Package 'outliertree'

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as.list.outlieroutputs

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as.list.outlieroutputs

Convert outlier outputs to R list

Description

Converts outliers results (an object as returned from predict.outliertree or from extract.training.outliers) to an R list which can be modified programatically.

Usage

```
## S3 method for class 'outlieroutputs' as.list(x, ...)
```

Arguments

- X Outliers flagged by an outlier.tree model, returned by e.g. the prediction function.
- ... Not used.

Value

The same outliers as a list of lists.

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check.outlierness.bounds

Check values that could potentially flag an observation as outlier

Description

Returns, for each numeric/date/timestamp column, a range of values *outside* of which observations could potentially be flagged as being an outlier in some cluster, and for categorical/ordinal/boolean columns, the factor levels that can be flagged as being an outlier in some cluster. If the lower bound is higher than the upper bound, it means any value can potentially be flagged as outlier.

Usage

```
check.outlierness.bounds(outlier_tree_model)
```

Arguments

outlier_tree_model

An Outlier Tree model object as generated by 'outlier.tree'.

Value

A list with column as the names and the bounds or categories as values.

```
extract.training.outliers
```

Extract outliers found in training data

Description

Extracts outliers from a model generated by 'outlier.tree' if it was passed parameter 'save_outliers' = 'TRUE'.

Usage

```
extract.training.outliers(outlier_tree_model)
```

Arguments

```
outlier_tree_model
```

An Outlier Tree object as returned by 'outlier.tree'.

Value

A data frame with the outliers, which can be pretty-printed by function 'print' from this same package.

hypothyroid

Data about thyroid hormones for anonymous patients

Description

This data contains several obvious outliers from misspellings in data entry. From Garavan Institute. For more details see link at the bottom.

Usage

```
data(hypothyroid)
```

Format

An object of class data. frame with 2772 rows and 23 columns.

Source

```
http://archive.ics.uci.edu/ml/datasets/thyroid+disease
```

outlier.tree

Outlier Tree

Description

Fit Outlier Tree model to normal data with perhaps some outliers.

Usage

```
outlier.tree(
  df,
 max_depth = 4L,
 min_gain = 0.01,
 z_norm = 2.67,
  z_outlier = 8,
  pct_outliers = 0.01,
 min_size_numeric = 25L,
 min_size_categ = 50L,
  categ_split = "binarize",
  categ_outliers = "tail",
  numeric_split = "raw",
  cols_ignore = NULL,
  follow_all = FALSE,
  gain_as_pct = TRUE,
  save_outliers = FALSE,
  outliers_print = 10L,
```

```
min_decimals = 2L,
 nthreads = parallel::detectCores()
)
```

Arguments

df

Data Frame with regular (i.e. non-outlier) data that might contain some outliers. See details for allowed column types.

max_depth

Maximum depth of the trees to grow. Can also pass zero, in which case it will only look for outliers with no conditions (i.e. takes each column as a 1-d distribution and looks for outliers in there independently of the values in other columns).

min_gain

Minimum gain that a split has to produce in order to consider it (both in terms of looking for outliers in each branch, and in considering whether to continue branching from them). Note that default value for GritBot is 1e-6, with 'gain as pct' = 'FALSE', but it's recommended to pass higher values (e.g. 1e-1) when using 'gain_as_pct' = 'FALSE'.

z_norm

Maximum Z-value (from standard normal distribution) that can be considered as a normal observation. Note that simply having values above this will not automatically flag observations as outliers, nor does it assume that columns follow normal distributions. Also used for categorical and ordinal columns for building approximate confidence intervals of proportions.

z_outlier

Minimum Z-value that can be considered as an outlier. There must be a large gap in the Z-value of the next observation in sorted order to consider it as outlier, given by (z_outlier - z_norm). Decreasing this parameter is likely to result in more observations being flagged as outliers. Ignored for categorical and ordinal columns.

pct_outliers

Approximate max percentage of outliers to expect in a given branch.

min_size_numeric

Minimum size that branches need to have when splitting a numeric column. In order to look for outliers in a given branch for a numeric column, it must have a minimum of twice this number of observations.

min_size_categ Minimum size that branches need to have when splitting a categorical or ordinal column. In order to look for outliers in a given branch for a categorical, ordinal, or boolean column, it must have a minimum of twice this number of observations.

categ_split

How to produce categorical-by-categorical splits. Options are:

- "binarize": Will binarize the target variable according to whether it's equal to each present category within it (greater/less for ordinal), and split each binarized variable separately.
- "bruteforce": Will evaluate each possible binary split of the categories (that is, it evaluates 2ⁿ potential splits every time). Note that trying this when there are many categories in a column will result in exponential computation time that might never finish.
- "separate": Will create one branch per category of the splitting variable (this is how GritBot handles them).

categ_outliers How to look for outliers in categorical variables. Options are:

• "tail": Will try to flag outliers if there is a large gap between proportions in sorted order, and this gap is unexpected given the prior probabilities. Such criteria tends to sometimes flag too many uninteresting outliers, but is able to detect more cases and recognize outliers when there is no single dominant category.

• "majority": Will calculate an equivalent to z-value according to the number of observations that do not belong to the non-majority class, according to formula ' $(n-n_maj)/(n * p_prior) < 1/z_outlier^2$ '. Such criteria tends to miss many interesting outliers and will only be able to flag outliers in large sample sizes. This is the approach used by GritBot.

numeric_split How to determine the split point in numeric variables. Options are:

- "mid": Will calculate the midpoint between the largest observation that goes to the '<=' branch and the smallest observation that goes to the '>' branch.
- "raw": Will set the split point as the value of the largest observation that goes to the '<=' branch.

This doesn't affect how outliers are determined in the training data passed in 'df', but it does affect the way in which they are presented and the way in which new outliers are detected when using 'predict'. "mid" is recommended for continuous-valued variables, while "raw" will provide more readable explanations for counts data at the expense of perhaps slightly worse generalizability to unseen data.

cols_ignore Vector containing columns which will not be split, but will be evaluated for usage in splitting other columns. Can pass either a logical (boolean) vector with the same number of columns as 'df', or a character vector of column names (must match with those of 'df'). Pass 'NULL' to use all columns.

> Whether to continue branching from each split that meets the size and gain criteria. This will produce exponentially many more branches, and if depth is large, might take forever to finish. Will also produce a lot more spurious outiers. Not recommended.

> Whether the minimum gain above should be taken in absolute terms, or as a percentage of the standard deviation (for numerical columns) or shannon entropy (for categorical columns). Taking it in absolute terms will prefer making more splits on columns that have a large variance, while taking it as a percentage might be more restrictive on them and might create deeper trees in some columns. For GritBot this parameter would always be 'FALSE'. Recommended to pass higher values for 'min_gain' when passing 'FALSE' here. Not that when 'gain as pct' = 'FALSE', the results will be sensitive to the scales of variables.

> Whether to store outliers detected in 'df' in the object that is returned. These outliers can then be extracted from the returned object through function 'extract.training.outliers'.

outliers_print Maximum number of flagged outliers in the training data to print after fitting the model. Pass zero or 'NULL' to avoid printing any. Outliers can be printed from resulting data frame afterwards through the 'predict' method, or through

follow_all

gain_as_pct

save_outliers

the 'print' method (on the extracted outliers, not on the model object) if passing 'save outliers' = 'TRUE'.

min_decimals Minimum number of decimals to use when printing numeric values for the

flagged outliers. The number of decimals will be dynamically increased according to the relative magnitudes of the values being reported. Ignored when

passing 'outliers_print=0' or 'outliers_print=FALSE'.

nthreads Number of parallel threads to use. When fitting the model, it will only use up to

one thread per column, while for prediction it will use up to one thread per row. The more threads that are used, the more memory will be required and allocated, so using more threads will not always lead to better speed. Can be changed after

the object is already initialized.

Details

Explainable outlier detection through decision-tree grouping. Tries to detect outliers by generating decision trees that attempt to "predict" the values of each column based on each other column, testing in each branch of every tried split (if it meets some minimum criteria) whether there are observations that seem too distant from the others in a 1-D distribution for the column that the split tries to "predict" (unlike other methods, this will not generate a score for each observation).

Splits are based on gain, while outlierness is based on confidence intervals. Similar in spirit to the GritBot software developed by RuleQuest research.

Supports columns of types numeric (either as type 'numeric' or 'integer'), categorical (either as type 'character' or 'factor' with unordered levels), boolean (as type 'logical'), and ordinal (as type 'factor' with ordered levels as checked by 'is.ordered'). Can handle missing values in any of them. Can also pass dates/timestamps that will get converted to numeric but shown as dates/timestamps in the output. Offers option to set columns to be used only for generating conditions without looking at outliers in them.

Infinite values will be taken into consideration when the column is used to split another column (that is, +inf will go into the branch that is greater than something, -inf into the other branch), but when a column is the target of the split, they will be taken as missing - that is, it will not report infinite values as outliers.

Value

An object with the fitted model that can be used to detect more outliers in new data, and from which outliers in the training data can be extracted (when passing 'save_outliers' = 'TRUE').

References

- GritBot software: https://www.rulequest.com/gritbot-info.html
- Cortes, David. "Explainable outlier detection through decision tree conditioning." arXiv preprint arXiv:2001.00636 (2020).

See Also

predict.outliertree extract.training.outliers hypothyroid

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Examples

```
library(outliertree)
### example dataset with interesting outliers
data(hypothyroid)
### fit the model and get a print of outliers
model <- outlier.tree(hypothyroid,</pre>
  outliers_print=10,
  save_outliers=TRUE,
  nthreads=1)
### extract outlier info as R list
outliers <- extract.training.outliers(model)</pre>
summary(outliers)
### information for row 745 (list of lists)
outliers[[745]]
### outliers can be sliced too
outliers[700:1000]
### use custom row names
df.w.names <- hypothyroid</pre>
row.names(df.w.names) <- paste0("rownum", 1:nrow(hypothyroid))</pre>
outliers.w.names <- predict(model, df.w.names, return_outliers=TRUE, nthreads=1)</pre>
outliers.w.names[["rownum745"]]
```

predict.outliertree Pro

Predict method for Outlier Tree

Description

Predict method for Outlier Tree

Usage

```
## S3 method for class 'outliertree'
predict(
  object,
  newdata,
  outliers_print = 15L,
  min_decimals = 2L,
  return_outliers = TRUE,
  nthreads = object$nthreads,
  ...
)
```

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Arguments

object An Outlier Tree object as returned by 'outlier.tree'.

newdata A Data Frame in which to look for outliers according to the fitted model.

outliers_print How many outliers to print. Pass zero or 'NULL' to avoid printing them. Must

pass at least one of 'outliers_print' and 'return_outliers'.

min_decimals Minimum number of decimals to use when printing numeric values for the

flagged outliers. The number of decimals will be dynamically increased according to the relative magnitudes of the values being reported. Ignored when

passing 'outliers_print=0' or 'outliers_print=FALSE'.

return_outliers

Whether to return the outliers in an R object (otherwise will just print them).

nthreads Number of parallel threads to use. Parallelization is done by rows.

... Not used.

Details

Note that after loading a serialized object from 'outlier.tree' through 'readRDS' or 'load', it will only de-serialize the underlying C++ object upon running 'predict' or 'print', so the first run will be slower, while subsequent runs will be faster as the C++ object will already be in-memory.

Value

If passing 'return_outliers' = 'TRUE', will return a list of lists with the outliers and their information (each row is an entry in the first list, with the same names as the rows in the input data frame), which can be printed into a human-readable format after-the-fact through functions 'print' and 'summary' (they do the same thing). Otherwise, will not return anything, but will print the outliers if any are detected. Note that, while the object that is returned will display a short summary of only some observations when printing it in the console, it actually contains information for all rows, and can be subsetted to obtain information specific to one row.

See Also

outlier.tree print.outlieroutputs

Examples

```
library(outliertree)
### random data frame with an obvious outlier
nrows = 100
set.seed(1)
df = data.frame(
   numeric_col1 = c(rnorm(nrows - 1), 1e6),
   numeric_col2 = rgamma(nrows, 1),
   categ_col = sample(c('categA', 'categB', 'categC'),
        size = nrows, replace = TRUE)
   )
### test data frame with another obvious outlier
```

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```
nrows\_test = 50
df_test = data.frame(
   numeric_col1 = rnorm(nrows_test),
   numeric\_col2 = c(-1e6, rgamma(nrows\_test - 1, 1)),
   categ_col = sample(c('categA', 'categB', 'categC'),
        size = nrows_test, replace = TRUE)
)
### fit model on training data
outliers_model = outlier.tree(df, outliers_print=FALSE, nthreads=1)
### find the test outlier
test_outliers = predict(outliers_model, df_test,
    outliers_print = 1, return_outliers = TRUE,
   nthreads = 1)
### retrieve the outlier info (for row 1) as an R list
test_outliers[[1]]
### to turn it into a 6-column table:
# dt = t(data.table::as.data.table(test_outliers))
```

Description

Pretty-prints outliers as output by the 'predict' function from an Outlier Tree model (as generated by function 'outlier.tree'), or by 'extract.training.outliers'. Same as function 'summary'.

Usage

```
## S3 method for class 'outlieroutputs'
print(x, outliers_print = 15L, min_decimals = 2L, only_these_rows = NULL, ...)
```

Arguments

x Outliers as returned by predict method on an object from 'outlier.tree'.

outliers_print Maximum number of outliers to print.

outlier s_prime mamman number of outliers to prime.

Minimum number of decimals to use when printing numeric values for the flagged outliers. The number of decimals will be dynamically increased according to the relative magnitudes of the values being reported. Ignored when passing 'outliers_print=0' or 'outliers_print=FALSE'.

only_these_rows

min_decimals

Specific rows to print (either numbers if the row names in the original data frame were null, or the row names they had if non-null). Pass 'NULL' to print information about potentially all rows

.. Not used.

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Value

The same input 'x' that was passed (as 'invisible').

See Also

outlier.tree predict.outliertree

Examples

```
### Example re-printing results for selected rows
library(outliertree)
data("hypothyroid")
### Fit model
otree <- outlier.tree(hypothyroid,</pre>
 nthreads=1,
 outliers_print=0)
### Store predictions
pred <- predict(otree,</pre>
 hypothyroid,
 outliers_print=0,
 return_outliers=TRUE,
 nthreads=1)
### Print stored predictions
### Row 531 is an outlier, but 532 is not
print(pred, only\_these\_rows = c(531, 532))
```

print.outliertree

Print summary information from Outlier Tree model

Description

Displays general statistics from a fitted Outlier Tree model (same as 'summary'). For printing the outliers discovered, use function 'print' on the returned outliers (e.g. from 'predict'), not on the model object iself.

Usage

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

Arguments

x An Outlier Tree model as produced by function 'outlier.tree'.

... Not used.

Details

Note that after loading a serialized object from 'outlier.tree' through 'readRDS' or 'load', it will only de-serialize the underlying C++ object upon running 'predict' or 'print', so the first run will be slower, while subsequent runs will be faster as the C++ object will already be in-memory.

Value

The same input 'x' that was passed (as 'invisible').

summary.outlieroutputs

Print outliers in human-readable format

Description

Pretty-prints outliers as output by the 'predict' function from an Outlier Tree model (as generated by function 'outlier.tree'), or by 'extract.training.outliers'. Same as function 'print' (see the documentation of 'print' for more information about the parameters).

Usage

```
## S3 method for class 'outlieroutputs'
summary(object, outliers_print = 15, ...)
```

Arguments

```
object Outliers as returned by predict method on an object from 'outlier.tree'.

outliers_print Maximum number of outliers to print.

Not used.
```

Value

The same 'object' input, returned invisibly.

See Also

print.outlieroutputs

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summary.outliertree

Print summary information from Outlier Tree model

Description

Displays general statistics from a fitted Outlier Tree model (same as 'print'). For printing the outliers discovered, use function 'print' on the returned outliers (e.g. from 'predict'), not on the model object iself.

Usage

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

Arguments

object An Outlier Tree model as produced by function 'outlier.tree'.
... Not used.

Value

The same input 'object' that was passed (as 'invisible').

See Also

print.outliertree

titanic

Data about passengers of the RMS Titanic

Description

This dataset contains information about 1,309 of the passengers who boarded the RMS Titanic ship for its last trip in the year 1912 (spoiler: the ship doesn't reach its destination).

Usage

```
data(titanic)
```

Format

An object of class data. frame with 1309 rows and 14 columns.

Source

https://github.com/jbryer/CompStats/raw/master/Data/titanic3.csv

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```
variable.names.outliertree
```

Get Variable Names for OutlierTree Model

Description

Returns the variable names from the data to which an OutlierTree model was fitted.

Columns will be returned in the following order according to their types, regardless of the order that they had in theoriginal input data:

- 1. Numeric.
- 2. Date.
- 3. Timestamp (POSIXct, POSIXlt).
- 4. Categorical.
- 5. Boolean / logical.
- 6. Ordinal.

Usage

```
## S3 method for class 'outliertree'
variable.names(object, ...)
```

Arguments

object An OutlierTree model as returned by function outlier.tree.
... Not used.

Value

A character vector with the variable names.

[.outlieroutputs

Slice or sub-set outliers

Description

Select given rows from outlier results.

Usage

```
## S3 method for class 'outlieroutputs' x[i]
```

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Arguments

- x An object of class 'outlieroutputs', as returned by e.g. predict.outliertree.
- i Rows to select. Can pass numbers or strings. Works the same as when selecting elements from a list.

Value

An object of class 'outlieroutputs' containing only the selected rows.

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