-of-bag'.
  - other: For other tree-related values see [ODT](#).
- oobErr: 'out-of-bag' error for forest, classification error rate for classification or mean square error for regression.
- oobConfusionMat: 'out-of-bag' confusion matrix for forest.
- type, Levels and NodeRotateFun are important parameters for building the tree.



- paramList: Parameters in a named list to be used by NodeRotateFun.
- data: The list of data related parameters used to build the tree.
- tree: The list of tree related parameters used to build the tree.
- forest: The list of forest related parameters used to build the tree.

### Author(s)

YU Liu and Yingcun Xia

### References

- Zhan H, Liu Y, Xia Y. Consistency of The Oblique Decision Tree and Its Random Forest[J]. arXiv preprint arXiv:2211.12653, 2022.
- Tomita T M, Browne J, Shen C, et al. Sparse projection oblique randomer forests[J]. Journal of machine learning research, 2020, 21(104).

### See Also

[predict.ODRF](#) [print.ODRF](#) [ODRF.error](#) [VarImp](#)

### Examples

```
#Classification with Oblique Decision Random Forest
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODRF(varieties_of_wheat~,train_data,type='i-classification')
pred <- predict(tree,test_data[, -8])
#estimation error
(mean(pred!=test_data[, 8]))

#Regression with Oblique Decision Random Forest
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])

tree = ODRF(Density~,train_data,type='regression')
pred <- predict(tree,test_data[, -1])
#estimation error
mean((pred-test_data[, 1])^2)

### Train ODRF on one-of-K encoded categorical data ###
Xcol1=sample(c("A","B","C"),100,replace = TRUE)
Xcol2=sample(c("1","2","3","4","5"),100,replace = TRUE)
Xcon=matrix(rnorm(100*3),100,3)
X=data.frame(Xcol1,Xcol2,Xcon)
Xcat=c(1,2)
catLabel=NULL
y=as.factor(sample(c(0,1),100,replace = TRUE))
```

```

tree = ODRF(y~X,type='g-classification')

numCat <- apply(X[,Xcat,drop = FALSE], 2, function(x) length(unique(x)))
X1 <- matrix(0, nrow = nrow(X), ncol = sum(numCat)) # initialize training data matrix X
catLabel <- vector("list", length(Xcat))
names(catLabel)<- colnames(X)[Xcat]
col.idx <- 0L
# one-of-K encode each categorical feature and store in X
for (j in 1:length(Xcat)) {
  catMap <- (col.idx + 1L):(col.idx + numCat[j])
  # convert categorical feature to K dummy variables
  catLabel[[j]]=levels(as.factor(X[,Xcat[j]]))
  X1[, catMap] <- (matrix(X[,Xcat[j]],nrow(X),numCat[j])==matrix(catLabel[[j]],nrow(X),
    numCat[j],byrow = TRUE))+0
  col.idx <- col.idx + numCat[j]
}
X=cbind(X1,X[,~Xcat])

#Print the result after processing of category variables
X
#   1 2 3 4 5 6 7 8          X1          X2          X3
#1 0 1 0 0 1 0 0 0 -0.81003483  0.7900958 -1.94504333
#2 0 0 1 0 0 0 0 1 -0.02528851 -0.5143964 -0.18628226
#3 1 0 0 1 0 0 0 0  1.15532067  2.0236020  1.02942500
#4 1 0 0 0 0 1 0 0  1.18598589  1.0594630  0.42990019
#5 1 0 0 1 0 0 0 0 -0.21695438  1.5145973  0.09316665
#6 0 0 1 0 0 0 0 1 -1.11507717 -0.5775602  0.09918911
catLabel
# $Xcol1
# [1] "A" "B" "C"
# $Xcol2
# [1] "1" "2" "3" "4" "5"

```

---

ODRF.error

*oblique decision random forest error*


---

## Description

the error of class ODRF at different number of trees.

## Usage

```
ODRF.error(ppForest, data, newdata = NULL, ...)
```

## Arguments

ppForest	an object of class ODRF, as that created by the function <a href="#">ODRF</a> .
data	Training data of class data.frame in which to interpret the variables named in the formula.If data is missing it is obtained from the current environment by formula.
newdata	A data frame or matrix containing new data.

**Value**

OOB error and test error, classification error rate for classification or  $RPE(MSE/\text{mean}((y_{\text{test}} - \text{mean}(y))^2))$  for regression.

**See Also**

[ODRF plot.ODRF.error](#)

**Examples**

```
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

forest = ODRF(varieties_of_wheat~.,train_data,type='i-classification')
error=ODRF.error(forest,train_data,test_data)
```

---

ODT

---

*Classification and Regression with Oblique Decision Tree*


---

**Description**

ODT uses a linear combination of predictors as partitioning variables to create classification and regression tree.

**Usage**

```
ODT(
  formula,
  data = NULL,
  type = NULL,
  NodeRotateFun = "RotMatPPO",
  FunDir = getwd(),
  paramList = NULL,
  MaxDepth = Inf,
  numNode = Inf,
  MinLeaf = 5,
  Levels = NULL,
  subset = NULL,
  weights = NULL,
  na.action = na.fail,
  catLabel = NULL,
  Xcat = 0,
  Xscale = "Min-max",
  TreeRandRotate = FALSE,
  ...
)
```

**Arguments**

formula	Object of class formula with a response but no interaction terms describing the model to fit. If this is a data frame, it is taken as the model frame. (see <a href="#">model.frame</a> )
data	Training data of class data.frame in which to interpret the variables named in the formula. If data is missing it is obtained from the current environment by formula.
type	The criterion used for splitting the nodes. 'g-classification': gini impurity index(default) and 'i-classification': information gain for classification; 'regression': mean square error for regression. y is a factor then is a classification.
NodeRotateFun	Name of the function of class character that implements a linear combination of predictor variables in the split node. Default is "RotMatPPO" with model="PPR". (see <a href="#">RotMatPPO</a> ) Users can define this function, for details see <a href="#">RotMatMake</a> .
FunDir	The path to the function of the user-defined NodeRotateFun. (default current Workspace)
paramList	Parameters in a named list to be used by NodeRotateFun. If left unchanged, default values will be populated, for details see <a href="#">defaults</a> .
MaxDepth	The maximum depth of the tree (default Inf).
numNode	The number of nodes used by the tree (default Inf).
MinLeaf	Minimal node size. Default 1 for classification, 5 for regression.
Levels	The category label of the response variable when type is not equal to 'regression'.
subset	An index vector indicating which rows should be used. (NOTE: If given, this argument must be named.)
weights	A vector of length same as data that are positive weights.(default NULL)
na.action	A function to specify the action to be taken if NAs are found. (NOTE: If given, this argument must be named.)
catLabel	A category labels of class list in prediction variables. (for details see Examples)
Xcat	A class vector is used to indicate which variables are class variables. The default Xcat=0 means that no special treatment is given to category variables. When Xcat=NULL, the variable x that satisfies the condition (length(unique(x))<10) & (n>20) is judged to be a category variable
Xscale	Predictor variable standardization methods. "Min-max", "Quantile", "No" denote Min-max transformation, Quantile transformation and No transformation, respectively. (default "Min-max")
TreeRandRotate	If or not to randomly rotate the Training data before building the tree. (default FALSE)
...	optional parameters to be passed to the low level function.

**Value**

An object of class ODT, which is a list with the following components:

- call: The original call to ODT.

- **terms**: An object of class `c("terms", "formula")` (see [terms.object](#)) summarizing the formula. Used by various methods, but typically not of direct relevance to users.
  - **structure**: A set of tree structure data records.
    - nodeRotaMat**: Record the split variables (first column), split node serial number (second column) and rotation direction (third column) for each node. (The first column and the third column are 0 means leaf nodes)
  - **nodeNumLabel**: Record each leaf node's category for classification or predicted value for regression (second column is data size). (Each column is 0 means it is not a leaf node)
  - **nodeCutValue**: Record the split point of each node. (0 means leaf nodes)
  - **nodeCutIndex**: Record the index values of the partitioning variables selected based on the partition criterion type.
  - **childNode**: Record the number of child nodes after each splitting.
  - **nodeDepth**: Record the depth of the tree where each node is located.
- **type**, **Levels** and **NodeRotateFun** are important parameters for building the tree.
- **paramList**: Parameters in a named list to be used by **NodeRotateFun**.
- **data**: The list of data related parameters used to build the tree.
- **tree**: The list of tree related parameters used to build the tree.

### Author(s)

YU Liu and Yingcun Xia

### References

Zhan H, Liu Y, Xia Y. Consistency of The Oblique Decision Tree and Its Random Forest[J]. arXiv preprint arXiv:2211.12653, 2022.

### See Also

[predict.ODT](#) [print.ODT](#) [plot.ODT](#) [plot\\_ODT\\_depth](#)

### Examples

```
#Classification with Oblique Decision Tree
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODT(varieties_of_wheat~,train_data,type='i-classification')
pred <- predict(tree,test_data[,8])
#estimation error
(mean(pred!=test_data[,8]))

#Regression with Oblique Decision Tree
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])
```

```

tree = ODT(Density~.,train_data,type='regression')
pred <- predict(tree,test_data[,-1])
#estimation error
mean((pred-test_data[,1])^2)

### Train ODT on one-of-K encoded categorical data ###
Xcol1=sample(c("A","B","C"),100,replace = TRUE)
Xcol2=sample(c("1","2","3","4","5"),100,replace = TRUE)
Xcon=matrix(rnorm(100*3),100,3)
X=data.frame(Xcol1,Xcol2,Xcon)
Xcat=c(1,2)
catLabel=NULL
y=as.factor(sample(c(0,1),100,replace = TRUE))
tree = ODT(y~X,type='g-classification')

numCat <- apply(X[,Xcat,drop = FALSE], 2, function(x) length(unique(x)))
X1 <- matrix(0, nrow = nrow(X), ncol = sum(numCat)) # initialize training data matrix X
catLabel <- vector("list", length(Xcat))
names(catLabel)<- colnames(X)[Xcat]
col.idx <- 0L
# one-of-K encode each categorical feature and store in X
for (j in 1:length(Xcat)) {
  catMap <- (col.idx + 1L):(col.idx + numCat[j])
  # convert categorical feature to K dummy variables
  catLabel[[j]]=levels(as.factor(X[,Xcat[j]]))
  X1[, catMap] <- (matrix(X[,Xcat[j]],nrow(X),numCat[j])==matrix(catLabel[[j]],nrow(X),
    numCat[j],byrow = TRUE))+0
  col.idx <- col.idx + numCat[j]
}
X=cbind(X1,X[,Xcat])

#Print the result after processing of category variables
X
#   1 2 3 4 5 6 7 8          X1          X2          X3
#1 0 1 0 0 1 0 0 0 -0.81003483  0.7900958 -1.94504333
#2 0 0 1 0 0 0 0 1 -0.02528851 -0.5143964 -0.18628226
#3 1 0 0 1 0 0 0 0  1.15532067  2.0236020  1.02942500
#4 1 0 0 0 0 1 0 0  1.18598589  1.0594630  0.42990019
#5 1 0 0 1 0 0 0 0 -0.21695438  1.5145973  0.09316665
#6 0 0 1 0 0 0 0 1 -1.11507717 -0.5775602  0.09918911
catLabel
#Xcol1
#[1] "A" "B" "C"
#Xcol2
#[1] "1" "2" "3" "4" "5"

```

---

online

---

*A machine learning algorithm for training class ODT and ODRF.*


---

## Description

The [ODT](#) and [ODRF](#) are constantly updated by multiple batches of data to optimize the model. [online](#) is a S3 method for class `ODT` and `ODRF`.

**Usage**

```
online(obj, ...)
```

**Arguments**

`obj` an object of class ODT or ODRF.  
`...` For other parameters related to class `obj`, see ODT or ODRF.

**Value**

object of class ODT or ODRF.

**See Also**

[ODT](#) [ODRF](#) [online.ODT](#) [online.ODRF](#)

---

online.ODRF	<i>A machine learning algorithm for training class ODRF.</i>
-------------	--

---

**Description**

The [ODRF](#) is constantly updated by multiple batches of data to optimize the model.

**Usage**

```
## S3 method for class 'ODRF'
online(ppForest, data, weights = NULL)
```

**Arguments**

`data` Training data of class `data.frame` is used to update the object of class ODT.  
`weights` A vector of length same as `data` that are positive weights. (default NULL)  
`obj` An object of class ODRF.

**Value**

The same result as ODRF.

**See Also**

[ODRF](#) [prune.ODRF](#)

## Examples

```
#Classification with Oblique Decision Random Forest
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODRF(varieties_of_wheat~,train_data[seq(floor(nrow(train_data)/2)),],type='i-classification')
tree = online(tree,train_data[-seq(floor(nrow(train_data)/2)),])
pred <- predict(tree,test_data[, -8])
#estimation error
(mean(pred!=test_data[,8]))

#Regression with Oblique Decision Random Forest
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])

tree = ODRF(Density~,train_data[seq(floor(nrow(train_data)/2)),],type='regression')
tree = online(tree,train_data[-seq(floor(nrow(train_data)/2)),])
pred <- predict(tree,test_data[, -8])
#estimation error
mean((pred-test_data[,1])^2)
```

---

online.ODT

*A machine learning algorithm for training class ODT.*


---

## Description

The [ODT](#) is constantly updated by multiple batches of data to optimize the model.

## Usage

```
## S3 method for class 'ODT'
online(ppTree, data, weights = NULL)
```

## Arguments

data	Training data of class <code>data.frame</code> is used to update the object of class <code>ODT</code> .
weights	A vector of length same as data that are positive weights. (default <code>NULL</code> )
obj	an object of class <code>ODT</code> .

## Value

The same result as `ODT`.

## See Also

[ODT prune.ODT](#)



## Examples

```
#Classification with Oblique Decision Tree
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODT(varieties_of_wheat~,train_data[seq(floor(nrow(train_data)/2)),],
type='i-classification')
tree = online(tree,train_data[-seq(floor(nrow(train_data)/2)),,])
pred <- predict(tree,test_data[, -8])
#estimation error
(mean(pred!=test_data[,8]))

#Regression with Oblique Decision Tree
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])

tree = ODT(Density~,train_data[seq(floor(nrow(train_data)/2)),],
type='regression')
tree = online(tree,train_data[-seq(floor(nrow(train_data)/2)),,])
pred <- predict(tree,test_data[, -8])
#estimation error
mean((pred-test_data[,1])^2)
```

---

plot.ODRF.error

*Plot method for ODRF objects*


---

## Description

Draw the error graph of class ODRF at different number of trees.

## Usage

```
## S3 method for class 'ODRF.error'
plot(
  Err,
  lty = 1,
  main = paste0("Oblique ", ifelse(Err$type == "regression", "Regression",
    "Classification"), " Forest"),
  ...
)
```

## Arguments

Err	Object of class <a href="#">ODRF.error</a> .
lty	a vector of line types, see <a href="#">par</a> .
main	main title of the plot.
...	arguments to be passed to methods.

**Value**

OOB error and test error, classification error rate for classification or  $RPE(MSE/\text{mean}((y_{\text{test}} - \text{mean}(y))^2))$  for regression.

**See Also**

[ODRF](#) [ODRF.error](#)

**Examples**

```
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

forest = ODRF(varieties_of_wheat~,train_data,type='i-classification')
error=ODRF.error(forest,train_data,test_data)
plot(error)
```

---

plot.ODT

*oblique decision tree plot*


---

**Description**

Draw oblique decision tree with tree structure. It is modified from a function in PPtreeViz library.

**Usage**

```
## S3 method for class 'ODT'
plot(
  ppTree,
  font.size = 17,
  width.size = 1,
  xadj = 0,
  main = paste0("Oblique ", ifelse(ppTree$type == "regression", "Regression",
    "Classification"), " Tree"),
  sub = NULL,
  ...
)
```

**Arguments**

ppTree	an object of class <a href="#">ODT</a> .
font.size	font size of plot
width.size	size of eclipse in each node.
xadj	The size of the left and right movement.
main	main title
sub	sub title
...	arguments to be passed to methods.

## References

Lee, EK(2017) PPtreeViz: An R Package for Visualizing Projection Pursuit Classification Trees, Journal of Statistical Software <doi:10.18637/jss.v083.i08>

## See Also

[ODT plot\\_ODT\\_depth](#)

## Examples

```
data(iris)
tree <- ODT(Species~., data = iris,type='i-classification')
plot(tree)
```

---

plot.prune.ODT	<i>prune oblique decision tree plot</i>
----------------	---

---

## Description

Draw the error graph of class ODT at different number of split nodes.

## Usage

```
## S3 method for class 'prune.ODT'
plot(
  ppTree,
  position = "topleft",
  main = paste0("Oblique ", ifelse(ppTree$type == "regression", "Regression",
    "Classification"), " Tree"),
  ...
)
```

## Arguments

ppTree	an object of class <a href="#">prune.ODT</a> .
position	Position of the curve label.
main	main title
...	arguments to be passed to methods.

## Value

Error of test data after each pruning, classification error rate for classification or  $RPE(MSE/\text{mean}((y_{\text{test}} - \text{mean}(y))^2))$  for regression.

## See Also

[ODT prune.ODT](#)

**Examples**

```

data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODT(varieties_of_wheat~,train_data,type='i-classification')
tree = prune(tree,train_data)
#oblique decision tree plot (default)
plot(tree)
#prune oblique decision tree plot
class(tree)="prune.ODT"
plot(tree)

```

plot.VarImp

*Variable Importance Plot***Description**

Dotchart of variable importance as measured by a Oblique Decision Random Forest.

**Usage**

```

## S3 method for class 'VarImp'
plot(
  varImp,
  nvar = min(30, nrow(varImp$varImp)),
  main = paste0("Oblique ", ifelse(varImp$type == "regression", "Regression",
    "Classification"), " Forest"),
  ...
)

```

**Arguments**

varImp	an object of class <a href="#">VarImp</a> .
nvar	How many variables to show?
main	plot title.
...	arguments to be passed to methods.

**Value**

A matrix of importance measure, first column for each predictor variable and second column is Increased error. classification error rate for classification or  $RPE(MSE/\text{mean}((y_{\text{test}} - \text{mean}(y))^2))$  for regression.

**See Also**

[ODRF VarImp](#)

**Examples**

```
data(breast_cancer)
forest = ODRF(diagnosis~.,seeds,type='i-classification')
varimp=VarImp(forest,seeds)
plot(varimp)
```

plot\_ODT\_depth

*oblique decision tree depth plot***Description**

Draw the error graph of class ODT at different depths.

**Usage**

```
plot_ODT_depth(
  formula,
  data = NULL,
  newdata = NULL,
  type = "i-classification",
  NodeRotateFun = "RotMatPPO",
  paramList = NULL,
  main = paste0("Oblique ", ifelse(type == "regression", "Regression",
    "Classification"), " Tree"),
  ...
)
```

**Arguments**

formula	Object of class formula with a response but no interaction terms describing the model to fit. If this is a data frame, it is taken as the model frame. (see <a href="#">model.frame</a> )
data	Training data of class data.frame in which to interpret the variables named in the formula.If data is missing it is obtained from the current environment by formula.
newdata	A data frame or matrix containing new data.
type	The criterion used for splitting the nodes. 'g-classification': gini impurity index(default) and 'i-classification': information gain for classification; 'regression': mean square error for regression.
NodeRotateFun	Name of the function of class character that implements a linear combination of predictor variables in the split node. Default is "RotMatPPO" with model="PPR". (see <a href="#">RotMatPPO</a> ) Users can define this function, for details see <a href="#">RotMatMake</a> .
paramList	Parameters in a named list to be used by ODT.
main	main title
...	arguments to be passed to methods.

Value

OOB error and test error, classification error rate for classification or  $RPE(MSE/mean((y_{test}-mean(y))^2))$  for regression.

See Also

[ODT plot.ODT](#)

Examples

```
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])
plot_ODT_depth(Density~.,train_data,test_data,type='regression')
```

---

PPO	<i>Projection Pursuit Optimization</i>
-----	--

---

Description

Find the optimal projection using various projectin pursuit indices with class information.

Usage

```
PPO(X, y, model = "PPR", type = "i-classification", weights = NULL, ...)
```

Arguments

X	An n by d numeric matrix (preferable) or data frame.
y	a n vector.
model	<p>model for projection pursuit.</p> <p>"PPR"(default): projection projection regression from <a href="#">ppr</a>. When y is a category label, it is expanded to K binary features.</p> <ul style="list-style-type: none"><li>• "Log": logistic regression from <a href="#">nnet</a>.</li><li>• "Rand": The random projection generated from -1, 1. The following models can only be used for classification, i.e. the type must be 'i-classification' or 'g-classification'.</li><li>• "LDA", "PDA", "Lr", "GINI", and "ENTROPY" from library PPtreeViz.<ul style="list-style-type: none"><li>• The following models from library <a href="#">Pursuit</a>.<ul style="list-style-type: none"><li>"holes": Holes index</li><li>– "cm": Central Mass index</li><li>– "holes": Holes index</li><li>– "friedmantukey": Friedman Tukey index</li><li>– "legendre": Legendre index</li><li>– "laguerrefourier": Laguerre Fourier index</li><li>– "hermite": Hermite index,</li></ul></li></ul></li></ul>

	<ul style="list-style-type: none"> <li>– "naturalhermite": Natural Hermite index</li> <li>– "kurtosismax": Maximum kurtosis index,</li> <li>– "kurtosismin": Minimum kurtosis index,</li> <li>– "moment": Moment index</li> <li>– "mf": MF index</li> <li>– "chi": Chi-square index</li> </ul>
type	The criterion used for splitting the variable. 'g-classification': gini impurity index (classification, default), 'i-classification': information gain (classification) or 'regression': mean square error (regression).
weights	A vector of length same as data that are positive weights. (default NULL)

### Value

optimal projection direction.

### References

Friedman, J. H. and Stuetzle, W. (1981). Projection pursuit regression. *Journal of the American Statistical Association*, 76, 817–823. doi: 10.2307/2287576.

Ripley, B. D. (1996) *Pattern Recognition and Neural Networks*. Cambridge.

Lee, YD, Cook, D., Park JW, and Lee, EK(2013) PPTree: Projection Pursuit Classification Tree, *Electronic Journal of Statistics*, 7:1369-1386.

COOK, D., LEE, E. K., BUJA, A., WICKHAM, H.. Grand tours, projection pursuit guided tours and manual controls. In Chen, Chunhouh, Hardle, Wolfgang, Unwin, e Antony (Eds.), *Handbook of data Visualization*, Springer Handbooks of Computational Statistics, chapter III.2, p. 295-314. Springer, 2008.

### See Also

[RotMatMake](#) [RotMatRand](#) [RotMatRF](#) [RotMatMake](#)

### Examples

```
#classification
data(iris)
(PP <- PPO(iris[,1:4],iris[,5],model = "Log",type='i-classification'))
(PP <- PPO(iris[,1:4],iris[,5],model = "PPR",type='i-classification'))
(PP <- PPO(iris[,1:4],iris[,5],model = "LDA",type='i-classification'))

#regression
data(body_fat)
(PP <- PPO(body_fat[,2:15],body_fat[,1],model = "Log",type='regression'))
(PP <- PPO(body_fat[,2:15],body_fat[,1],model = "Rand",type='regression'))
(PP <- PPO(body_fat[,2:15],body_fat[,1],model = "PPR",type='regression'))
```

---

predict.ODRF	<i>predict method for ODRF objects</i>
--------------	--

---

## Description

Prediction a oblique decision random forest based on an input matrix or data frame using [ODRF](#) function.

## Usage

```
## S3 method for class 'ODRF'
predict(ppForest, Xnew, type = "response", weight.tree = FALSE)
```

## Arguments

ppForest	an object of class ODRF, as that created by the function ODRF.
Xnew	an n by d numeric matrix (preferable) or data frame. The rows correspond to observations and columns correspond to features.
type	one of response, prob or tree, indicating the type of output: predicted values, matrix of class probabilities or predicted value for each tree.
weight.tree	Whether to weight the tree, if TRUE then use the out-of-bag error of the tree as the weight. (default FALSE)

## Value

A set of vectors in the following list:

- response: the predicted values of the new data.
- prob: matrix of class probabilities (one column for each class and one row for each input). If ppForest\$type is regression, a vector of tree weights is returned.
- tree: it is a matrix where each column contains prediction by a tree in the forest.

## References

- Zhan H, Liu Y, Xia Y. Consistency of The Oblique Decision Tree and Its Random Forest[J]. arXiv preprint arXiv:2211.12653, 2022.

## See Also

[ODRF](#)

## Examples

```
#Classification with Oblique Decision Tree
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODRF(varieties_of_wheat~,train_data,type='i-classification')
```



```

pred <- predict(tree,test_data[,-8])
#estimation error
(mean(pred!=test_data[,8]))

#Regression with Oblique Decision Tree
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])

tree = ODRF(Density~.,train_data,type='regression')
pred <- predict(tree,test_data[,-1])
#estimation error
mean((pred-test_data[,1])^2)

```

predict.ODT

*predict method for ODT objects***Description**

Prediction a oblique decision tree based on an input matrix or data frame using [ODT](#) function.

**Usage**

```

## S3 method for class 'ODT'
predict(ppTree, Xnew, leafnode = FALSE)

```

**Arguments**

ppTree	an object of class ODT, as that created by the function ODT.
Xnew	an n by d numeric matrix (preferable) or data frame. The rows correspond to observations and columns correspond to features.
leafnode	If or not output the leaf node sequence number that Xnew is partitioned. (default FALSE)

**Value**

A set of vectors in the following list:

- prediction: the predicted response of the new data.
- leafnode: the leaf node sequence number that the new data is partitioned.

**References**

- Zhan H, Liu Y, Xia Y. Consistency of The Oblique Decision Tree and Its Random Forest[J]. arXiv preprint arXiv:2211.12653, 2022.

**See Also**

[ODT](#)

**Examples**

```
#Classification with Oblique Decision Tree
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODT(varieties_of_wheat~.,train_data,type='i-classification')
pred <- predict(tree,test_data[, -8])
#estimation error
(mean(pred!=test_data[,8]))

#Regression with Oblique Decision Tree
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])

tree = ODT(Density~.,train_data,type='regression')
pred <- predict(tree,test_data[, -1])
#estimation error
mean((pred-test_data[,1])^2)
```

print.ODRF

*Print ODRF***Description**

Print contents of ODRF object.

**Usage**

```
## S3 method for class 'ODRF'
print(ppForest, ...)
```

**Arguments**

ppForest	an object of class <a href="#">ODRF</a> .
...	arguments to be passed to methods

**See Also**

[ODRF](#)

**Examples**

```
data(iris)
forest <- ODRF(Species~.,data = iris)
forest
```

---

print.ODT	<i>Print ODT result</i>
-----------	-------------------------

---

**Description**

Print the oblique decision tree result

**Usage**

```
## S3 method for class 'ODT'
print(ppTree, projection = FALSE, cutvalue = FALSE, verbose = TRUE, ...)
```

**Arguments**

ppTree	an object of class <a href="#">ODT</a> .
projection	print projection coefficients in each node if TRUE
cutvalue	print cutoff values in each node if TRUE
verbose	print if TRUE, no output if FALSE
...	arguments to be passed to methods

**References**

Lee, EK(2017) PPtreeViz: An R Package for Visualizing Projection Pursuit Classification Trees, Journal of Statistical Software <doi:10.18637/jss.v083.i08>

**See Also**

[ODT](#)

**Examples**

```
data(iris)
tree <- ODT(Species~.,data = iris)
tree
print(tree,projection=TRUE,cutvalue=TRUE)
```

---

prune	<i>Pruning of class ODT or ODRF.</i>
-------	--------------------------------------

---

**Description**

Prune ODT or ODRF from bottom to top with validation data based on prediction error. prune is a S3 method for class class ODT and ODRF.

**Usage**

```
prune(obj, ...)
```

**Arguments**

obj an object of class ODT or ODRF.  
 ... For other parameters related to class obj, see [ODT](#) or [ODRF](#).

**Value**

an object of class ODT and prune.ODT.

**See Also**

ODT [ODRF](#) prune.ODT [prune.ODRF](#)

---

prune.ODRF	<i>Pruning of class ODRF.</i>
------------	-------------------------------

---

**Description**

Prune ODRF from bottom to top with validation data based on prediction error.

**Usage**

```
## S3 method for class 'ODRF'
prune(ppForest, data, MaxDepth = 1, useOOB = TRUE)
```

**Arguments**

data validation data of class data.frame is used to prune the object of class ODRF. Note that when useOOB=TRUE, data must be the training data data in [ODRF](#)  
 MaxDepth The maximum depth of the tree after pruning. (Default 1)  
 useOOB Whether to use OOB for pruning. (Default TRUE)  
 obj an object of class [ODRF](#).

**Value**

an object of class ODRF and prune.ODRF.

ppForest The same result as ODRF.

- pruneError Error of validation data or OOB () after each pruning in each tree, classification error rate for classification or mean square error for regression.

**See Also**

[ODRF](#) [prune.ODT](#)

## Examples

```
#Classification with Oblique Decision Random Forest
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODRF(varieties_of_wheat~,train_data[seq(floor(nrow(train_data)/2)),],
type='i-classification')
tree = prune(tree,train_data[-seq(floor(nrow(train_data)/2)),])
pred <- predict(tree,test_data[, -8])
#estimation error
(mean(pred!=test_data[,8]))

#Regression with Oblique Decision Random Forest
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])

tree = ODRF(Density~,train_data[seq(floor(nrow(train_data)/2)),],
type='regression')
tree = prune(tree,train_data[-seq(floor(nrow(train_data)/2)),])
pred <- predict(tree,test_data[, -8])
#estimation error
mean((pred-test_data[,1])^2)
```

prune.ODT

*Pruning of class ODT.*

## Description

Prune ODT from bottom to top with validation data based on prediction error.

## Usage

```
## S3 method for class 'ODT'
prune(ppTree, data, MaxDepth = 1)
```

## Arguments

data	validation data of class <code>data.frame</code> is used to prune the object of class <code>ODT</code> .
MaxDepth	The maximum depth of the tree after pruning. (Default 1)
obj	an object of class <code>ODT</code> .

## Value

an object of class `ODT` and `prune.ODT`.

ppTree The same result as `ODT`.

- `pruneError` Error of validation data after each pruning, classification error rate for classification or  $RPE(MSE/mean((y_{test}-mean(y))^2))$  for regression.

**See Also**

[ODT plot.prune.ODT](#)

**Examples**

```
#Classification with Oblique Decision Tree
data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

tree = ODT(varieties_of_wheat~,train_data[seq(floor(nrow(train_data)/2)),],type='i-classification')
tree = prune(tree,train_data[-seq(floor(nrow(train_data)/2)),])
pred <- predict(tree,test_data[, -8])
#estimation error
(mean(pred!=test_data[,8]))

#Regression with Oblique Decision Tree
data(body_fat)
set.seed(221212)
train = sample(1:252,100)
train_data = data.frame(body_fat[train,])
test_data = data.frame(body_fat[-train,])

tree = ODT(Density~,train_data[seq(floor(nrow(train_data)/2)),],type='regression')
tree = prune(tree,train_data[-seq(floor(nrow(train_data)/2)),])
pred <- predict(tree,test_data[, -8])
#estimation error
mean((pred-test_data[,1])^2)
```

---

RotMatMake

*Create rotation matrix used to determine linear combination of mtry features.*

---

**Description**

Create any projection matrix with a self-defined projection matrix function and projection optimization model function

**Usage**

```
RotMatMake(
  X = NULL,
  y = NULL,
  RotMatFun = "RotMatPPO",
  PPFun = "PPO",
  FunDir = getwd(),
  paramList = NULL
)
```

**Arguments**

<code>X</code>	An $n$ by $d$ numeric matrix (preferable) or data frame.
<code>y</code>	a $n$ vector.
<code>RotMatFun</code>	a self-defined projection matrix function name, which can also be <a href="#">RotMatRand</a> and <a href="#">RotMatPPO</a> . Note that <code>(,...)</code> is necessary.
<code>PPFun</code>	a self-defined projection matrix function, which can also be <a href="#">PPO</a> . Note that <code>(,...)</code> is necessary.
<code>FunDir</code>	The path to the function of the user-defined <code>NodeRotateFun</code> . (default current Workspace)
<code>paramList</code>	Parameters in a named list to be used. If left unchanged, default values will be populated, for details see <a href="#">defaults</a> .
<code>...</code>	used to handle superfluous arguments passed in using <code>paramList</code> .

**Value**

A random matrix to use in running [ODT](#).

Variable: variables to be projected.

- Number: Number of variables to be projected.
- Coefficient: Coefficients of the projection matrix.

**See Also**

[RotMatPPO](#) [RotMatRand](#) [RotMatRF](#)

**Examples**

```
X <- matrix(rnorm(200),20,10)
y <- (rnorm(20)>0)+0
set.seed(220828)
(RotMat <- RotMatMake(X,y,"RotMatRand","PPO"))
(RotMat <- RotMatMake(X,y,"RotMatPPO","PPO",paramList=list(model="Log")))
```

```
## Self-defined projection matrix function and projection optimization model function.##
## Note that (,...) is necessary.
makeRotMat=function(dimX,dimProj,numProj,...){
  RotMat=matrix(1,dimProj*numProj,3)
  for (np in seq(numProj)) {
    RotMat[(dimProj*(np-1)+1):(dimProj*np),1]=sample(1:dimX,dimProj,replace = FALSE)
    RotMat[(dimProj*(np-1)+1):(dimProj*np),2]=np
  }
  return(RotMat)
}
```

```
makePP=function(dimProj,prob,...){
  pp=sample(c(1L, -1L),dimProj, replace = TRUE,prob = c(prob, 1-prob))
  return(pp)
}
```

```
RotMat <- RotMatMake(RotMatFun="makeRotMat",PPFun="makePP",
  paramList=list(dimX=8,dimProj=5,numProj=4,prob=0.5))
head(RotMat)
```

#	Variable	Number	Coefficient
#[1,]	3	1	-1
#[2,]	6	1	1
#[3,]	4	1	-1
#[4,]	7	1	-1
#[5,]	8	1	1
#[6,]	6	2	-1

RotMatPPO

*Create a Projection Matrix: RotMatPPO***Description**

Create a projection matrix using projection pursuit optimization ([PPO](#)).

**Usage**

```
RotMatPPO(
  X,
  y,
  model = "PPR",
  type = "i-classification",
  weights = NULL,
  dimProj = min(ceiling(length(y)^0.4), ceiling(ncol(X) * 2/3)),
  numProj = ifelse(dimProj == "Rand", max(5, sample(floor(ncol(X)/3), 1)), max(5,
    ceiling(ncol(X)/dimProj))),
  catLabel = NULL,
  ...
)
```

**Arguments**

<code>X</code>	An $n$ by $d$ numeric matrix (preferable) or data frame.
<code>y</code>	a $n$ vector.
<code>model</code>	model for projection pursuit.
<code>type</code>	The criterion used for splitting the variable. 'g-classification': gini impurity index (classification, default), 'i-classification': information gain (classification) or 'regression': mean square error (regression).
<code>weights</code>	A vector of length same as data that are positive weights. (default NULL)
<code>dimProj</code>	Number of variables to be projected, <code>dimProj=min(ceiling(n^0.4),ceiling(ncol(X)*2/3))</code> (default) or <code>dimProj="Rand"</code> : random from 1 to <code>ncol(X)</code> .
<code>numProj</code>	the number of desired columns in the projection matrix, is also the number of projection directions. When <code>dimProj="Rand"</code> default <code>numProj=max(5, sample(floor(ncol(X)/3), 1))</code> , otherwise default <code>numProj=max(5, ceiling(p0/dimProj))</code> .
<code>catLabel</code>	A category labels of class list in prediction variables <sup>14</sup> For details see Examples of <a href="#">ODRF</a> .
<code>...</code>	used to handle superfluous arguments passed in using <code>paramList</code> .



**Value**

A random matrix to use in running [ODT](#).

Variable: variables to be projected.

- Number: Number of variables to be projected.
- Coefficient: Coefficients of the projection matrix.

**See Also**

[RotMatMake](#) [RotMatRand](#) [RotMatRF](#)

**Examples**

```
X <- matrix(rnorm(200),20,10)
y <- (rnorm(20)>0)+0
set.seed(220828)
(RotMat <- RotMatPPO(X,y))
(RotMat <- RotMatPPO(X,y,dimProj="Rand"))
(RotMat <- RotMatPPO(X,y,dimProj=10,numProj=5))
```

---

RotMatRand

*Random Rotation Matrix*


---

**Description**

Generate various rotation matrices by different distributions.

**Usage**

```
RotMatRand(
  dimX,
  randDist = "Binary",
  numProj = ceiling(sqrt(dimX)),
  dimProj = "Rand",
  sparsity = ifelse(dimX >= 10, 3/dimX, 1/dimX),
  prob = 0.5,
  lambda = 1,
  catLabel = NULL,
  ...
)
```

**Arguments**

dimX	the number of dimensions.
randDist	The probability distribution of the random projection direction from. "Binary", "Norm", "Uniform" denote the B-1,1 binomial distribution (default), N(0,1) normal distribution and U(-1,1) uniform distribution, respectively.
numProj	the number of desired columns in the projection matrix, is also the number of projection directions.(default ceiling(sqrt(dimX)))
dimProj	Number of variables to be projected, default dimProj="Rand": random from 1 to ncol(X).

sparsity	a real number in $(0, 1)$ that specifies the distribution of non-zero elements in the random matrix. When sparsity="pois" means that non-zero elements are generated by the $p(\lambda)$ Poisson distribution.
prob	a probability $\in (0, 1)$ used for sampling from. $-1, 1$ where prob = 0 will only sample -1 and prob = 1 will only sample 1.
lambda	Parameter of the Poisson distribution (default 1).
catLabel	A category labels of class list in prediction variables¼for details see Examples of <a href="#">ODRF</a> .
...	used to handle superfluous arguments passed in using paramList.

**Value**

A random matrix to use in running [ODT](#).

Variable: variables to be projected.

- Number: Number of variables to be projected.
- Coefficient: Coefficients of the projection matrix.

**See Also**

[RotMatPPO](#) [RotMatRF](#) [RotMatMake](#)

**Examples**

```
set.seed(1)
paramList <- list(dimX = 8, numProj= 3, sparsity=0.25, prob=0.5)
(RotMat <- do.call(RotMatRand, paramList))
paramList <- list(dimX = 8,numProj= 3,sparsity="pois")
(RotMat <- do.call(RotMatRand, paramList))
paramList <- list(dimX = 8, randDist="Norm", dimProj=5)
(RotMat <- do.call(RotMatRand, paramList))
```

---

RotMatRF

*Create a Projection Matrix: Random Forest (RF)*

---

**Description**

Create a projection matrix with coefficient 1, so that ODRF (ODT) has the same partition variables as random forest (tree).

**Usage**

```
RotMatRF(dimX, numProj, catLabel = NULL, ...)
```

**Arguments**

dimX	the number of dimensions.
numProj	the number of desired columns in the projection matrix, is also the number of projection directions.(default ceiling(sqrt(dimX)))
catLabel	A category labels of class list in prediction variables¼for details see Examples of <a href="#">ODRF</a> .
...	used to handle superfluous arguments passed in using paramList.

**Value**

A random matrix to use in running [ODT](#).

Variable: variables to be projected.

- Number: Number of variables to be projected.
- Coefficient: Coefficients of the projection matrix.

**See Also**

[RotMatPPO](#) [RotMatRand](#) [RotMatMake](#)

**Examples**

```
paramList <- list(dimX = 8, numProj= 3, catLabel=NULL)
set.seed(2)
(RotMat <- do.call(RotMatRF, paramList))
```

---

seeds

*seeds Data Set*


---

**Description**

Measurements of geometrical properties of kernels belonging to three different varieties of wheat. A soft X-ray technique and GRAINS package were used to construct all seven, real-valued attributes.

**Format**

A data frame with 209 rows and 7 covariate variables and 1 response variable.

**Details**

The variables listed below, from left to right, are:

- area A
- perimeter P
- compactness  $C = 4 \cdot \pi \cdot A / P^2$
- length of kernel
- width of kernel
- asymmetry coefficient
- length of kernel groove
- varieties of wheat (1, 2, 3 for Kama, Rosa and Canadian respectively)

**Source**

<https://archive.ics.uci.edu/ml/datasets/seeds>

**References**

M. Charytanowicz, J. Niewczas, P. Kulczycki, P.A. Kowalski, S. Lukasik, S. Zak, 'A Complete Gradient Clustering Algorithm for Features Analysis of X-ray Images', in: Information Technologies in Biomedicine, Ewa Pietka, Jacek Kawa (eds.), Springer-Verlag, Berlin-Heidelberg, 2010, pp. 15-24.

**Examples**

```

data(seeds)
set.seed(221212)
train = sample(1:209,100)
train_data = data.frame(seeds[train,])
test_data = data.frame(seeds[-train,])

library(ODRF)

rf = ODRF(varieties_of_wheat~,train_data,type='i-classification')
pred <- predict(rf,test_data[,-8],weight = FALSE)$prediction
#estimation error
(mean(pred!=test_data[,8]))

tree = ODT(varieties_of_wheat~,train_data,type='i-classification')
pred <- predict(tree,test_data[,-8])
#estimation error
(mean(pred!=test_data[,8]))

```

VarImp

*oblique decision random forest variable importance***Description**

variable importance measure is computed from permuting OOB data.

**Usage**

```
VarImp(ppForest, data)
```

**Arguments**

ppForest	an object of class <a href="#">ODRF</a> .
data	Training data of class <code>data.frame</code> in which to interpret the variables named in the formula.If data is missing it is obtained from the current environment by formula.

**Details**

A note from `randomForest`, here are the definitions of the variable importance measures. The first measure is computed from permuting OOB data: For each tree, the prediction error on the out-of-bag portion of the data is recorded (error rate for classification, RPE for regression). Then the same is done after permuting each predictor variable. The difference between the two are then averaged over all trees, and normalized by the standard deviation of the differences. If the standard deviation of the differences is equal to 0 for a variable, the division is not done (but the average is almost always equal to 0 in that case).

**Value**

A matrix of importance measure, first column for each predictor variable and second column is Increased error. classification error rate for classification or  $RPE(MSE/\text{mean}((y_{\text{test}}-\text{mean}(y))^2))$  for regression.

**See Also**[ODRF plot.VarImp](#)**Examples**

```
data(breast_cancer)
forest = ODRF(diagnosis~.,seeds,type='i-classification')
varimp=VarImp(forest,seeds)
```

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