

Package ‘FFTrees’

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Type Package

Title Generate, Visualise, and Evaluate Fast-and-Frugal Decision Trees

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Description Create, visualize, and test fast-and-frugal decision trees (FFTs) using the algorithms and methods described in Phillips, Neth, Woike & Gaissmaier (2017). FFTs are simple and transparent decision trees for solving binary classification problems. FFTs can be preferable to more complex algorithms because they require very little information, are easy to communicate, and are robust against overfitting.

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Imports caret, rpart, randomForest, e1071, crayon, graphics, progress, scales, dplyr, knitr, magrittr, stringr, testthat, tibble, tidyselect

Suggests rmarkdown, spelling

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URL <https://CRAN.R-project.org/package=FFTrees>,
<https://github.com/ndphillips/FFTrees/>

BugReports <https://github.com/ndphillips/FFTrees/issues>

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add_stats	<i>Add decision statistics to data (containing counts of a 2x2 contingency table)</i>
-----------	---

Description

add_stats assumes the input of essential 2x2 frequency counts (as a data frame data with variable names "hi", "fa", "mi", and "cr") and uses them to compute various decision accuracy measures.

Usage

```
add_stats(
  data,
  sens.w = 0.5,
  cost.each = NULL,
  cost.outcomes = list(hi = 0, fa = 1, mi = 1, cr = 0)
)
```

Arguments

data	A data frame with (integer) values named "hi", "fa", "mi", and "cr".
sens.w	numeric. Sensitivity weight (for computing weighted accuracy, wacc).
cost.each	numeric. An optional fixed cost added to all outputs (e.g.; the cost of the cue).
cost.outcomes	list. A list of length 4 named "hi", "fa", "mi", "cr", and specifying the costs of a hit, false alarm, miss, and correct rejection, respectively. E.g.; cost.outcomes = listc("hi" = 0, "fa" = 10, "mi" = 20, "cr" = 0) means that a false alarm and miss cost 10 and 20 units, respectively, while correct decisions incur no costs.

Details

Providing numeric values for cost.each (as a vector) and cost.outcomes (as a named list) allows computing cost information for the counts of corresponding classification decisions.

Value

A data frame with variables of computed accuracy and cost measures (but dropping inputs).

blood	<i>Blood donation data</i>
-------	----------------------------

Description

Blood donation data

Usage

blood

Format

A data frame containing 748 rows and 5 columns.

recency Months since last donation

frequency Total number of donations

total Total blood donated in c.c.

time Months since first donation

donation.crit Did he/she donated blood in March 2007?

...

Source

<https://archive.ics.uci.edu/ml/datasets/Blood+Transfusion+Service+Center>

See Also

Other datasets: [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

breastcancer	<i>Physiological data of patients tested for breast cancer</i>
--------------	--

Description

Physiological data of patients tested for breast cancer

Usage

breastcancer

Format

A data frame containing 699 patients (rows) and 9 variables (columns).

thickness Clump Thickness

cellsize.unif Uniformity of Cell Size

cellshape.unif Uniformity of Cell Shape

adhesion Marginal Adhesion

epithelial Single Epithelial Cell Size

nuclei.bare Bare Nuclei

chromatin Bland Chromatin

nucleoli Normal Nucleoli

mitoses Mitoses

diagnosis Is cancer present? TRUE or FALSE

...

Source

[https://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+\(Original\)](https://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+(Original))

See Also

Other datasets: [blood](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

car

Car acceptability data

Description

A dataset on car evaluations based on basic features, derived from a simple hierarchical decision model.

Usage

car

Format

A data frame containing 1728 cars (rows) and 7 variables (columns).

buying.price Numeric

maint.price Factor

doors Factor

persons Numeric

luggage Numeric
safety Factor
acceptability Factor
 ...

Details

The (yet to be binarized) criterion variable is a car's acceptability rating.

Source

<http://archive.ics.uci.edu/ml/datasets/Car+Evaluation>

References

Bohanec, M., Rajkovic, V. (1990): Expert system for decision making. *Sistemica* 1 (1), pp. 145-157.

See Also

Other datasets: [blood](#), [breastcancer](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

classtable	<i>Compute classification statistics for binary prediction and criterion (e.g.; truth) vectors</i>
------------	--

Description

The primary confusion matrix is computed by [confusionMatrix](#) of the **caret** package.

Usage

```
classtable(
  prediction_v = NULL,
  criterion_v,
  sens.w = 0.5,
  cost.v = NULL,
  correction = 0.25,
  cost.outcomes = list(hi = 0, fa = 1, mi = 1, cr = 0),
  na_prediction_action = "ignore"
)
```

Arguments

prediction_v	logical. A logical vector of predictions.
criterion_v	logical. A logical vector of (TRUE) criterion values.
sens.w	numeric. Sensitivity weight parameter (from 0 to 1, for computing wacc). Default: sens.w = .50.
cost.v	list. An optional list of additional costs to be added to each case.
correction	numeric. Correction added to all counts for calculating dprime.
cost.outcomes	list. A list of length 4 with names 'hi', 'fa', 'mi', and 'cr' specifying the costs of a hit, false alarm, miss, and correct rejection, respectively. For instance, cost.outcomes = listc("hi" = 0, "fa" = 10, "mi" = 20, "cr" = 0) means that a false alarm and miss cost 10 and 20, respectively, while correct decisions have no cost.
na_prediction_action	not sure.

 comp.pred

A wrapper for competing classification algorithms.

Description

comp.pred provides a wrapper for many classification algorithms — such as CART (rpart::rpart), logistic regression (glm), support vector machines (svm::svm), and random forests (randomForest::randomForest).

Usage

```
comp.pred(
  formula,
  data.train,
  data.test = NULL,
  algorithm = NULL,
  model = NULL,
  new.factors = "exclude"
)
```

Arguments

formula	a formula
data.train	dataframe. A training dataset.
data.test	dataframe. A testing dataset.
algorithm	string. An algorithm in the set "lr" – logistic regression, "cart" – decision trees, "rlr" – regularized logistic regression, "svm" – support vector machines, "rf" – random forests
model	model. An optional existing model applied to test data
new.factors	string. What should be done if new factor values are discovered in the test set? "exclude" = exclude (i.e.; remove these cases), "base" = predict the base rate of the criterion.

contraceptive	<i>Contraceptive use data</i>
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Description

A subset of the 1987 National Indonesia Contraceptive Prevalence Survey.

Usage

contraceptive

Format

A data frame containing 1473 rows and 10 columns.

wife.age Numeric

wife.edu Factor

hus.ed Factor

children Numeric

wife.rel Numeric

wife.work Factor

hus.occ Factor

sol Factor

media Numeric

cont.crit numeric

...

Details

The samples describe married women who were either not pregnant or do not know if they were at the time of interview.

The problem consists in predicting a woman's current contraceptive method choice (here: binarized `cont.crit`) based on her demographic and socio-economic characteristics.

Source

<https://archive.ics.uci.edu/ml/datasets/Contraceptive+Method+Choice>

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

creditapproval	<i>Credit approval data</i>
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Description

Credit approval data

Usage

```
creditapproval
```

Format

A data frame containing 690 rows and 15 columns

Source

<https://archive.ics.uci.edu/ml/datasets/Credit+Approval>

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

factclean	<i>Clean factor variables in prediction data</i>
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Description

Clean factor variables in prediction data

Usage

```
factclean(data.train, data.test, show.warning = T)
```

Arguments

<code>data.train</code>	A training dataset
<code>data.test</code>	A testing dataset
<code>show.warning</code>	logical

fertility

Fertility data

Description

This dataset describes a sample of 100 volunteers providing a semen sample that was analyzed according to the WHO 2010 criteria.

Usage

fertility

Format

A data frame containing 100 rows and 10 columns.

Details

Sperm concentration are related to socio-demographic data, environmental factors, health status, and life habits.

The binary criterion variable is diagnosis: Normal (N) vs. altered (O).

Source

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

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

FFTrees

Main function to create and apply fast-and-frugal trees (FFTs)

Description

FFTrees is the workhorse function of the **FFTrees** package for creating fast-and-frugal trees (FFTs).

FFTs are decision algorithms for solving binary classification tasks, i.e., they predict the values of a binary criterion variable based on 1 or multiple predictor variables (cues).

Using FFTrees on data usually generates a range of FFTs and corresponding summary statistics (as an FFTrees object) that can then be printed, plotted, and examined further.

The criterion and predictor variables are specified in [formula](#) notation. Based on the settings of data and data.test, FFTs are trained on a (required) training dataset and tested on an (optional) test dataset.

Create fast-and-frugal trees (FFTs).

Usage

```

FFTrees(
  formula = NULL,
  data = NULL,
  data.test = NULL,
  algorithm = "ifan",
  max.levels = NULL,
  sens.w = 0.5,
  cost.outcomes = NULL,
  cost.cues = NULL,
  stopping.rule = "exemplars",
  stopping.par = 0.1,
  goal = NULL,
  goal.chase = NULL,
  goal.threshold = "bacc",
  numthresh.method = "o",
  numthresh.n = 10,
  decision.labels = c("False", "True"),
  main = NULL,
  train.p = 1,
  rounding = NULL,
  repeat.cues = TRUE,
  my.tree = NULL,
  tree.definitions = NULL,
  do.comp = TRUE,
  do.cart = TRUE,
  do.lr = TRUE,
  do.rf = TRUE,
  do.svm = TRUE,
  object = NULL,
  force = FALSE,
  quiet = FALSE,
  comp = NULL,
  rank.method = NULL,
  store.data = NULL,
  verbose = NULL
)

```

Arguments

<code>formula</code>	formula. A formula specifying a binary criterion variable (as logical) as a function of 1 or more predictor variables (cues).
<code>data</code>	dataframe. A training dataset.
<code>data.test</code>	dataframe. An optional testing dataset with the same structure as <code>data</code> .
<code>algorithm</code>	character. The algorithm used to create FFTs. Can be 'ifan', 'dfan'.
<code>max.levels</code>	integer. The maximum number of levels considered for the trees. Because all permutations of exit structures are considered, the larger <code>max.levels</code> is, the

	more trees will be created.
<code>sens.w</code>	numeric. A number from 0 to 1 indicating how to weight sensitivity relative to specificity. Only relevant when <code>goal = 'wacc'</code> .
<code>cost.outcomes</code>	A list of length 4 with names 'hi', 'fa', 'mi', and 'cr' specifying the costs of a hit, false alarm, miss, and correct rejection, respectively. E.g.; <code>cost.outcomes = listc("hi" = 0, "fa" = 10, "mi" = 20, "cr" = 0)</code> means that a false alarm and miss cost 10 and 20, respectively, while correct decisions have no costs.
<code>cost.cues</code>	A list containing costs for each cue. Each element should have a name corresponding to a column in data, and each entry should be a single (positive) number. Cues not present in <code>cost.cues</code> are assumed to have no costs (i.e., a value of 0).
<code>stopping.rule</code>	character. A string indicating the method to stop growing trees. "levels" means the tree grows until a certain level; "exemplars" means the tree grows until a certain number of unclassified exemplars remain; "statdelta" means the tree grows until the change in the criterion statistic is less than a specified level.
<code>stopping.par</code>	numeric. A number indicating the parameter for the stopping rule. For <code>stopping.rule "levels"</code> , this is the number of levels. For stopping rule "exemplars", this is the smallest percentage of exemplars allowed in the last level.
<code>goal</code>	character. A string indicating the statistic to maximize when selecting final trees: "acc" = overall accuracy, "bacc" = balanced accuracy, "wacc" = weighted accuracy.
<code>goal.chase</code>	character. A string indicating the statistic to maximize when constructing trees: "acc" = overall accuracy, "bacc" = balanced accuracy, "wacc" = weighted accuracy, "cost" = cue costs.
<code>goal.threshold</code>	character. A string indicating the statistic to maximize when calculating cue thresholds: "acc" = overall accuracy, "bacc" = balanced accuracy, "wacc" = weighted accuracy. Default: <code>goal.threshold = "bacc"</code> .
<code>numthresh.method</code>	character. How should thresholds for numeric cues be determined? "o" will optimize thresholds, while "m" will always use the median.
<code>numthresh.n</code>	integer. Number of numeric thresholds to try.
<code>decision.labels</code>	string. A vector of strings of length 2 indicating labels for negative and positive cases. E.g.; <code>decision.labels = c("Healthy", "Diseased")</code> .
<code>main</code>	string. An optional label for the dataset. Passed on to other functions, like <code>plot.FFTrees</code> , and <code>print.FFTrees</code> .
<code>train.p</code>	numeric. What percentage of the data to use for training when <code>data.test</code> is not specified? For example, <code>train.p = .5</code> will randomly split data into a 50% training set and a 50% test set. <code>train.p = 1</code> , the default, uses all data for training.
<code>rounding</code>	integer. An integer indicating digit rounding for non-integer numeric cue thresholds. The default is NULL which means no rounding. A value of 0 rounds all possible thresholds to the nearest integer, 1 rounds to the nearest .1 (etc.).

<code>repeat.cues</code>	logical. Can cues occur multiple times within a tree?
<code>my.tree</code>	string. A string representing a verbal description of an FFT, i.e., an FFT in words. For example, <code>my.tree = "If age > 20, predict TRUE. If sex = {m}, predict FALSE. Otherwise, predict TRUE."</code>
<code>tree.definitions</code>	dataframe. An optional hard-coded definition of trees (see details below). If specified, no new trees are created.
<code>do.comp</code> , <code>do.cart</code> , <code>do.lr</code> , <code>do.rf</code> , <code>do.svm</code>	logical. Should alternative algorithms be created for comparison? All TRUE by default. Options are: <code>cart</code> = regular (non-frugal) trees with rpart ; <code>lr</code> = logistic regression with glm ; <code>rf</code> = random forests with randomForest ; <code>svm</code> = support vector machines with e1071 . Specifying <code>comp = FALSE</code> sets all these arguments to FALSE.
<code>object</code>	FFTrees. An optional existing FFTrees object. When specified, no new trees are fitted and the existing trees are applied to <code>data</code> and <code>data.test</code> .
<code>force</code>	logical. Setting <code>force = TRUE</code> forces some parameters (like <code>goal</code>) to be as specified by the user even when the algorithm thinks those specifications don't make sense. Default is <code>force = FALSE</code> .
<code>quiet</code>	logical. Should progress reports be printed? Can be helpful for diagnosis when the function is running slowly. Default is <code>quiet = FALSE</code> (i.e., show progress).
<code>comp</code> , <code>rank.method</code> , <code>store.data</code> , <code>verbose</code>	Deprecated arguments (unused or replaced, to be retired in future releases).

Value

An FFTrees object with the following elements:

criterion_name The name of the binary criterion variable (as character).

cue_names The names of all potential predictor variables (cues) in the data (as character).

formula The [formula](#) specified when creating the FFTs.

trees A list of FFTs created, with further details contained in `n`, `best`, `definitions`, `inwords`, `stats`, `level_stats`, and `decisions`.

data The original training and test data (if available).

params A list of defined control parameters (e.g.; `algorithm`, `goal`).

competition Models and classification statistics for competitive classification algorithms: Regularized logistic regression, CART, and random forest.

cues A list of cue information, with further details contained in `thresholds` and `stats`.

See Also

[print.FFTrees](#) for printing FFTs; [plot.FFTrees](#) for plotting FFTs; [summary.FFTrees](#) for summarizing FFTs; [inwords](#) for obtaining a verbal description of FFTs; [showcues](#) for plotting cue accuracies.

Examples

```
# Create fast-and-frugal trees (FFTs) for heart disease:
heart.fft <- FFTrees(formula = diagnosis ~ .,
                    data = heart.train,
                    data.test = heart.test,
                    main = "Heart Disease",
                    decision.labels = c("Healthy", "Diseased")
                    )

# Print a summary of the result:
heart.fft

# Plot an FFT applied to training data:
plot(heart.fft, stats = FALSE)
plot(heart.fft)

# Apply FFT to (new) testing data:
plot(heart.fft, data = "test")
plot(heart.fft, data = "test", tree = 2) # Plot Tree #2

# Predict classes and probabilities for new data:
predict(heart.fft, newdata = heartdisease)
predict(heart.fft, newdata = heartdisease, type = "prob")

# Create custom trees with my.tree:
custom.fft <- FFTrees(
  formula = diagnosis ~ .,
  data = heartdisease,
  my.tree = "If chol > 300, predict True.
            If sex = {m}, predict False,
            If age > 70, predict True, otherwise predict False."
)

# Plot the (pretty terrible) custom tree:
plot(custom.fft)
```

FFTrees.guide

*Open the **FFTrees** package guide*

Description

Open the **FFTrees** package guide

Usage

```
FFTrees.guide()
```

Value

No return value, called for side effects.

fftrees_cuerank	<i>Calculate thresholds that optimize some statistic (goal) for cues in data</i>
-----------------	--

Description

Calculate thresholds that optimize some statistic (goal) for cues in data

Usage

```
fftrees_cuerank(x = NULL, newdata = NULL, data = "train", rounding = NULL)
```

Arguments

x	An FFTrees object.
newdata	dataframe.
data	dataframe.
rounding	integer.

Value

A data frame containing thresholds and marginal classification statistics for each cue

fftrees_ffttowords	<i>Describe a fast-and-frugal tree (FFT) in words</i>
--------------------	---

Description

fftrees_ffttowords provides a verbal description of an FFT (in an FFTrees object).

fftrees_ffttowords is the complement to [fftrees_wordstoefftrees](#), which parses a verbal description of an FFT into an FFTrees object.

The final sentence (or tree node) of the FFT's description always predicts positive criterion values (i.e., TRUE instances) first, before predicting negative criterion values (i.e., FALSE instances). Note that this may require a reversal of cue directions (if the original tree description predicted FALSE instances before predicting TRUE instances).

Usage

```
fftrees_ffttowords(x = NULL, digits = 2)
```

Arguments

x An FFTrees object created with [FFTrees](#).
 digits How many digits to round numeric values (as integer)?

Value

A list of string vectors.

See Also

[fftrees_wordstoefftrees](#) for converting a verbal description of an FFT into an FFTrees object; [fftrees_create](#) for creating FFTrees objects; [fftrees_grow_fan](#) for creating FFTs by applying algorithms to data; [print.FFTrees](#) for printing FFTs; [plot.FFTrees](#) for plotting FFTs; [summary.FFTrees](#) for summarizing FFTs; [FFTrees](#) for creating FFTs from and applying them to data.

Examples

```
heart.fft <- FFTrees(diagnosis ~ .,
  data = heartdisease,
  decision.labels = c("Healthy", "Disease")
)

inwords(heart.fft)
```

 fftrees_fitcomp

Fit competitive algorithms

Description

`fftrees_fitcomp` fits competitive algorithms for binary classification tasks (e.g., LR, CART, RF, SVM) to the data and parameters specified in an FFTrees object.

`fftrees_fitcomp` is called by the main [FFTrees](#) function when creating FFTs from and applying them to data (unless `do.comp = FALSE`).

Usage

```
fftrees_fitcomp(x)
```

Arguments

x An FFTrees object.

See Also

[FFTrees](#) for creating FFTs from and applying them to data.

fftrees_grow_fan *Grow fast-and-frugal trees (FFTs) using the fan algorithm*

Description

fftrees_grow_fan is called by [fftrees_define](#) to create new FFTs by applying the fan algorithms (specifically, either ifan or dfan) to data.

Usage

```
fftrees_grow_fan(x, repeat.cues = TRUE)
```

Arguments

x An FFTrees object.
repeat.cues logical.

See Also

[fftrees_create](#) for creating FFTrees objects; [fftrees_define](#) for defining FFTs; [fftrees_grow_fan](#) for creating FFTs by applying algorithms to data; [fftrees_wordstoftrees](#) for creating FFTs from verbal descriptions; [FFTrees](#) for creating FFTs from and applying them to data.

fftrees_ranktrees *Rank FFTs by current goal*

Description

fftrees_ranktrees ranks trees in an FFTrees object x based on the current goal (either "cost" or as specified in x\$params\$goal).

fftrees_ranktrees is called by the main [FFTrees](#) function when creating FFTs from and applying them to data.

Usage

```
fftrees_ranktrees(x, data = "train")
```

Arguments

x An FFTrees object.
data character. Default is data = "train".

See Also

[FFTrees](#) for creating FFTs from and applying them to data.

```
fftrees_threshold_factor_grid
```

Perform a grid search over factor and return accuracy statistics for a given factor cue

Description

Perform a grid search over factor and return accuracy statistics for a given factor cue

Usage

```
fftrees_threshold_factor_grid(
  thresholds = NULL,
  cue_v = NULL,
  criterion_v = NULL,
  directions = "=",
  sens.w = 0.5,
  cost.outcomes = list(hi = 0, fa = 1, mi = 1, cr = 0),
  cost.each = 0,
  goal.threshold = "bacc"
)
```

Arguments

thresholds	numeric. A vector of factor thresholds to consider.
cue_v	numeric. Feature/cue values.
criterion_v	logical. A logical vector of (TRUE) criterion values.
directions	character. Character vector of threshold directions to consider.
sens.w	numeric. Sensitivity weight parameter (from 0 to 1, for computing wacc). Default: sens.w = .50.
cost.outcomes	list. A list of length 4 with names 'hi', 'fa', 'mi', and 'cr' specifying the costs of a hit, false alarm, miss, and correct rejection, respectively. For instance, cost.outcomes = listc("hi" = 0, "fa" = 10, "mi" = 20, "cr" = 0) means that a false alarm and miss cost 10 and 20, respectively, while correct decisions have no cost.
cost.each	numeric.
goal.threshold	character.

Value

A data frame containing accuracy statistics for several factor thresholds

See Also

[fftrees_threshold_numeric_grid](#) for numeric cues.

```
fftrees_threshold_numeric_grid
```

Perform a grid search over thresholds and return accuracy statistics for a given numeric cue

Description

Perform a grid search over thresholds and return accuracy statistics for a given numeric cue

Usage

```
fftrees_threshold_numeric_grid(
  thresholds,
  cue_v,
  criterion_v,
  directions = c(">", "<="),
  sens.w = 0.5,
  cost.each = 0,
  cost.outcomes = list(hi = 0, fa = 1, mi = 1, cr = 0),
  goal.threshold = "bacc"
)
```

Arguments

<code>thresholds</code>	numeric. A vector of thresholds to consider.
<code>cue_v</code>	numeric. Feature values.
<code>criterion_v</code>	logical. A logical vector of (TRUE) criterion values.
<code>directions</code>	character. Possible directions to consider.
<code>sens.w</code>	numeric. Sensitivity weight parameter (from 0 to 1, for computing wacc). Default: <code>sens.w = .50</code> .
<code>cost.each</code>	numeric. Cost to add to each value (e.g.; cost of the cue).
<code>cost.outcomes</code>	list. A list of length 4 with names 'hi', 'fa', 'mi', and 'cr' specifying the costs of a hit, false alarm, miss, and correct rejection, respectively. For instance, <code>cost.outcomes = listc("hi" = 0, "fa" = 10, "mi" = 20, "cr" = 0)</code> means that a false alarm and miss cost 10 and 20, respectively, while correct decisions have no cost.
<code>goal.threshold</code>	character. A string indicating the statistic to maximize when calculating cue thresholds: "acc" = overall accuracy, "wacc" = weighted accuracy, "bacc" = balanced accuracy.

Value

A data frame containing accuracy statistics for several numeric thresholds.

See Also

[fftrees_threshold_factor_grid](#) for factor cues.

fftrees_wordstoftrees

Convert a text description of an FFT into an FFTrees object

Description

fftrees_wordstoftrees converts a verbal description of an FFT (provided as a text string) into a tree definition (of an FFTrees object).

fftrees_wordstoftrees is the complement function to [fftrees_ffttowords](#), which converts a tree definition (of an FFTrees object) into a verbal description.

To increase robustness, the parsing of fftrees_wordstoftrees allows for lower- or uppercase spellings (but not typographical variants) and ignores the else-part of the final sentence (i.e., the part beginning with "otherwise").

Usage

```
fftrees_wordstoftrees(x, my.tree)
```

Arguments

x	An FFTrees object.
my.tree	string. A verbal string defining an FFT.

Value

An FFTrees object with a new tree definition as described by my.tree.

See Also

[fftrees_ffttowords](#) for converting FFTs into verbal descriptions; [print.FFTrees](#) for printing FFTs; [plot.FFTrees](#) for plotting FFTs; [summary.FFTrees](#) for summarizing FFTs; [FFTrees](#) for creating FFTs from and applying them to data.

forestfires	<i>Forest fires data</i>
-------------	--------------------------

Description

A dataset of forest fire statistics.

Usage

forestfires

Format

A data frame containing 517 rows and 13 columns.

X Integer -x-axis spatial coordinate within the Montesinho park map: 1 to 9

Y Integer - y-axis spatial coordinate within the Montesinho park map: 2 to 9

month Factor - month of the year: "jan" to "dec"

day Factor -day of the week: "mon" to "sun"

FFMC Numeric -FFMC index from the FWI system: 18.7 to 96.20

DMC Numeric - DMC index from the FWI system: 1.1 to 291.3

DC Numeric - DC index from the FWI system: 7.9 to 860.6

ISI Numeric - ISI index from the FWI system: 0.0 to 56.10

temp Numeric - temperature in Celsius degrees: 2.2 to 33.30

RH Numeric - relative humidity in percent: 15.0 to 100

wind Numeric - wind speed in km/h: 0.40 to 9.40

rain Numeric - outside rain in mm/m2 : 0.0 to 6.4

area Numeric - the burned area of the forest (in ha): 0.00 to 1090.84

...

Source

<http://archive.ics.uci.edu/ml/datasets/Forest+Fires>

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

heart.cost	<i>Cue costs for the heartdisease data</i>
------------	--

Description

This data further characterizes the variables (cues) in the [heartdisease](#) dataset.

Usage

```
heart.cost
```

Format

A data frame containing 153 rows and 14 columns.

cue The name of the cue

cost The cost of the cue

...

Source

<https://archive.ics.uci.edu/ml/machine-learning-databases/heart-disease/costs/>

See Also

[heartdisease](#) dataset.

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

heart.test	<i>Heart disease testing data</i>
------------	-----------------------------------

Description

Testing data for a [heartdisease](#) data. This subset is used to test the prediction performance of a model trained on the [heart.train](#) data. The dataset [heartdisease](#) contains both datasets.

Usage

```
heart.test
```

Format

A data frame containing 153 rows and 14 columns (see [heartdisease](#) for details).

Source

<https://archive.ics.uci.edu/ml/datasets/Heart+Disease>

See Also

[heartdisease](#) dataset.

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

heart.train

Heart disease training data

Description

Training data for a binary prediction model (here: FFT) on (a subset of) the [heartdisease](#) data. The complementary subset for model testing is [heart.test](#). The data in [heartdisease](#) contains both subsets.

Usage

```
heart.train
```

Format

A data frame containing 150 rows and 14 columns (see [heartdisease](#) for details).

Source

<https://archive.ics.uci.edu/ml/datasets/Heart+Disease>

See Also

[heartdisease](#) dataset.

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

heartdisease	<i>Heart disease data</i>
--------------	---------------------------

Description

A dataset predicting the diagnosis of 303 patients tested for heart disease.

Usage

heartdisease

Format

A data frame containing 303 rows and 14 columns, with the following variables:

diagnosis True value of binary criterion: TRUE = Heart disease, FALSE = No Heart disease

age Age (in years)

sex Sex, 1 = male, 0 = female

cp Chest pain type: ta = typical angina, aa = atypical angina, np = non-anginal pain, a = asymptomatic

trestbps Resting blood pressure (in mm Hg on admission to the hospital)

chol Serum cholesterol in mg/dl

fbs Fasting blood sugar > 120 mg/dl: 1 = true, 0 = false

restecg Resting electrocardiographic results. "normal" = normal, "abnormal" = having ST-T wave abnormality (T wave inversions and/or ST elevation or depression of > 0.05 mV), "hypertrophy" = showing probable or definite left ventricular hypertrophy by Estes' criteria.

thalach Maximum heart rate achieved

exang Exercise induced angina: 1 = yes, 0 = no

oldpeak ST depression induced by exercise relative to rest

slope The slope of the peak exercise ST segment.

ca Number of major vessels (0-3) colored by fluoroscopy

thal "normal" = normal, "fd" = fixed defect, "rd" = reversible defect

...

Source

<https://archive.ics.uci.edu/ml/datasets/Heart+Disease>

See Also

[heart.cost](#) dataset for cost information.

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

inwords	<i>Provide a verbal description of an FFT</i>
---------	---

Description

inwords generates and provides a verbal description of a fast-and-frugal tree (FFT) from an FFTrees object.

Usage

```
inwords(x, tree = 1)
```

Arguments

x	An FFTrees object.
tree	The tree to display (as numeric).

Value

A verbal description of an FFT (as a string).

See Also

[print.FFTrees](#) for printing FFTs; [plot.FFTrees](#) for plotting FFTs; [summary.FFTrees](#) for summarizing FFTs; [FFTrees](#) for creating FFTs from and applying them to data.

iris.v	<i>Iris data</i>
--------	------------------

Description

A famous dataset from R.A. Fisher (1936) simplified to predict only the virginica class (i.e., as a binary classification problem).

Usage

```
iris.v
```

Format

A data frame containing 150 rows and 4 columns.

Source

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

References

Fisher, R.A. (1936): The use of multiple measurements in taxonomic problems. *Annual Eugenics*, 7, Part II, pp. 179–188.

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

 mushrooms

Mushrooms data

Description

Data describing poisonous vs. non-poisonous mushrooms.

Usage

mushrooms

Format

A data frame containing 8,124 rows and 23 columns.

See <http://archive.ics.uci.edu/ml/machine-learning-databases/mushroom/agaricus-lepiota.names> for column descriptions.

poisonous logical criterion variable

cshape character

csurface character

ccolor character

bruises character

odor numeric

gattach character

gspace character

gsize character

gcolor character

sshape character

sroot character

ssaring character

ssbring character

scaring character

scbring character
vtype character
vcolor character
ringnum character
ringtype character
sporepc character
population character
habitat character
 ...

Details

This dataset includes descriptions of hypothetical samples corresponding to 23 species of gilled mushrooms in the Agaricus and Lepiota Family. Each species is classified as poisonous (True or False). The Guide clearly states that there is no simple rule for determining the edibility of a mushroom; no rule like “leaflets three, let it be” for Poisonous Oak and Ivy.

Source

<http://archive.ics.uci.edu/ml/datasets/Mushroom>

References

Mushroom records drawn from The Audubon Society Field Guide to North American Mushrooms (1981). G.H. Lincoff (Pres.), New York: A.A. Knopf.

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [sonar](#), [titanic](#), [voting](#), [wine](#)

plot.FFTrees

Plot an FFTrees object

Description

plot.FFTrees visualizes an FFTrees object created by the [FFTrees](#) function.

plot.FFTrees is the main plotting function of the **FFTrees** package and called when evaluating the generic [plot](#) on an FFTrees object.

plot.FFTrees visualizes a selected FFT, key data characteristics, and various aspects of classification performance.

As x may not contain test data, plot.FFTrees by default plots the performance characteristics for training data (i.e., fitting), rather than for test data (i.e., for prediction). When test data is available, specifying data = "test" plots prediction performance.

Whenever the sensitivity weight (`sens.w`) is set to its default of `sens.w = 0.50`, a level shows *balanced* accuracy (`bacc`). If, however, `sens.w` deviates from its default, the level shows the tree's *weighted* accuracy value (`wacc`) and the current `sens.w` value (below the level).

Many aspects of the plot (e.g., its panels) and the FFT's appearance (e.g., labels of its nodes and exits) can be customized by setting corresponding arguments.

Usage

```
## S3 method for class 'FFTrees'
plot(
  x = NULL,
  data = "train",
  what = "tree",
  tree = 1,
  main = NULL,
  cue.labels = NULL,
  decision.labels = NULL,
  cue.cex = NULL,
  threshold.cex = NULL,
  decision.cex = 1,
  comp = TRUE,
  stats = TRUE,
  show.header = NULL,
  show.tree = NULL,
  show.confusion = NULL,
  show.levels = NULL,
  show.roc = NULL,
  show.icons = NULL,
  show.iconguide = NULL,
  hlines = TRUE,
  label.tree = NULL,
  label.performance = NULL,
  n.per.icon = NULL,
  which.tree = NULL,
  level.type = "bar",
  decision.names = NULL,
  ...
)
```

Arguments

<code>x</code>	An FFTrees object created by the <code>FFTrees</code> function.
<code>data</code>	The data in <code>x</code> to be plotted (as a string); must be either <code>'train'</code> (for fitting performance) or <code>'test'</code> (for prediction performance). By default, <code>data = 'train'</code> (as <code>x</code> may not contain test data).
<code>what</code>	What should be plotted (as a string)? <code>'tree'</code> (the default) shows details of one tree (specified by <code>tree</code>); <code>'cues'</code> shows the marginal accuracy of cues in ROC

	space; 'roc' shows the performance of tree(s) (and comparison algorithms) in ROC space.
tree	The tree to be plotted (as an integer, only valid when the corresponding tree argument is non-empty). Default: tree = 1. To plot the best training or best test tree with respect to the goal specified during FFT construction, use "best.train" or "best.test", respectively.
main	The main plot label (as a character string).
cue.labels	An optional string of labels for the cues / nodes (as character vector).
decision.labels	A character vector of length 2 indicating the content-specific names for noise and signal predictions/exits.
cue.cex	The size of the cue labels (as numeric).
threshold.cex	The size of the threshold labels (as numeric).
decision.cex	The size of the decision labels (as numeric).
comp	Should the performance of competitive algorithms (e.g.; logistic regression, random forests, etc.) be shown in the ROC plot (if available, as logical)?
stats	Should statistical information be plotted (as logical)? If FALSE, only the tree diagram (without any reference to statistics) will be plotted.
show.header	Show header with basic data properties (in top panel, as logical)?
show.tree	Show nodes and exits of FFT (in middle panel, as logical)?
show.confusion	Show 2x2 confusion matrix (in bottom panel, as logical)?
show.levels	Show performance levels (in bottom panel, as logical)?
show.roc	Show ROC curve (in bottom panel, as logical)?
show.icons	Show exit cases as icon arrays (in middle panel, as logical)?
show.iconguide	Show icon guide (in middle panel, as logical)?
hlines	Show horizontal panel separation lines (as logical)? Default: hlines = TRUE.
label.tree	Label for the FFT (optional, as character string).
label.performance	Labels for the performance section (optional, as character string).
n.per.icon	Number of cases per icon (as numeric).
which.tree	deprecated argument, included for backwards compatibility, use "tree" instead.
level.type	How should bottom levels be drawn (as a string)? Can be "bar" (the default) or "line".
decision.names	deprecated argument.
...	Graphical parameters (passed either to <code>showcues</code> when what = 'cues' or to <code>title</code> when what = 'roc').

Value

A plot visualizing and describing an FFT.

See Also

[showcues](#) for plotting cue accuracies; [print.FFTrees](#) for printing FFTs; [summary.FFTrees](#) for summarizing FFTs; [FFTrees](#) for creating FFTs from and applying them to data.

Other plot functions: [showcues\(\)](#)

Examples

```
# Create FFTs (for heartdisease data):
heart.fft <- FFTrees(formula = diagnosis ~ .,
                    data = heartdisease
                    )

# Visualize the default FFT (Tree #1):
plot(heart.fft,
     main = "Heart Disease Diagnosis",
     decision.labels = c("Absent", "Present")
     )

# Visualize FFT #2 (with customized labels):
plot(heart.fft,
     tree = 2,
     main = "An FFT for heart disease diagnosis",
     cue.labels = c("1. thal?", "2. cp?", "3. ca?", "4. exang"),
     decision.labels = c("ok", "sick"),
     show.header = FALSE,
     show.confusion = FALSE,
     show.levels = FALSE,
     show.roc = FALSE
     )

# Visualize cue accuracies:
plot(heart.fft, what = "cues")

# For more details, see
vignette("FFTrees_plot", package = "FFTrees")
```

predict.FFTrees

Predict classification outcomes or probabilities from data

Description

`predict.FFTrees` predicts binary classification outcomes or their probabilities from newdata for an `FFTrees` object.

Usage

```
## S3 method for class 'FFTrees'
predict(
  object = NULL,
  newdata = NULL,
  tree = 1,
  type = "class",
  sens.w = NULL,
  method = "laplace",
  data = NULL,
  ...
)
```

Arguments

object	An FFTrees object created by the FFTrees function.
newdata	dataframe. A data frame of test data.
tree	integer. Which tree in the object should be used? By default, tree = 1 is used.
type	string. What should be predicted? Can be "class", which returns a vector of class predictions, "prob" which returns a matrix of class probabilities, or "both" which returns a matrix with both class and probability predictions.
sens.w, data	deprecated
method	string. Method of calculating class probabilities. Either 'laplace', which applies the Laplace correction, or 'raw' which applies no correction.
...	Additional arguments passed on to predict.

Value

Either a logical vector of predictions, or a matrix of class probabilities.

See Also

[print.FFTrees](#) for printing FFTs; [plot.FFTrees](#) for plotting FFTs; [summary.FFTrees](#) for summarizing FFTs; [FFTrees](#) for creating FFTs from and applying them to data.

Examples

```
# Create training and test data:
set.seed(100)
breastcancer <- breastcancer[sample(nrow(breastcancer)), ]
breast.train <- breastcancer[1:150, ]
breast.test <- breastcancer[151:303, ]

# Create an FFTrees object from the training data:
breast.fft <- FFTrees(
  formula = diagnosis ~ .,
  data = breast.train
)
```

```
# Predict classes for test data:
breast.fft.pred <- predict(breast.fft,
  newdata = breast.test
)

# Predict class probabilities for test data:
breast.fft.pred <- predict(breast.fft,
  newdata = breast.test,
  type = "prob"
)
```

print.FFTrees *Print basic information of fast-and-frugal trees (FFTs)*

Description

print.FFTrees prints basic information on FFTs for an FFTrees object x.

As x may not contain test data, print.FFTrees by default prints the performance characteristics for training data (i.e., fitting), rather than for test data (i.e., for prediction). When test data is available, specify data = "test" to print prediction performance.

Usage

```
## S3 method for class 'FFTrees'
print(x = NULL, tree = 1, data = "train", ...)
```

Arguments

x	An FFTrees object created by FFTrees .
tree	The tree to be printed (as an integer, only valid when the corresponding tree argument is non-empty). Default: tree = 1. To print the best training or best test tree with respect to the goal specified during FFT construction, use "best.train" or "best.test", respectively.
data	The data in x to be printed (as a string); must be either 'train' (for fitting performance) or 'test' (for prediction performance). By default, data = 'train' (as x may not contain test data).
...	additional arguments passed to print.

Value

Prints summary information about an FFT to the console.

See Also

[plot.FFTrees](#) for plotting FFTs; [summary.FFTrees](#) for summarizing FFTs; [inwords](#) for obtaining a verbal description of FFTs; [FFTrees](#) for creating FFTs from and applying them to data.

select_best_tree	Select the best tree (from the current set)
------------------	---

Description

select_best_tree selects (looks up and identifies) the best tree from the set (or “fan”) of FFTs contained in the current FFTrees object `x`, an existing type of data (`'train'` or `'test'`), and a goal for which corresponding statistics are available in the designated data type (in `x$trees$stats`).

Usage

```
select_best_tree(x, data, goal)
```

Arguments

<code>x</code>	An FFTrees object.
<code>data</code>	character. Must be either "train" or "test".
<code>goal</code>	character. A goal to maximize or minimize when selecting a tree from an existing <code>x</code> (for which values exist in <code>x\$trees\$stats</code>).

Details

Importantly, `select_best_tree` only identifies and selects from the set of *existing* trees with known statistics, rather than creating new trees or computing new cue thresholds. More specifically, `goal` is used for identifying and selecting the best of an existing set of FFTs, but not for computing new cue thresholds (see `goal.threshold` and `fftrees_cuerank()`) or creating new trees (see `goal.chase` and `fftrees_ranktrees()`).

Value

An integer denoting the tree that maximizes/minimizes `goal` in `data`.

See Also

[FFTrees](#) for creating FFTs from and applying them to data.

showcues	<i>Visualize cue accuracies (as points in ROC space)</i>
----------	--

Description

showcues plots the cue accuracies of an FFTrees object created by the [FFTrees](#) function (as points in ROC space).

If the optional arguments `cue.accuracies` and `alt.goal` are specified, their values take precedence over the corresponding settings of an FFTrees object `x` (but do not change `x`).

showcues is called when the main [plot.FFTrees](#) function is set to what = "cues".

Usage

```
showcues(
  x = NULL,
  cue.accuracies = NULL,
  alt.goal = NULL,
  main = NULL,
  top = 5,
  quiet = FALSE,
  ...
)
```

Arguments

<code>x</code>	An FFTrees object created by the FFTrees function.
<code>cue.accuracies</code>	An optional data frame specifying cue accuracies directly (without specifying FFTrees object <code>x</code>).
<code>alt.goal</code>	An optional alternative goal to sort the current cue accuracies (without using the goal of FFTrees object <code>x</code>).
<code>main</code>	A main plot title (as character string).
<code>top</code>	How many of the top cues should be highlighted (as an integer)?
<code>quiet</code>	Should user feedback messages be printed (as logical)? Default: <code>quiet = FALSE</code> (i.e., show messages).
<code>...</code>	Graphical parameters (passed to plot).

Value

A plot showing cue accuracies (of an FFTrees object) (as points in ROC space).

See Also

[print.FFTrees](#) for printing FFTs; [plot.FFTrees](#) for plotting FFTs; [summary.FFTrees](#) for summarizing FFTs; [FFTrees](#) for creating FFTs from and applying them to data.

Other plot functions: [plot.FFTrees\(\)](#)

Examples

```
# Create fast-and-frugal trees (FFTs) for heart disease:
heart.fft <- FFTrees(formula = diagnosis ~ .,
                    data = heart.train,
                    data.test = heart.test,
                    main = "Heart Disease",
                    decision.labels = c("Healthy", "Diseased")
                    )

# Show cue accuracies (in ROC space):
showcues(heart.fft,
        main = "Predicting heart disease")
```

sonar

Sonar data

Description

Sonar data

Usage

sonar

Format

A data frame containing 208 rows and 60 columns.

Source

[https://archive.ics.uci.edu/ml/datasets/Connectionist+Bench+\(Sonar,+Mines+vs.+Rocks\)](https://archive.ics.uci.edu/ml/datasets/Connectionist+Bench+(Sonar,+Mines+vs.+Rocks))

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [titanic](#), [voting](#), [wine](#)

summary.FFTrees	<i>Summarize an FFTrees object</i>
-----------------	------------------------------------

Description

summary.FFTrees summarizes key contents of an FFTrees object.

Usage

```
## S3 method for class 'FFTrees'
summary(object, tree = NULL, ...)
```

Arguments

object	An FFTrees object.
tree	The tree to summarize (as an integer, but may be a vector). If tree = NULL (as per default) or exceeding the possible range 1:object\$trees\$n, information on all trees in object is returned.
...	Additional arguments (currently ignored).

Details

summary.FFTrees returns an invisible list containing two elements:

1. definitions and corresponding performance measures of trees;
2. stats on decision frequencies, derived probabilities, and costs (separated by train and test).

A header prints descriptive information of the FFTrees object (to the console): Its main title, number of trees (object\$trees\$n), and the name of the criterion variable (object\$criterion_name).

Per default, information on all available trees is shown and returned. Specifying tree filters the output list elements for the corresponding tree(s). When only a single tree is specified, the printed header includes a verbal description of the corresponding tree.

While summary.FFTrees provides key details about the specified tree(s), the individual decisions (stored in object\$trees\$decisions) are not shown or returned.

Value

An invisible list with elements containing the definitions and performance stats of the FFT(s) specified by tree(s).

See Also

[print.FFTrees](#) for printing FFTs; [plot.FFTrees](#) for plotting FFTs; [inwords](#) for obtaining a verbal description of FFTs; [FFTrees](#) for creating FFTs from and applying them to data.

titanic	<i>Titanic survival data</i>
---------	------------------------------

Description

Data indicating who survived on the Titanic.

Usage

```
titanic
```

Format

A data frame containing 2,201 rows and 4 columns.

class Factor - Class (first, second, third, or crew)

age Factor - Age group (child or adult)

sex Factor - Sex (male or female)

survived Factor - Whether the passenger survived (1) or not (0)

...

Details

See [Titanic](#) of the R **datasets** package for details and the same data (in a 4-dimensional table).

Source

<https://www.encyclopedia-titanica.org>

References

Dawson, Robert J. MacG. (1995), The 'Unusual Episode' Data Revisited. *Journal of Statistics Education*, 3. doi: 10.1080/10691898.1995.11910499.

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [voting](#), [wine](#)

voting

Voting data

Description

A dataset of votes for each of the U.S. House of Representatives Congressmen on the 16 key votes identified by the CQA.

Usage

voting

Format

A data frame containing 435 rows and 16 columns.

Details

The CQA lists nine different types of votes: voted for, paired for, and announced for (these three simplified to yea), voted against, paired against, and announced against (these three simplified to nay), voted present, voted present to avoid conflict of interest, and did not vote or otherwise make a position known (these three simplified to an unknown disposition).

The binary criterion variable used here is `party.crit`.

Source

<https://archive.ics.uci.edu/ml/datasets/Congressional+Voting+Records>

References

Congressional Quarterly Almanac, 98th Congress, 2nd session 1984, Volume XL: Congressional Quarterly Inc. Washington, D.C., 1985.

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [wine](#)

wine

Wine tasting data

Description

Chemical and tasting data from wines in North Portugal.

Usage

wine

Format

A data frame containing 6497 rows and 13 columns.

Source

<http://archive.ics.uci.edu/ml/datasets/Wine+Quality>

See Also

Other datasets: [blood](#), [breastcancer](#), [car](#), [contraceptive](#), [creditapproval](#), [fertility](#), [forestfires](#), [heart.cost](#), [heart.test](#), [heart.train](#), [heartdisease](#), [iris.v](#), [mushrooms](#), [sonar](#), [titanic](#), [voting](#)

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