

# Package ‘outliertree’

August 6, 2022

**Type** Package

**Title** Explainable Outlier Detection Through Decision Tree Conditioning

**Version** 1.8.1-1

**Author** David Cortes

**Maintainer** David Cortes <david.cortes.rivera@gmail.com>

**URL** <https://github.com/david-cortes/outliertree>

**BugReports** <https://github.com/david-cortes/outliertree/issues>

**Description** Outlier detection method that flags suspicious values within observations, contrasting them against the normal values in a user-readable format, potentially describing conditions within the data that make a given outlier more rare.

Full procedure is described in Cortes (2020) <[arXiv:2001.00636](https://arxiv.org/abs/2001.00636)>.

Loosely based on the 'GritBot' <<https://www.rulequest.com/gritbot-info.html>> software.

**License** GPL (>= 3)

**Imports** Rcpp (>= 1.0.1)

**Depends** R (>= 3.5.0)

**Suggests** knitr, rmarkdown, kableExtra, data.table

**LinkingTo** Rcpp, Rcereal

**LazyData** true

**VignetteBuilder** knitr

**RoxygenNote** 7.1.2

**NeedsCompilation** yes

**Repository** CRAN

**Date/Publication** 2022-08-06 20:00:02 UTC

## R topics documented:

<code>as.list.outlieroutputs</code> . . . . .	2
<code>check.outlierness.bounds</code> . . . . .	3
<code>extract.training.outliers</code> . . . . .	3

hypothyroid . . . . .	4
outlier.tree . . . . .	4
predict.outliertree . . . . .	8
print.outlieroutputs . . . . .	10
print.outliertree . . . . .	11
summary.outlieroutputs . . . . .	12
summary.outliertree . . . . .	12
titanic . . . . .	13
unpack.outlier.tree . . . . .	13
[.outlieroutputs . . . . .	14

<b>Index</b>	<b>16</b>
--------------	-----------

---

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 <a href="#">outlier.tree</a> model, returned by e.g. the prediction function.
...	Not used.

## Value

The same outliers as a list of lists.

---

`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

<code>df</code>	Data Frame with regular (i.e. non-outlier) data that might contain some outliers. See details for allowed column types.
<code>max_depth</code>	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).
<code>min_gain</code>	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 <code>'gain_as_pct' = 'FALSE'</code> , but it's recommended to pass higher values (e.g. 1e-1) when using <code>'gain_as_pct' = 'FALSE'</code> .
<code>z_norm</code>	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.
<code>z_outlier</code>	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.
<code>pct_outliers</code>	Approximate max percentage of outliers to expect in a given branch.
<code>min_size_numeric</code>	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.
<code>min_size_categ</code>	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.
<code>categ_split</code>	How to produce categorical-by-categorical splits. Options are: <ul style="list-style-type: none"> <li><code>"binarize"</code> : 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.</li> <li><code>"bruteforce"</code> : Will evaluate each possible binary split of the categories (that is, it evaluates <math>2^n</math> 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.</li> <li><code>"separate"</code> : Will create one branch per category of the splitting variable (this is how GritBot handles them).</li> </ul>

- `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.
- `follow_all` 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 outliers. Not recommended.
- `gain_as_pct` 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.
- `save_outliers` 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

	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 outlieriness 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](#) [unpack.outlier.tree](#)

## Examples

```
library(outliertree)

### example dataset with interesting outliers
data(hypothyroid)

### fit the model and get a print of outliers
model <- outlier.tree(hypothyroid,
  outliers_print=10,
  save_outliers=TRUE,
  nthreads=1)

### extract outlier info as R list
outliers <- extract.training.outliers(model)
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
row.names(df.w.names) <- paste0("rownum", 1:nrow(hypothyroid))
outliers.w.names <- predict(model, df.w.names, return_outliers=TRUE)
outliers.w.names[["rownum745"]]
```

---

predict.outliertree    *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,
  ...
)
```



**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).
...	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](#) [unpack.outlier.tree](#)

**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
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)

### 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))

```

---

```
print.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 ‘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.
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’.
only_these_rows	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.

## 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,
  nthreads=1,
  outliers_print=0)

### Store predictions
pred <- predict(otree,
  hypothyroid,
  outliers_print=0,
  return_outliers=TRUE)

### 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 itself.

**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.

**See Also**

[print.outlieroutputs](#)

---

```
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 itself.

**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>

---

unpack.outlier.tree

*Unpack Outlier Tree model after de-serializing*

---

**Description**

After persisting an outlier tree model object through ‘saveRDS’, ‘save’, or restarting a session, the underlying C++ objects that constitute the outlier tree model and which live only on the C++ heap memory are not saved along, thus not restored after loading a saved model through ‘readRDS’ or ‘load’.

The model object however keeps serialized versions of the C++ objects as raw bytes, from which the C++ objects can be reconstructed, and are done so automatically after calling ‘predict’, ‘print’, or ‘summary’ on the freshly-loaded object from ‘readRDS’ or ‘load’.

This function allows de-serializing the object bytes without invoking any extra side effects or computations, akin to XGBoost’s ‘xgb.Booster.complete’ or CatBoost’s ‘catboost.restore\_handle’.

**Usage**

```
unpack.outlier.tree(model)
```

**Arguments**

`model` An Outlier Tree object as returned by `'outlier.tree'`, which has been just loaded from a disk file through `'readRDS'`, `'load'`, or a session restart.

**Details**

If the model is going to be used in a production system, it's possible after de-serialization to delete the raw bytes in order to save memory (e.g. `'otree$obj_from_cpp$serialized_obj <- NULL'`). The memory will however not be freed automatically, as it's managed by R's garbage collector.

**Value**

No return value. Object is modified in-place.

**Examples**

```
### Warning: this example will generate a temporary .Rds
### file in your temp folder, and will then delete it
library(outliertree)
set.seed(1)
df <- as.data.frame(matrix(rnorm(1000), nrow = 250))
otree <- outlier.tree(df, outliers_print=0, nthreads=1)
temp_file <- file.path(tempdir(), "otree.Rds")
saveRDS(otree, temp_file)
otree2 <- readRDS(temp_file)
file.remove(temp_file)

cat("Pointer after loading model is this: \n")
print(otree2$obj_from_cpp$ptr_model)

### now unpack the raw bytes
unpack.outlier.tree(otree2)
cat("Pointer after unpacking is this: \n")
print(otree2$obj_from_cpp$ptr_model)
```

---

[.outlieroutputs      *Slice or sub-set outliers*

---

**Description**

Select given rows from outlier results.

**Usage**

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

**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.

# Index

## \* datasets

- hypothyroid, 4
- titanic, 13

[.outlieroutputs, 14

as.list.outlieroutputs, 2

check.outlierness.bounds, 3

extract.training.outliers, 2, 3, 7

hypothyroid, 4, 7

outlier.tree, 2, 4, 9, 11

predict.outliertree, 2, 7, 8, 11, 15

print.outlieroutputs, 9, 10, 12

print.outliertree, 11, 13

summary.outlieroutputs, 12

summary.outliertree, 12

titanic, 13

unpack.outlier.tree, 7, 9, 13