

# Package ‘ocf’

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**Type** Package

**Title** Ordered Correlation Forest

**Version** 1.0.0

**Description** Nonparametric estimator for ordered non-numeric outcomes. The estimator modifies a standard random forest splitting criterion to build a collection of forests, each estimating the conditional probability of a single class. The package also implements a nonparametric estimator of the covariates’ marginal effects.

**License** GPL-3

**Encoding** UTF-8

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## R topics documented:

<code>marginal_effects</code> . . . . .	2
<code>mean_squared_error</code> . . . . .	4

multinomial_ml . . . . .	6
ocf . . . . .	8
ordered_ml . . . . .	10
predict.mml . . . . .	11
predict.ocf . . . . .	13
predict.oml . . . . .	14
print.ocf . . . . .	16
print.ocf.marginal . . . . .	17
summary.ocf . . . . .	18
summary.ocf.marginal . . . . .	19
tree_info . . . . .	20

<b>Index</b>	<b>22</b>
--------------	-----------

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marginal_effects	<i>Marginal Effects for Ordered Correlation Forest</i>
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## Description

Nonparametric estimation of marginal effects using an `ocf` object.

## Usage

```
marginal_effects(
  object,
  data = NULL,
  which_covariates = c(),
  eval = "atmean",
  bandwidth = 0.1,
  inference = FALSE
)
```

## Arguments

<code>object</code>	An <code>ocf</code> object.
<code>data</code>	Data set of class <code>data.frame</code> to estimate marginal effects. It must contain at least the same covariates used to train the forests. If <code>NULL</code> , marginal effects are estimated on <code>object\$full_data</code> .
<code>which_covariates</code>	Character vector storing the names of the covariates for which marginal effect estimation is desired. If empty (the default), marginal effects are estimated for all covariates.
<code>eval</code>	Evaluation point for marginal effects. Either <code>"mean"</code> , <code>"atmean"</code> or <code>"atmedian"</code> .
<code>bandwidth</code>	How many standard deviations <code>x_up</code> and <code>x_down</code> differ from <code>x</code> .
<code>inference</code>	Whether to extract weights and compute standard errors. The weights extraction considerably slows down the program.

## Details

`marginal_effects` can estimate mean marginal effects, marginal effects at the mean, or marginal effects at the median, according to the `eval` argument.

The routine assumes that covariates with more than ten unique values are continuous. Otherwise, covariates are assumed to be discrete.

## Value

Object of class `ocf.marginal`.

## Author(s)

Riccardo Di Francesco

## See Also

[ocf](#)

## Examples

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Fit ocf. Use large number of trees.
forests <- ocf(y, X, n.trees = 4000)

## Marginal effects at the mean.
me <- marginal_effects(forests, eval = "atmean")
print(me)
summary(me)

## LATEX.
print(me, latex = TRUE)

## Compute standard errors. This requires honest forests.
honest_forests <- ocf(y, X, n.trees = 4000, honesty = TRUE)
honest_me <- marginal_effects(honest_forests, eval = "atmean", inference = TRUE)
honest_me$standard.errors
honest_me$p.values # These are not corrected for multiple hypotheses testing!

print(honest_me, latex = TRUE)
```

---

mean\_squared\_error      *Accuracy Measures for Ordered Probability Predictions*

---

### Description

Accuracy measures for evaluating ordered probability predictions.

### Usage

```
mean_squared_error(y, predictions, use.true = FALSE)
```

```
mean_absolute_error(y, predictions, use.true = FALSE)
```

```
mean_ranked_score(y, predictions, use.true = FALSE)
```

```
classification_error(y, predictions)
```

### Arguments

y	Either the observed outcome vector or a matrix of true probabilities.
predictions	Predictions.
use.true	If TRUE, then the program treats y as a matrix of true probabilities.

### Details

#### MSE, MAE, and RPS:

When calling one of `mean_squared_error`, `mean_absolute_error`, or `mean_ranked_score`, `predictions` must be a matrix of predicted class probabilities, with as many rows as observations in `y` and as many columns as classes of `y`.

If `use.true == FALSE`, the mean squared error (MSE), the mean absolute error (MAE), and the mean ranked probability score (RPS) are computed as follows:

$$MSE = \frac{1}{n} \sum_{i=1}^n \sum_{m=1}^M (1(Y_i = m) - \hat{p}_m(x))^2$$

$$MAE = \frac{1}{n} \sum_{i=1}^n \sum_{m=1}^M |1(Y_i = m) - \hat{p}_m(x)|$$

$$RPS = \frac{1}{n} \sum_{i=1}^n \frac{1}{M-1} \sum_{m=1}^M (1(Y_i \leq m) - \hat{p}_m^*(x))^2$$

If `use.true == TRUE`, the MSE, the MAE, and the RPS are computed as follows (useful for simulation studies):

$$MSE = \frac{1}{n} \sum_{i=1}^n \sum_{m=1}^M (p_m(x) - \hat{p}_m(x))^2$$

$$MSE = \frac{1}{n} \sum_{i=1}^n \sum_{m=1}^M |p_m(x) - \hat{p}_m(x)|$$

$$RPS = \frac{1}{n} \sum_{i=1}^n \frac{1}{M-1} \sum_{m=1}^M (p_m^*(x) - \hat{p}_m^*(x))^2$$

where:

$$p_m(x) = P(Y_i = m | X_i = x)$$

$$p_m^*(x) = P(Y_i \leq m | X_i = x)$$

#### Classification error:

When calling [classification\\_error](#), predictions must be a vector of predicted class labels.

Classification error (CE) is computed as follows:

$$CE = \frac{1}{n} \sum_{i=1}^n 1(Y_i \neq \hat{Y}_i)$$

where  $Y_i$  are the observed class labels.

#### Value

The MSE, the MAE, the RPS, or the CE of the method.

#### Author(s)

Riccardo Di Francesco

#### See Also

[mean\\_ranked\\_score](#)

#### Examples

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])
```

```

## Training-test split.
train_idx <- sample(seq_len(length(y)), floor(length(y) * 0.5))

y_tr <- y[train_idx]
X_tr <- X[train_idx, ]

y_test <- y[-train_idx]
X_test <- X[-train_idx, ]

## Fit ocf on training sample.
forests <- ocf(y_tr, X_tr)

## Accuracy measures on test sample.
predictions <- predict(forests, X_test)

mean_squared_error(y_test, predictions$probabilities)
mean_ranked_score(y_test, predictions$probabilities)
classification_error(y_test, predictions$classification)

```

---

multinomial\_ml

*Multinomial Machine Learning*


---

## Description

Estimation strategy to estimate conditional choice probabilities for ordered non-numeric outcomes.

## Usage

```
multinomial_ml(y = NULL, X = NULL, learner = "forest", scale = TRUE)
```

## Arguments

y	Outcome vector.
X	Covariate matrix (no intercept).
learner	String, either "forest" or "l1". Selects the base learner to estimate each expectation.
scale	Logical, whether to scale the covariates. Ignored if learner is not "l1".

## Details

Multinomial machine learning expresses conditional choice probabilities as expectations of binary variables:

$$p_m(X_i) = \mathbb{E}[1(Y_i = m) | X_i]$$

This allows us to estimate each expectation separately using any regression algorithm to get an estimate of conditional probabilities.

`multinomial_ml` combines this strategy with either regression forests or penalized logistic regression with an L1 penalty, according to the user-specified parameter learner.

If `learner == "l1"`, the penalty parameters are chosen via 10-fold cross-validation and `model.matrix` is used to handle non-numeric covariates. Additionally, if `scale == TRUE`, the covariates are scaled to have zero mean and unit variance.

### Value

Object of class `mml`.

### Author(s)

Riccardo Di Francesco

### See Also

[ordered\\_ml](#), [ocf](#)

### Examples

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Training-test split.
train_idx <- sample(seq_len(length(y)), floor(length(y) * 0.5))

y_tr <- y[train_idx]
X_tr <- X[train_idx, ]

y_test <- y[-train_idx]
X_test <- X[-train_idx, ]

## Fit multinomial machine learning on training sample using two different learners.
multinomial_forest <- multinomial_ml(y_tr, X_tr, learner = "forest")
multinomial_l1 <- multinomial_ml(y_tr, X_tr, learner = "l1")

## Predict out of sample.
predictions_forest <- predict(multinomial_forest, X_test)
predictions_l1 <- predict(multinomial_l1, X_test)
```

```
## Compare predictions.
cbind(head(predictions_forest), head(predictions_l1))
```

---

ocf

*Ordered Correlation Forest*


---

## Description

Nonparametric estimator for ordered non-numeric outcomes. The estimator modifies a standard random forest splitting criterion to build a collection of forests, each estimating the conditional probability of a single class.

## Usage

```
ocf(
  y = NULL,
  X = NULL,
  honesty = FALSE,
  honesty.fraction = 0.5,
  inference = FALSE,
  alpha = 0,
  n.trees = 2000,
  mtry = ceiling(sqrt(ncol(X))),
  min.node.size = 5,
  max.depth = 0,
  replace = FALSE,
  sample.fraction = ifelse(replace, 1, 0.5),
  n.threads = 1
)
```

## Arguments

y	Outcome vector.
X	Covariate matrix (no intercept).
honesty	Whether to grow honest forests.
honesty.fraction	Fraction of honest sample. Ignored if honesty = FALSE.
inference	Whether to extract weights and compute standard errors. The weights extraction considerably slows down the routine. honesty = TRUE is required for valid inference.
alpha	Controls the balance of each split. Each split leaves at least a fraction alpha of observations in the parent node on each side of the split.
n.trees	Number of trees.



<code>mtry</code>	Number of covariates to possibly split at in each node. Default is the square root of the number of covariates.
<code>min.node.size</code>	Minimal node size.
<code>max.depth</code>	Maximal tree depth. A value of 0 corresponds to unlimited depth, 1 to "stumps" (one split per tree).
<code>replace</code>	If TRUE, grow trees on bootstrap subsamples. Otherwise, trees are grown on random subsamples drawn without replacement.
<code>sample.fraction</code>	Fraction of observations to sample.
<code>n.threads</code>	Number of threads. Zero corresponds to the number of CPUs available.

**Value**

Object of class `ocf`.

**Author(s)**

Riccardo Di Francesco

**See Also**

[marginal\\_effects](#)

**Examples**

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Training-test split.
train_idx <- sample(seq_len(length(y)), floor(length(y) * 0.5))

y_tr <- y[train_idx]
X_tr <- X[train_idx, ]

y_test <- y[-train_idx]
X_test <- X[-train_idx, ]

## Fit ocf on training sample.
forests <- ocf(y_tr, X_tr)

## We have compatibility with generic S3-methods.
print(forests)
summary(forests)
```

```

predictions <- predict(forests, X_test)
head(predictions$probabilities)
table(y_test, predictions$classification)

## Compute standard errors. This requires honest forests.
honest_forests <- ocf(y_tr, X_tr, honesty = TRUE, inference = TRUE)
head(honest_forests$predictions$standard.errors)

```

---

 ordered\_ml

*Ordered Machine Learning*


---

### Description

Estimation strategy to estimate conditional choice probabilities for ordered non-numeric outcomes.

### Usage

```
ordered_ml(y = NULL, X = NULL, learner = "forest", scale = TRUE)
```

### Arguments

y	Outcome vector.
X	Covariate matrix (no intercept).
learner	String, either "forest" or "l1". Selects the base learner to estimate each expectation.
scale	Logical, whether to scale the covariates. Ignored if learner is not "l1".

### Details

Ordered machine learning expresses conditional choice probabilities as the difference between the cumulative probabilities of two adjacent classes, which in turn can be expressed as conditional expectations of binary variables:

$$p_m(X_i) = \mathbb{E}[1(Y_i \leq m) | X_i] - \mathbb{E}[1(Y_i \leq m-1) | X_i]$$

Then we can separately estimate each expectation using any regression algorithm and pick the difference between the m-th and the (m-1)-th estimated surfaces to estimate conditional probabilities.

`ordered_ml` combines this strategy with either regression forests or penalized logistic regression with an L1 penalty, according to the user-specified parameter `learner`.

If `learner == "forest"`, then the `orf` function is called from an external package, as this estimator has already been proposed by Lechner and Okasa (2019).

If `learner == "l1"`, the penalty parameters are chosen via 10-fold cross-validation and `model.matrix` is used to handle non-numeric covariates. Additionally, if `scale == TRUE`, the covariates are scaled to have zero mean and unit variance.

**Value**

Object of class oml.

**Author(s)**

Riccardo Di Francesco

**See Also**

[multinomial\\_ml](#), [ocf](#)

**Examples**

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Training-test split.
train_idx <- sample(seq_len(length(y)), floor(length(y) * 0.5))

y_tr <- y[train_idx]
X_tr <- X[train_idx, ]

y_test <- y[-train_idx]
X_test <- X[-train_idx, ]

## Fit ordered machine learning on training sample using two different learners.
ordered_forest <- ordered_ml(y_tr, X_tr, learner = "forest")
ordered_l1 <- ordered_ml(y_tr, X_tr, learner = "l1")

## Predict out of sample.
predictions_forest <- predict(ordered_forest, X_test)
predictions_l1 <- predict(ordered_l1, X_test)

## Compare predictions.
cbind(head(predictions_forest), head(predictions_l1))
```

**Description**

Prediction method for class mml.

**Usage**

```
## S3 method for class 'mml'  
predict(object, data = NULL, ...)
```

**Arguments**

object	An mml object.
data	Data set of class data.frame. It must contain the same covariates used to train the base learners. If data is NULL, then object\$X is used.
...	Further arguments passed to or from other methods.

**Details**

If object\$learner == "l1", then `model.matrix` is used to handle non-numeric covariates. If we also have object\$scaling == TRUE, then data is scaled to have zero mean and unit variance.

**Value**

Matrix of predictions.

**Author(s)**

Riccardo Di Francesco

**See Also**

[multinomial\\_ml](#), [ordered\\_ml](#)

**Examples**

```
## Load data from orf package.  
set.seed(1986)  
  
library(orf)  
data(odata)  
odata <- odata[1:100, ] # Subset to reduce elapsed time.  
  
y <- as.numeric(odata[, 1])  
X <- as.matrix(odata[, -1])  
  
## Training-test split.  
train_idx <- sample(seq_len(length(y)), floor(length(y) * 0.5))  
  
y_tr <- y[train_idx]  
X_tr <- X[train_idx, ]
```

```

y_test <- y[-train_idx]
X_test <- X[-train_idx, ]

## Fit multinomial machine learning on training sample using two different learners.
multinomial_forest <- multinomial_ml(y_tr, X_tr, learner = "forest")
multinomial_l1 <- multinomial_ml(y_tr, X_tr, learner = "l1")

## Predict out of sample.
predictions_forest <- predict(multinomial_forest, X_test)
predictions_l1 <- predict(multinomial_l1, X_test)

## Compare predictions.
cbind(head(predictions_forest), head(predictions_l1))

```

---

predict.ocf

*Prediction Method for ocf Objects*

---

### Description

Prediction method for class `ocf`.

### Usage

```

## S3 method for class 'ocf'
predict(object, data = NULL, type = "response", ...)

```

### Arguments

<code>object</code>	An <code>ocf</code> object.
<code>data</code>	Data set of class <code>data.frame</code> . It must contain at least the same covariates used to train the forests. If <code>data</code> is <code>NULL</code> , then <code>object\$full_data</code> is used.
<code>type</code>	Type of prediction. Either <code>"response"</code> or <code>"terminalNodes"</code> .
<code>...</code>	Further arguments passed to or from other methods.

### Details

If `type == "response"`, the routine returns the predicted conditional class probabilities and the predicted class labels. If forests are honest, the predicted probabilities are honest.

If `type == "terminalNodes"`, the IDs of the terminal node in each tree for each observation in `data` are returned.

### Value

Desired predictions.

**Author(s)**

Riccardo Di Francesco

**See Also**

[ocf](#), [marginal\\_effects](#)

**Examples**

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Training-test split.
train_idx <- sample(seq_len(length(y)), floor(length(y) * 0.5))

y_tr <- y[train_idx]
X_tr <- X[train_idx, ]

y_test <- y[-train_idx]
X_test <- X[-train_idx, ]

## Fit ocf on training sample.
forests <- ocf(y_tr, X_tr)

## Predict on test sample.
predictions <- predict(forests, X_test)
head(predictions$probabilities)
predictions$classification

## Get terminal nodes.
predictions <- predict(forests, X_test, type = "terminalNodes")
predictions$forest.1[1:10, 1:20] # Rows are observations, columns are forests.
```

---

predict.oml

*Prediction Method for oml Objects*

---

**Description**

Prediction method for class oml.

**Usage**

```
## S3 method for class 'oml'  
predict(object, data = NULL, ...)
```

**Arguments**

object	An oml object.
data	Data set of class <code>data.frame</code> . It must contain the same covariates used to train the base learners. If data is <code>NULL</code> , then <code>object\$X</code> is used.
...	Further arguments passed to or from other methods.

**Details**

If `object$learner == "l1"`, then `model.matrix` is used to handle non-numeric covariates. If we also have `object$scaling == TRUE`, then data is scaled to have zero mean and unit variance.

**Value**

Matrix of predictions.

**Author(s)**

Riccardo Di Francesco

**See Also**

[multinomial\\_ml](#), [ordered\\_ml](#)

**Examples**

```
## Load data from orf package.  
set.seed(1986)  
  
library(orf)  
data(odata)  
odata <- odata[1:100, ] # Subset to reduce elapsed time.  
  
y <- as.numeric(odata[, 1])  
X <- as.matrix(odata[, -1])  
  
## Training-test split.  
train_idx <- sample(seq_len(length(y)), floor(length(y) * 0.5))  
  
y_tr <- y[train_idx]  
X_tr <- X[train_idx, ]  
  
y_test <- y[-train_idx]  
X_test <- X[-train_idx, ]
```

```
## Fit ordered machine learning on training sample using two different learners.
ordered_forest <- ordered_ml(y_tr, X_tr, learner = "forest")
ordered_l1 <- ordered_ml(y_tr, X_tr, learner = "l1")

## Predict out of sample.
predictions_forest <- predict(ordered_forest, X_test)
predictions_l1 <- predict(ordered_l1, X_test)

## Compare predictions.
cbind(head(predictions_forest), head(predictions_l1))
```

---

print.ocf

*Print Method for ocf Objects*

---

## Description

Prints an [ocf](#) object.

## Usage

```
## S3 method for class 'ocf'
print(x, ...)
```

## Arguments

`x` An [ocf](#) object.  
`...` Further arguments passed to or from other methods.

## Value

Prints an [ocf](#) object.

## Author(s)

Riccardo Di Francesco

## See Also

[ocf](#)

## Examples

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
```



```
odata <- odata[1:200, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Fit ocf.
forests <- ocf(y, X)

## Print.
print(forests)
```

---

print.ocf.marginal      *Print Method for ocf.marginal Objects*

---

## Description

Prints an `ocf.marginal` object.

## Usage

```
## S3 method for class 'ocf.marginal'
print(x, latex = FALSE, ...)
```

## Arguments

<code>x</code>	An <code>ocf.marginal</code> object.
<code>latex</code>	If TRUE, prints LATEX code.
<code>...</code>	Further arguments passed to or from other methods.

## Details

Compilation of the LATEX code requires the following packages: `booktabs`, `float`, `adjustbox`. If standard errors have been estimated, they are printed in parenthesis below each point estimate.

## Value

Prints an `ocf.marginal` object.

## Author(s)

Riccardo Di Francesco

## See Also

[ocf](#), [marginal\\_effects](#).

## Examples

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Fit ocf. Use large number of trees.
forests <- ocf(y, X, n.trees = 4000)

## Marginal effects at the mean.
me <- marginal_effects(forests, eval = "atmean")
print(me)
print(me, latex = TRUE)

## Add standard errors.
honest_forests <- ocf(y, X, n.trees = 4000, honesty = TRUE)
honest_me <- marginal_effects(honest_forests, eval = "atmean", inference = TRUE)
print(honest_me, latex = TRUE)
```

---

summary.ocf

*Summary Method for ocf Objects*

---

## Description

Summarizes an [ocf](#) object.

## Usage

```
## S3 method for class 'ocf'
summary(object, ...)
```

## Arguments

**object**            An [ocf](#) object.  
**...**             Further arguments passed to or from other methods.

## Value

Summarizes an [ocf](#) object.

## Author(s)

Riccardo Di Francesco

**See Also**[ocf, marginal\\_effects](#)**Examples**

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Fit ocf.
forests <- ocf(y, X)

## Summary.
summary(forests)
```

---

summary.ocf.marginal *Summary Method for ocf.marginal Objects*

---

**Description**

Summarizes an ocf.marginal object.

**Usage**

```
## S3 method for class 'ocf.marginal'
summary(object, latex = FALSE, ...)
```

**Arguments**

object	An ocf.marginal object.
latex	If TRUE, prints LATEX code.
...	Further arguments passed to or from other methods.

**Details**

Compilation of the LATEX code requires the following packages: booktabs, float, adjustbox. If standard errors have been estimated, they are printed in parenthesis below each point estimate.

**Value**

Summarizes an ocf.marginal object.

**Author(s)**

Riccardo Di Francesco

**See Also**

[ocf](#), [marginal\\_effects](#).

**Examples**

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:100, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Fit ocf. Use large number of trees.
forests <- ocf(y, X, n.trees = 4000)

## Marginal effects at the mean.
me <- marginal_effects(forests, eval = "atmean")
summary(me)
summary(me, latex = TRUE)

## Add standard errors.
honest_forests <- ocf(y, X, n.trees = 4000, honesty = TRUE)
honest_me <- marginal_effects(honest_forests, eval = "atmean", inference = TRUE)
summary(honest_me, latex = TRUE)
```

---

tree\_info

*Tree Information in Readable Format*

---

**Description**

Extracts tree information from a `ocf` forest object.

**Usage**

```
tree_info(object, tree = 1)
```

**Arguments**

<code>object</code>	ocf forest object.
<code>tree</code>	Number of the tree of interest.

**Details**

Nodes and variables IDs are 0-indexed, i.e., node 0 is the root node.

All values smaller than or equal to `splitval` go to the left and all values larger go to the right.

**Value**

A data.frame with the following columns:

<code>nodeID</code>	Node IDs.
<code>leftChild</code>	IDs of the left child node.
<code>rightChild</code>	IDs of the right child node.
<code>splitvarID</code>	IDs of the splitting variable.
<code>splitvarName</code>	Name of the splitting variable.
<code>splitval</code>	Splitting value.
<code>terminal</code>	Logical, TRUE for terminal nodes.
<code>prediction</code>	One column with the predicted conditional class probabilities.

**Author(s)**

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**See Also**

[ocf](#)

**Examples**

```
## Load data from orf package.
set.seed(1986)

library(orf)
data(odata)
odata <- odata[1:200, ] # Subset to reduce elapsed time.

y <- as.numeric(odata[, 1])
X <- as.matrix(odata[, -1])

## Fit ocf.
forests <- ocf(y, X)

## Extract information from tenth tree of first forest.
info <- tree_info(forests$forests.info$forest.1, tree = 10)
head(info)
```

# Index

classification\_error, [5](#)  
classification\_error  
    (mean\_squared\_error), [4](#)

marginal\_effects, [2](#), [3](#), [9](#), [14](#), [17](#), [19](#), [20](#)  
mean\_absolute\_error, [4](#)  
mean\_absolute\_error  
    (mean\_squared\_error), [4](#)  
mean\_ranked\_score, [4](#), [5](#)  
mean\_ranked\_score (mean\_squared\_error),  
    [4](#)  
mean\_squared\_error, [4](#), [4](#)  
model.matrix, [7](#), [10](#), [12](#), [15](#)  
multinomial\_m1, [6](#), [7](#), [11](#), [12](#), [15](#)

ocf, [2](#), [3](#), [7](#), [8](#), [11](#), [13](#), [14](#), [16–21](#)  
ordered\_m1, [7](#), [10](#), [10](#), [12](#), [15](#)  
orf, [10](#)

predict.mml, [11](#)  
predict.ocf, [13](#)  
predict.oml, [14](#)  
print.ocf, [16](#)  
print.ocf.marginal, [17](#)

summary.ocf, [18](#)  
summary.ocf.marginal, [19](#)

tree\_info, [20](#)