Package 'BT'

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Title (Adaptive) Boosting Trees Algorithm

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Depends R (>= 4.0)

Imports rpart, stats, statmod, parallel

Suggests rmarkdown, knitr, testthat (>= 3.0.0)

Description Performs (Adaptive) Boosting Trees for Poisson distributed response variables, using loglink function.

The code approach is similar to the one used in 'gbm'/'gbm3'. Moreover, each tree in the expansion is built thanks to the 'rpart' package.

This package is based on following books and articles

Denuit, M., Hainaut, D., Trufin, J. (2019) <doi:10.1007/978-3-030-25820-7> Denuit, M., Hainaut, D., Trufin, J. (2019) <doi:10.1007/978-3-030-57556-4> Denuit, M., Hainaut, D., Trufin, J. (2019) <doi:10.1007/978-3-030-25827-6> Denuit, M., Hainaut, D., Trufin, J. (2022) <doi:10.1080/03461238.2022.2037016> Denuit, M., Huyghe, J., Trufin, J. (2022) <https://dial.uclouvain.be/pr/boreal/fr/ object/boreal%3A244325/datastream/PDF_01/view> Denuit, M., Trufin, J., Verdebout, T. (2022) <https: //dial.uclouvain.be/pr/boreal/fr/object/boreal%3A268577>.

URL https://github.com/GiregWillame/BT/

BugReports https://github.com/GiregWillame/BT/issues/

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ΒT

(Adaptive) Boosting Trees (ABT/BT) Algorithm.

Description

Performs the (Adaptive) Boosting Trees algorithm. This code prepares the inputs and calls the function BT_call. Each tree in the process is built thanks to the rpart function. In case of cross-validation, this function prepares the folds and performs multiple calls to the fitting function BT_call.

Usage

```
BT(
  formula = formula(data),
  data = list(),
  tweedie.power = 1,
  ABT = TRUE,
  n.iter = 100,
  train.fraction = 1,
  interaction.depth = 4,
  shrinkage = 1,
  bag.fraction = 1,
  colsample.bytree = NULL,
  keep.data = TRUE,
  is.verbose = FALSE,
  cv.folds = 1,
  folds.id = NULL,
  n.cores = 1,
```

```
tree.control = rpart.control(xval = 0, maxdepth = (if (!is.null(interaction.depth)) {
```

```
interaction.depth
} else {
    10
}), cp = -Inf, minsplit = 2),
weights = NULL,
seed = NULL,
....
)
```

Arguments

| formula | a symbolic description of the model to be fit. Note that the offset isn't supported in this algorithm. Instead, everything is performed with a log-link function and a direct relationship exist between response, offset and weights. | |
|-------------------|--|--|
| data | an optional data frame containing the variables in the model. By default the variables are taken from environment(formula), typically the environment from which BT is called. If keep.data=TRUE in the initial call to BT then BT stores a copy with the object (up to the variables used). | |
| tweedie.power | Experimental parameter currently not used - Set to 1 referring to Poisson distribution. | |
| ABT | a boolean parameter. If ABT=TRUE an adaptive boosting tree algorithm is built whereas if ABT=FALSE an usual boosting tree algorithm is run. By default, it is set to TRUE. | |
| n.iter | the total number of iterations to fit. This is equivalent to the number of trees and the number of basis functions in the additive expansion. Please note that the initialization is not taken into account in the n.iter. More explicitly, a weighted average initializes the algorithm and then n.iter trees are built. Moreover, note that the bag.fraction, colsample.bytree, are not used for this initializing phase. By default, it is set to 100. | |
| train.fraction | the first train.fraction * nrows(data) observations are used to fit the BT and the remainder are used for computing out-of-sample estimates (also known as validation error) of the loss function. By default, it is set to 1 meaning no out-of-sample estimates. | |
| interaction.depth | | |
| | the maximum depth of variable interactions: 1 builds an additive model, 2 builds a model with up to two-way interactions, etc. This parameter can also be inter- preted as the maximum number of non-terminal nodes. By default, it is set to 4. Please note that if this parameter is NULL, all the trees in the expansion are built based on the tree.control parameter only, independently of the ABT value. This option is devoted to advanced users only and allows them to benefit from the full flexibility of the implemented algorithm. | |
| shrinkage | a shrinkage parameter (in the interval (0,1]) applied to each tree in the expansion. Also known as the learning rate or step-size reduction. By default, it is set to 1. | |
| bag.fraction | the fraction of independent training observations randomly selected to propose the next tree in the expansion. This introduces randomness into the model fit. If | |

bag.fraction<1 then running the same model twice will result in similar but different fits. Please note that if this parameter is used the BTErrors\$training.error corresponds to the normalized in-bag error and the out-of-bag improvements are computed and stored in BTErrors\$oob.improvement. See BTFit for more details. By default, it is set to 1. colsample.bytree each tree will be trained on a random subset of colsample.bytree number of features. Each tree will consider a new random subset of features from the formula, adding variability to the algorithm and reducing computation time. colsample.bytree will be bounded between 1 and the number of features considered in the formula. By default, it is set to NULL meaning no effect. keep.data a boolean variable indicating whether to keep the data frames. This is particularly useful if one wants to keep track of the initial data frames and is further used for predicting in case any data frame is specified. Note that in case of cross-validation, if keep.data=TRUE the initial data frames are saved whereas the cross-validation samples are not. By default, it is set to FALSE. is.verbose if is.verbose=TRUE, the BT will print out the algorithm progress. By default, it is set to FALSE. cv.folds a positive integer representing the number of cross-validation folds to perform. If cv. folds>1 then BT, in addition to the usual fit, will perform a cross-validation and calculate an estimate of generalization error returned in BTErrors\$cv.error. By default, it is set to 1 meaning no cross-validation. folds.id an optional vector of values identifying what fold each observation is in. If supplied, this parameter prevails over cv.folds. By default, folds.id = NULL meaning that no folds are defined. n.cores the number of cores to use for parallelization. This parameter is used during the cross-validation. This parameter is bounded between 1 and the maximum number of available cores. By default, it is set to 1 leading to a sequential approach. tree.control for advanced user only. It allows to define additional tree parameters that will be used at each iteration. See rpart.control for more information. weights optional vector of weights used in the fitting process. These weights must be positive but do not need to be normalized. By default, it is set to NULL which corresponds to an uniform weight of 1 for each observation. seed optional number used as seed. Please note that if cv.folds>1, the parLapply function is called. Therefore, the seed (if defined) used inside each fold will be a multiple of the seed parameter. not currently used. . . .

Details

The NA values are currently dropped using na.omit.

Value

a BTFit object.

Author(s)

Gireg Willame <gireg.willame@gmail.com>

This package is inspired by the gbm3 package. For more details, see https://github.com/gbm-developers/gbm3/.

References

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries I: GLMs and Extensions, *Springer Actuarial*.

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries II: Tree-Based Methods and Extensions, *Springer Actuarial*.

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M. Denuit, J. Trufin and T. Verdebout (2022). Boosting on the responses with Tweedie loss functions. Paper submitted for publication.

See Also

BTFit,BTCVFit,BT_call,BT_perf,predict.BTFit,summary.BTFit,print.BTFit,.BT_cv_errors.

Examples

```
## Load dataset.
dataset <- BT::BT_Simulated_Data</pre>
## Fit a Boosting Tree model.
BT_algo <- BT(formula = Y_normalized ~ Age + Sport + Split + Gender, # formula
              data = dataset, # data
              ABT = FALSE, # Classical Boosting Tree
              n.iter = 200,
              train.fraction = 0.8,
              interaction.depth = 3,
              shrinkage = 0.01,
              bag.fraction = 0.5,
              colsample.bytree = 2, # 2 explanatory variable used at each iteration.
              keep.data = FALSE, # Do not keep a data copy.
              is.verbose = FALSE, # Do not print progress.
              cv.folds = 3, # 3-cv will be performed.
              folds.id = NULL ,
              n.cores = 1,
              weights = ExpoR, # <=> Poisson model on response Y with ExpoR in offset.
              seed = NULL)
```

```
## Determine the model performance and plot results.
best_iter_val <- BT_perf(BT_algo, method='validation')</pre>
best_iter_oob <- BT_perf(BT_algo, method='00B', oobag.curve = TRUE)</pre>
best_iter_cv <- BT_perf(BT_algo, method ='cv', oobag.curve = TRUE)</pre>
best_iter <- best_iter_val</pre>
## Variable influence and plot results.
# Based on the first iteration.
variable_influence1 <- summary(BT_algo, n.iter = 1)</pre>
# Using all iterations up to best_iter.
variable_influence_best_iter <- summary(BT_algo, n.iter = best_iter)</pre>
## Print results : call, best_iters and summarized relative influence.
print(BT_algo)
## Model predictions.
# Predict on the link scale, using only the best_iter tree.
pred_single_iter <- predict(BT_algo, newdata = dataset,</pre>
                             n.iter = best_iter, type = 'link', single.iter = TRUE)
# Predict on the response scale, using the first best_iter.
pred_best_iter <- predict(BT_algo, newdata = dataset,</pre>
                           n.iter = best_iter, type = 'response')
```

Description

These are objects representing CV fitted boosting trees.

Details

CV (Adaptive) Boosting Tree Model Object.

Value

a list of BTFit objects with each element corresponding to a specific BT fit on a particular fold

Structure

The following components must be included in a legitimate BTCVFit object.

Author(s)

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This package is inspired by the gbm3 package. For more details, see https://github.com/gbm-developers/gbm3/.

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BTFit

References

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries I: GLMs and Extensions, *Springer Actuarial*.

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries II: Tree-Based Methods and Extensions, *Springer Actuarial*.

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries III: Neural Networks and Extensions, *Springer Actuarial*.

M. Denuit, D. Hainaut and J. Trufin (2022). **Response versus gradient boosting trees, GLMs and neural networks under Tweedie loss and log-link**. Accepted for publication in *Scandinavian Actuarial Journal*.

M. Denuit, J. Huyghe and J. Trufin (2022). **Boosting cost-complexity pruned trees on Tweedie responses: The ABT machine for insurance ratemaking**. Paper submitted for publication.

M. Denuit, J. Trufin and T. Verdebout (2022). Boosting on the responses with Tweedie loss functions. Paper submitted for publication.

See Also

BT.

BTFit BTFit

Description

These are objects representing fitted boosting trees.

Details

Boosting Tree Model Object.

Value

| BTInit | an object of class BTInit containing the initial fitted value initFit, the initial training.error and the initial validation.error if any. |
|--------------|---|
| BTErrors | an object of class BTErrors containing the vectors of errors for each iteration performed (excl. the initialization). More precisely, it contains the training.error, validation.error if train.fraction<1 and the oob.improvement if bag.fraction < 1. Moreover, if a cross-validation approach was performed, a vector of cross-validation errors cv.error as a function of boosting iteration is also stored in this object. |
| BTIndivFits | an object of class BTIndivFits containing the list of each individual tree fitted at each boosting iteration. |
| distribution | the Tweedie power (and so the distribution) that has been used to perform the algorithm. It will currently always output 1. |

| var.names | a vector containing the names of the explanatory variables. |
|---------------|---|
| response | the name of the target/response variable. |
| W | a vector containing the weights used. |
| seed | the used seed, if any. |
| BTData | if keep.data=TRUE, an object of class BTData containing the training.set and validation.set (can be NULL if not used). These data frames are reduced to the used variables, that are the response and explanatory variables. Note that in case of cross-validation, even if keep.data=TRUE the folds will not be kept. In fact, only the data frames related to the original fit (i.e. on the whole training set) will be saved. |
| BTParams | an object of class BTParams containing all the (Adaptive) boosting tree parameters. More precisely, it contains the ABT, train.fraction, shrinkage, interaction.depth, bag.fraction, n.iter, colsample.bytree and tree.control parameter values. |
| keep.data | the keep.data parameter value. |
| is.verbose | the is.verbose parameter value. |
| fitted.values | the training set fitted values on the score scale using all the n.iter (and initial- ization) iterations. |
| cv.folds | the number of cross-validation folds. Set to 1 if no cross-validation performed. |
| call | the original call to the BT algorithm. |
| Terms | the model.frame terms argument. |
| folds | a vector of values identifying to which fold each observation is in. This argument is not present if there is no cross-validation. On the other hand, it corresponds to folds.id if it was initially defined by the user. |
| cv.fitted | a vector containing the cross-validation fitted values, if a cross-validation was performed. More precisely, for a given observation, the prediction will be fur- nished by the cv-model for which this specific observation was out-of-fold. See predict.BTCVFit for more details. |

Structure

The following components must be included in a legitimate BTFit object.

Author(s)

Gireg Willame < gireg.willame@gmail.com>

This package is inspired by the gbm3 package. For more details, see https://github.com/gbm-developers/gbm3/.

References

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries I: GLMs and Extensions, *Springer Actuarial*.

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries II: Tree-Based Methods and Extensions, *Springer Actuarial*.

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M. Denuit, J. Trufin and T. Verdebout (2022). Boosting on the responses with Tweedie loss functions. Paper submitted for publication.

See Also

BT.

BT_call

(Adaptive) Boosting Trees (ABT/BT) fit.

Description

Fit a (Adaptive) Boosting Trees algorithm. This is for "power" users who have a large number of variables and wish to avoid calling model.frame which can be slow in this instance. This function is in particular called by BT. It is mainly split in two parts, the first one considers the initialization (see BT_callInit) whereas the second performs all the boosting iterations (see BT_callBoosting). By default, this function does not perform input checks (those are all done in BT) and all the parameters should be given in the right format. We therefore suppose that the user is aware of all the choices made.

Usage

```
BT_call(
  training.set,
  validation.set,
  tweedie.power,
  respVar,
  w,
  explVar,
  ABT,
  tree.control,
  train.fraction,
  interaction.depth,
  bag.fraction,
  shrinkage,
  n.iter,
  colsample.bytree,
  keep.data,
  is.verbose
```

```
BT_callInit(training.set, validation.set, tweedie.power, respVar, w)
```

```
BT_callBoosting(
  training.set,
 validation.set,
  tweedie.power,
 ABT,
  tree.control,
  interaction.depth,
  bag.fraction,
  shrinkage,
  n.iter,
  colsample.bytree,
  train.fraction,
  keep.data,
  is.verbose,
  respVar,
 w,
 explVar
)
```

Arguments

| training.set | a data frame containing all the related variables on which one wants to fit the algorithm. | |
|-------------------|--|--|
| validation.set | a held-out data frame containing all the related variables on which one wants to assess the algorithm performance. This can be NULL. | |
| tweedie.power | Experimental parameter currently not used - Set to 1 referring to Poisson distribution. | |
| respVar | the name of the target/response variable. | |
| W | a vector of weights. | |
| explVar | a vector containing the name of explanatory variables. | |
| ABT | a boolean parameter. If ABT=TRUE an adaptive boosting tree algorithm is built whereas if ABT=FALSE an usual boosting tree algorithm is run. | |
| tree.control | allows to define additional tree parameters that will be used at each iteration. See rpart.control for more information. | |
| train.fraction | the first train.fraction * nrows (data) observations are used to fit the BT and the remainder are used for computing out-of-sample estimates (also known as validation error) of the loss function. It is mainly used to report the value in the BTFit object. | |
| interaction.depth | | |
| | the maximum depth of variable interactions: 1 builds an additive model, 2 builds a model with up to two-way interactions, etc. This parameter can also be inter- preted as the maximum number of non-terminal nodes. By default, it is set to 4. | |

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)

| | Please note that if this parameter is NULL, all the trees in the expansion are built based on the tree.control parameter only. This option is devoted to advanced users only and allows them to benefit from the full flexibility of the implemented algorithm. |
|----------------|---|
| bag.fraction | the fraction of independent training observations randomly selected to propose the next tree in the expansion. This introduces randomness into the model fit. If bag.fraction<1 then running the same model twice will result in similar but different fits. BT uses the R random number generator, so set.seed ensures the same model can be reconstructed. Please note that if this parameter is used the BTErrors\$training.error corresponds to the normalized in-bag error. |
| shrinkage | a shrinkage parameter applied to each tree in the expansion. Also known as the learning rate or step-size reduction. |
| n.iter | the total number of iterations to fit. This is equivalent to the number of trees and the number of basis functions in the additive expansion. Please note that the initialization is not taken into account in the n.iter. More explicitly, a weighted average initializes the algorithm and then n.iter trees are built. Moreover, note that the bag.fraction, colsample.bytree, are not used for this initializing phase. |
| colsample.bytr | ee |
| | each tree will be trained on a random subset of colsample.bytree number of features. Each tree will consider a new random subset of features from the formula, adding variability to the algorithm and reducing computation time. colsample.bytree will be bounded between 1 and the number of features considered. |
| keep.data | a boolean variable indicating whether to keep the data frames. This is particu- larly useful if one wants to keep track of the initial data frames and is further used for predicting in case any data frame is specified. Note that in case of cross-validation, if keep.data=TRUE the initial data frames are saved whereas the cross-validation samples are not. |
| is.verbose | if is.verbose=TRUE, the BT will print out the algorithm progress. |

Value

a **BTFit** object.

Author(s)

Gireg Willame < gireg.willame@gmail.com>

This package is inspired by the gbm3 package. For more details, see https://github.com/gbm-developers/gbm3/.

References

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries I: GLMs and Extensions, *Springer Actuarial*.

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries II: Tree-Based Methods and Extensions, *Springer Actuarial*.

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M. Denuit, J. Trufin and T. Verdebout (2022). Boosting on the responses with Tweedie loss functions. Paper submitted for publication.

See Also

BTFit, BTCVFit, BT_perf, predict.BTFit, summary.BTFit, print.BTFit, .BT_cv_errors.

| BT_devTweedie | Deviance function for the Tweedie family. | |
|---------------|---|--|
|---------------|---|--|

Description

Compute the deviance for the Tweedie family case.

Usage

```
BT_devTweedie(y, mu, tweedieVal, w = NULL)
```

Arguments

| У | a vector containing the observed values. |
|------------|---|
| mu | a vector containing the fitted values. |
| tweedieVal | a numeric representing the Tweedie Power. It has to be a positive number outside of the interval]0,1[. |
| W | an optional vector of weights. |

Details

This function computes the Tweedie related deviance. The latter is defined as:

$$\begin{split} d(y,mu,w) &= w(y-mu)^2, if twee die Val = 0; \\ d(y,mu,w) &= 2w(ylog(y/mu)+mu-y), if twee die Val = 1; \\ d(y,mu,w) &= 2w(log(mu/y)+y/mu-1), if twee die Val = 2; \\ d(y,mu,w) &= 2w(max(y,0)^{(2}-p)/((1-p)(2-p))-ymu^{(1-p)}/(1-p)+mu^{(2-p)}/(2-p)), else die Val = 2; \end{split}$$

Value

A vector of individual deviance contribution.

BT_more

Author(s)

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This package is inspired by the gbm3 package. For more details, see https://github.com/gbm-developers/gbm3/.

References

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries I: GLMs and Extensions, *Springer Actuarial*.

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M. Denuit, J. Trufin and T. Verdebout (2022). Boosting on the responses with Tweedie loss functions. Paper submitted for publication.

See Also

BT, BT_call.

BT_more

Perform additional boosting iterations.

Description

Method to perform additional iterations of the Boosting Tree algorithm, starting from an initial BTFit object. This does not support further cross-validation. Moreover, this approach is only allowed if keep.data=TRUE in the original call.

Usage

```
BT_more(BTFit_object, new.n.iter = 100, is.verbose = FALSE, seed = NULL)
```

Arguments

| BTFit_object | a BTFit object. |
|--------------|---|
| new.n.iter | number of new boosting iterations to perform. |
| is.verbose | a logical specifying whether or not the additional fitting should run "noisely" with feedback on progress provided to the user. |
| seed | optional seed used to perform the new iterations. By default, no seed is set. |

Returns a new BTFit object containing the initial call as well as the new iterations performed.

Author(s)

Gireg Willame <gireg.willame@gmail.com>

This package is inspired by the gbm3 package. For more details, see https://github.com/gbm-developers/gbm3/.

References

M. Denuit, D. Hainaut and J. Trufin (2019). Effective Statistical Learning Methods for Actuaries I: GLMs and Extensions, *Springer Actuarial*.

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M. Denuit, J. Trufin and T. Verdebout (2022). Boosting on the responses with Tweedie loss functions. Paper submitted for publication.

See Also

BT, BTFit.

BT_perf

Performance assessment.

Description

Function to compute the performances of a fitted boosting tree.

Usage

```
BT_perf(
   BTFit_object,
   plot.it = TRUE,
   oobag.curve = FALSE,
   overlay = TRUE,
   method,
   main = ""
)
```

BT_perf

Arguments

| BTFit_object | a BTFit object resulting from an initial call to BT |
|--------------|--|
| plot.it | a boolean indicating whether to plot the performance measure. Setting plot.it = TRUE creates two plots. The first one plots the object\$BTErrors\$training.error (in black) as well as the object\$BTErrors\$validation.error (in red) and/or the object\$BTErrors\$cv.error (in green) depending on the method and parametriza- tion. These values are plotted as a function of the iteration number. The scale of the error measurement, shown on the left vertical axis, depends on the arguments used in the initial call to BT and the chosen method. |
| oobag.curve | indicates whether to plot the out-of-bag performance measures in a second plot. Note that this option makes sense if the bag.fraction was properly defined in the initial call to BT. |
| overlay | if set to TRUE and oobag.curve=TRUE then a right y-axis is added and the esti- mated cumulative improvement in the loss function is plotted versus the iteration number. |
| method | indicates the method used to estimate the optimal number of boosting itera- tions. Setting method = "OOB" computes the out-of-bag estimate and method = "validation" uses the validation dataset to compute an out-of-sample esti- mate. Finally, setting method = " cv " extracts the optimal number of iterations using cross-validation, if BT was called with $cv.folds > 1$. If missing, a guess- ing method is applied. |
| main | optional parameter that allows the user to define specific plot title. |

Value

Returns the estimated optimal number of iterations. The method of computation depends on the method argument.

Author(s)

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This package is inspired by the gbm3 package. For more details, see https://github.com/gbm-developers/gbm3/.

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See Also

BT, BT_call.

BT_Simulated_Data Simulated Database.

Description

A simulated database used for examples and vignettes. The variables are related to a motor insurance pricing context.

Usage

BT_Simulated_Data

Format

A simulated data frame with 50,000 rows and 7 columns, containing simulation of different policyholders:

Gender Gender, varying between male and female.

Age Age, varying from 18 to 65 years old.

- **Split** Noisy variable, not used to simulate the response variable. It allows to assess how the algorithm handle these features.
- Sport Car type, varying between yes (sport car) or no.

ExpoR Yearly exposure-to-risk, varying between 0 and 1.

Y Yearly claim number, simulated thanks to Poisson distribution.

Y_normalized Yearly claim frequency, corresponding to the ratio between Y and ExpoR.

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predict.BTFit

Description

Predicted values based on a boosting tree model object.

Usage

```
## S3 method for class 'BTFit'
predict(object, newdata, n.iter, type = "link", single.iter = FALSE, ...)
```

Arguments

| object | a BTFit object. |
|-------------|---|
| newdata | data frame of observations for which to make predictions. If missing or not a data frame, if keep.data=TRUE in the initial fit then the original training set will be used. |
| n.iter | number of boosting iterations used for the prediction. This parameter can be a vector in which case predictions are returned for each iteration specified. |
| type | the scale on which the BT makes the predictions. Can either be "link" or "re- sponse". Note that, by construction, a log-link function is used during the fit. |
| single.iter | if single.iter=TRUE then predict.BTFit returns the predictions from the sin- gle tree n.iter. |
| | not currently used. |

Details

predict.BTFit produces a predicted values for each observation in newdata using the first n.iter boosting iterations. If n.iter is a vector then the result is a matrix with each column corresponding to the BT predictions with n.iter[1] boosting iterations, n.iter[2] boosting iterations, and so on.

As for the fit, the predictions do not include any offset term. In the Poisson case, please remind that a weighted approach is initially favored.

Value

Returns a vector of predictions. By default, the predictions are on the score scale. If type = "response", then BT converts back to the same scale as the outcome. Note that, a log-link is supposed by construction.

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M. Denuit, J. Trufin and T. Verdebout (2022). Boosting on the responses with Tweedie loss functions. Paper submitted for publication.

See Also

BT, BTFit.

print.BTFit Printing function.

Description

Function to print the BT results.

Usage

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

Arguments

| Х | a BTFit object. |
|---|------------------------------------|
| | arguments passed to print.default. |

Details

Print the different input parameters as well as obtained results (best iteration/performance & relative influence) given the chosen approach.

Value

No value returned.

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See Also

BT, .BT_relative_influence, BT_perf.

summary.BTFit Summary of a BTFit object.

Description

Computes the relative influence of each variable in the BTFit object.

Usage

```
## S3 method for class 'BTFit'
summary(
    object,
    cBars = length(object$var.names),
    n.iter = object$BTParams$n.iter,
    plot_it = TRUE,
    order_it = TRUE,
    method = .BT_relative_influence,
    normalize = TRUE,
    ...
)
```

Arguments

| object | a BTFit object. |
|-----------|--|
| cBars | the number of bars to plot. If order=TRUE only the variables with the cBars largest relative influence will appear in the barplot. If order=FALSE then the first cBars variables will appear in the barplot. |
| n.iter | the number of trees used to compute the relative influence. Only the first n.iter trees will be used. |
| plot_it | an indicator as to whether the plot is generated. |
| order_it | an indicator as to whether the plotted and/or returned relative influences are sorted. |
| method | the function used to compute the relative influence. Currently, only .BT_relative_influence is available (default value as well). |
| normalize | if TRUE returns the normalized relative influence. |
| | additional argument passed to the plot function. |

Details

Please note that the relative influence for variables having an original **negative** relative influence is forced to 0.

Value

Returns a data frame where the first component is the variable name and the second one is the computed relative influence, normalized to sum up to 100. Depending on the plot_it value, the relative influence plot will be performed.

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See Also

BT, .BT_relative_influence.

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