Package ‘C50’

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Description

Fit classification tree models or rule-based models using Quinlan’s C5.0 algorithm

Usage

```r
## Default S3 method:
C5.0(
  x,
  y,
  trials = 1,
  rules = FALSE,
  weights = NULL,
  control = C5.0Control(),
  costs = NULL,
  ...
)
```

```r
## S3 method for class 'formula'
C5.0(formula, data, weights, subset, na.action = na.pass, ...)
```

Arguments

- `x` a data frame or matrix of predictors.
- `y` a factor vector with 2 or more levels
- `trials` an integer specifying the number of boosting iterations. A value of one indicates that a single model is used.
- `rules` A logical: should the tree be decomposed into a rule-based model?
- `weights` an optional numeric vector of case weights. Note that the data used for the case weights will not be used as a splitting variable in the model (see http://www.rulequest.com/see5-win.html#CASEWEIGHT for Quinlan’s notes on case weights).
- `control` a list of control parameters; see `C5.0Control()`
costs

a matrix of costs associated with the possible errors. The matrix should have \( C \) columns and rows where \( C \) is the number of class levels.

... 

other options to pass into the function (not currently used with default method)

formula

a formula, with a response and at least one predictor.

data

an optional data frame in which to interpret the variables named in the formula.

subset

optional expression saying that only a subset of the rows of the data should be used in the fit.

na.action

a function which indicates what should happen when the data contain NA. The default is to include missing values since the model can accommodate them.

Details

This model extends the C4.5 classification algorithms described in Quinlan (1992). The details of the extensions are largely undocumented. The model can take the form of a full decision tree or a collection of rules (or boosted versions of either).

When using the formula method, factors and other classes are preserved (i.e. dummy variables are not automatically created). This particular model handles non-numeric data of some types (such as character, factor and ordered data).

The cost matrix should by \( C \times C \), where \( C \) is the number of classes. Diagonal elements are ignored. Columns should correspond to the true classes and rows are the predicted classes. For example, if \( C = 3 \) with classes Red, Blue and Green (in that order), a value of 5 in the (2,3) element of the matrix would indicate that the cost of predicting a Green sample as Blue is five times the usual value (of one). Note that when costs are used, class probabilities cannot be generated using \texttt{predict.C5.0()}.

Internally, the code will attempt to halt boosting if it appears to be ineffective. For this reason, the value of \texttt{trials} may be different from what the model actually produced. There is an option to turn this off in \texttt{C5.0Control()}. 

Value

An object of class \texttt{C5.0} with elements:

- \texttt{boostResults}: a parsed version of the boosting table(s) shown in the output
- \texttt{call}: the function call
- \texttt{caseWeights}: not currently supported.
- \texttt{control}: an echo of the specifications from \texttt{C5.0Control()}
- \texttt{cost}: the text version of the cost matrix (or "")
- \texttt{costMatrix}: an echo of the model argument
- \texttt{dims}: original dimensions of the predictor matrix or data frame
- \texttt{levels}: a character vector of factor levels for the outcome
- \texttt{names}: a string version of the names file
- \texttt{output}: a string version of the command line output
- \texttt{predictors}: a character vector of predictor names
- \texttt{rbm}: a logical for rules
rules: a character version of the rules file

size: an integer vector of the tree/rule size (or sizes in the case of boosting)

tree: a string version of the tree file

trials: a named vector with elements Requested (an echo of the function call) and Actual (how many the model used)

Note

The command line version currently supports more data types than the R port. Currently, numeric, factor and ordered factors are allowed as predictors.

Author(s)

Original GPL C code by Ross Quinlan, R code and modifications to C by Max Kuhn, Steve Weston and Nathan Coulter

References


See Also

C5.0Control(), summary.C5.0(), predict.C5.0(), C5imp()

Examples

```r
library(modeldata)
data(mlc_churn)

treeModel <- C5.0(x = mlc_churn[1:3333, -20], y = mlc_churn$churn[1:3333])
treeModel
summary(treeModel)

ruleModel <- C5.0(churn ~ ., data = mlc_churn[1:3333, ], rules = TRUE)
ruleModel
summary(ruleModel)
```
C5.0Control

Control for C5.0 Models

Description

Various parameters that control aspects of the C5.0 fit.

Usage

C5.0Control(
  subset = TRUE,
  bands = 0,
  winnow = FALSE,
  noGlobalPruning = FALSE,
  CF = 0.25,
  minCases = 2,
  fuzzyThreshold = FALSE,
  sample = 0,
  seed = sample.int(4096, size = 1) - 1L,
  earlyStopping = TRUE,
  label = "outcome"
)

Arguments

subset A logical: should the model evaluate groups of discrete predictors for splits? Note: the C5.0 command line version defaults this parameter to FALSE, meaning no attempted groupings will be evaluated during the tree growing stage.

bands An integer between 2 and 1000. If TRUE, the model orders the rules by their affect on the error rate and groups the rules into the specified number of bands. This modifies the output so that the effect on the error rate can be seen for the groups of rules within a band. If this options is selected and rules = FALSE, a warning is issued and rules is changed to TRUE.

winnow A logical: should predictor winnowing (i.e feature selection) be used?

noGlobalPruning A logical to toggle whether the final, global pruning step to simplify the tree.

CF A number in (0, 1) for the confidence factor.

minCases an integer for the smallest number of samples that must be put in at least two of the splits.

fuzzyThreshold A logical toggle to evaluate possible advanced splits of the data. See Quinlan (1993) for details and examples.

sample A value between (0, .999) that specifies the random proportion of the data should be used to train the model. By default, all the samples are used for model training. Samples not used for training are used to evaluate the accuracy of the model in the printed output.
An integer for the random number seed within the C code.

A logical to toggle whether the internal method for stopping boosting should be used.

A character label for the outcome used in the output. @return A list of options.

Author(s)

Original GPL C code by Ross Quinlan, R code and modifications to C by Max Kuhn, Steve Weston and Nathan Coulter

References


See Also

C5.0(), predict.C5.0(), summary.C5.0(), C5imp()

Examples

library(modeldata)
data(mlc_churn)

treeModel <- C5.0(x = mlc_churn[1:3333, -20],
                  y = mlc_churn$churn[1:3333],
                  control = C5.0Control(winnow = TRUE))

summary(treeModel)

C5imp

Variable Importance Measures for C5.0 Models

Description

This function calculates the variable importance (aka attribute usage) for C5.0 models.

Usage

C5imp(object, metric = "usage", pct = TRUE, ...)

Arguments

object an object of class C5.0
metric either 'usage' or 'splits' (see Details below)
pct a logical: should the importance values be converted to be between 0 and 100?
... other options (not currently used)
Details

By default, C5.0 measures predictor importance by determining the percentage of training set samples that fall into all the terminal nodes after the split (this is used when `metric = "usage"`). For example, the predictor in the first split automatically has an importance measurement of 100 percent. Other predictors may be used frequently in splits, but if the terminal nodes cover only a handful of training set samples, the importance scores may be close to zero. The same strategy is applied to rule-based models as well as the corresponding boosted versions of the model.

There is a difference in the attribute usage numbers between this output and the nominal command line output. Although the calculations are almost exactly the same (we do not add 1/2 to everything), the C code does not display that an attribute was used if the percentage of training samples covered by the corresponding splits is very low. Here, the threshold was lowered and the fractional usage is shown.

When `metric = "splits"`, the percentage of splits associated with each predictor is calculated.

Value

a data frame with a column `Overall` with the predictor usage values. The row names indicate the predictor.

Author(s)

Original GPL C code by Ross Quinlan, R code and modifications to C by Max Kuhn, Steve Weston and Nathan Coulter

References


See Also

`C5.0()`, `C5.0Control()`, `summary.C5.0()`, `predict.C5.0()`

Examples

```r
library(modeldata)
data(mlc_churn)
treeModel <- C5.0(x = mlc_churn[1:3333, -20], y = mlc_churn$churn[1:3333])
C5imp(treeModel)
C5imp(treeModel, metric = "splits")
```
plot.C5.0

Plot a decision tree

Description

Plot a decision tree.

Usage

```
## S3 method for class 'C5.0'
plot(x, trial = 0, subtree = NULL, ...)
```

Arguments

- `x`: an object of class C5.0
- `trial`: an integer for how many boosting iterations are used for prediction. NOTE: the internals of C5.0 are zero-based so to get the initial decision tree you must use `trial = 0`. If `trial` is set too large, it is reset to the largest value and a warning is given.
- `subtree`: an optional integer that can be used to isolate nodes below the specified split. See `partykit::party()` for more details.
- `...`: options passed to `partykit::plot.party()`

Value

No value is returned; a plot is rendered.

Author(s)

Mark Culp, Max Kuhn

References


See Also

`C5.0()`, `partykit::party()`

Examples

```r
mod1 <- C5.0(Species ~ ., data = iris)
plot(mod1)
plot(mod1, subtree = 3)
```
mod2 <- C5.0(Species ~ ., data = iris, trials = 10)
plot(mod2) ## should be the same as above

## plot first weighted tree
plot(mod2, trial = 1)

---

**predict.C5.0**

*Predict new samples using a C5.0 model*

**Description**

This function produces predicted classes or confidence values from a C5.0 model.

**Usage**

```r
## S3 method for class 'C5.0'
predict(
  object,  
  newdata = NULL,  
  trials = object$trials["Actual"],  
  type = "class",  
  na.action = na.pass,
  ...
)
```

**Arguments**

- `object` an object of class `C5.0`
- `newdata` a matrix or data frame of predictors
- `trials` an integer for how many boosting iterations are used for prediction. See the note below.
- `type` either "class" for the predicted class or "prob" for model confidence values.
- `na.action` when using a formula for the original model fit, how should missing values be handled?
- `...` other options (not currently used)

**Details**

Note that the number of trials in the object may be less than what was specified originally (unless `earlyStopping = FALSE` was used in `C5.0Control()`). If the number requested is larger than the actual number available, the maximum actual is used and a warning is issued.

Model confidence values reflect the distribution of the classes in terminal nodes or within rules.

For rule-based models (i.e. not boosted), the predicted confidence value is the confidence value from the most specific, active rule. Note that C4.5 sorts the rules, and uses the first active rule for prediction. However, the default in the original sources did not normalize the confidence values.
For example, for two classes it was possible to get confidence values of (0.3815, 0.8850) or (0.0000, 0.922), which do not add to one. For rules, this code divides the values by their sum. The previous values would be converted to (0.3012, 0.6988) and (0, 1). There are also cases where no rule is activated. Here, equal values are assigned to each class.

For boosting, the per-class confidence values are aggregated over all of the trees created during the boosting process and these aggregate values are normalized so that the overall per-class confidence values sum to one.

When the cost argument is used in the main function, class probabilities derived from the class distribution in the terminal nodes may not be consistent with the final predicted class. For this reason, requesting class probabilities from a model using unequal costs will throw an error.

**Value**

when type = “class”, a factor vector is returned. When type = “prob”, a matrix of confidence values is returned (one column per class).

**Author(s)**

Original GPL C code by Ross Quinlan, R code and modifications to C by Max Kuhn, Steve Weston and Nathan Coulter

**References**


**See Also**

C5.0(), C5.0Control(), summary.C5.0(), C5imp()

**Examples**

```r
library(modeldata)
data(mlc_churn)
treeModel <- C5.0(x = mlc_churn[1:3333, -20], y = mlc_churn$churn[1:3333])
predict(treeModel, mlc_churn[3334:3350, -20])
predict(treeModel, mlc_churn[3334:3350, -20], type = "prob")
```
Summary of C5.0 Models

**Description**

This function prints out detailed summaries for C5.0 models.

**Usage**

```r
# S3 method for class 'C5.0'
summary(object, ...)  
```

**Arguments**

- `object`: an object of class C5.0
- `...`: other options (not currently used)

**Details**

The output of this function mirrors the output of the C5.0 command line version.

The terminal nodes have text indicating the number of samples covered by the node and the number that were incorrectly classified. Note that, due to how the model handles missing values, the sample numbers may be fractional.

There is a difference in the attribute usage numbers between this output and the nominal command line output. Although the calculations are almost exactly the same (we do not add 1/2 to everything), the C code does not display that an attribute was used if the percentage of training samples covered by the corresponding splits is very low. Here, the threshold was lowered and the fractional usage is shown.

**Value**

A list with values

- `output`: a single text string with the model output
- `comp2`: the call to this function

**Author(s)**

Original GPL C code by Ross Quinlan, R code and modifications to C by Max Kuhn, Steve Weston and Nathan Coulter

**References**

See Also

C5.0(), C5.0Control(), summary.C5.0(), C5imp()

Examples

library(modeldata)
data(mlc_churn)

treeModel <- C5.0(x = mlc_churn[1:3333, -20], y = mlc_churn$churn[1:3333])
summary(treeModel)
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