Package ‘TDMR’

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Type Package
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Description Tuned Data Mining in R ('TDMR') performs the complete tuning of a data mining task (predictive analytics, that is classification and regression). Preprocessing parameters and modeling parameters can be tuned simultaneously. It incorporates a variety of tuners (among them 'SPOT' and 'CMA' with package 'rCMA') and allows integration of additional tuners. Noise handling in the data mining optimization process is supported, see Koch et al. (2015) <doi:10.1016/j.asoc.2015.01.005>.
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  'tdmDispatchTuner.r' 'tdmEnvTMakeNew.r' 'tdmGeneralUtils.r'
  'tdmGraphicUtils.r' 'tdmMapDesign.r' 'tdmMetacostRf.r'
  'tdmModelingUtils.r' 'tdmOptsDefaults.r' 'tdmParaBootstrap.r'
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TDMR-package

Tuned Data Mining in R

Description

Tuned Data Mining in R

Details

Package: TDMR
Type: Package
Version: 2.0
Date: 04.05.2018
License: GPL (>= 2)
LazyLoad: yes

TDMR is a package for tuned data mining (predictive analytics, i.e. classification and regression). Its main features are:
1) A variety of tuners, with special emphasis on SPOT (a well-known R package for parameter tuning), but also CMA-ES (package rCMA) and other tuning algorithms.
2) Tuning of preprocessing parameters and model building parameters simultaneously. Preprocessing often includes feature generation.
3) Support for multiple tuning experiments (different settings, repetitions with different resamplings, ...).
4) Easy parallelization of those experiments with the help of R package parallel.
5) Extensibility: New tuning parameters, new preprocessing tools, model builders and even new tuners can be added easily.

The main entry point functions are tdmClassifyLoop, tdmRegressLoop, tdmTuneIt, and tdmBigLoop.
See `tdmOptsDefaultsSet` and `tdmDefaultsFill` for an overview of adjustable TDMR-parameters.

Author(s)

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), Patrick Koch

References

http://lwibs01.gm fh-koeln.de/blogs/ciop/research/tuned-data-mining/

---

**defaultOpts**

Default settings for the data mining part of TDMR (list opts).

**Description**

Sets suitable defaults for the data mining part of TDMR.

**Usage**

`defaultOpts()`

**Details**

With the call `setParams(myOpts, defaultOpts())` it is possible to extend a partial list `myOpts` to a list containing all `opts`-elements (the missing ones are taken from `defaultOpts()`). If `myOpts` has an element not present in `defaultOpts()`, this element is not taken and a warning is issued. With `setParams(myOpts, defaultOpts(), keepNotMatching=TRUE)` also elements of `myOpts` not present in `defaultOpts()` are taken (no warnings).

**Value**

a list with the elements according to `tdmOptsDefaultsSet`

**Author(s)**

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), Samineh Bagheri, THK, 2018

**See Also**

`setParams`, `defaultSC`
defaultSC

Default settings for the spotConfig part of TDMR.

Description

Sets suitable defaults for the spotConfig part of TDMR.

Usage

defaultSC()

Details

With the call setParams(mySC, defaultSC()) it is possible to extend a partial list mySC to a list containing all sC-elements (the missing ones are taken from defaultSC()). If mySC has an element not present in defaultSC(), this element is not taken and a warning is issued.

With setParams(mySC, defaultSC(), keepNotMatching=TRUE) also elements of mySC not present in defaultSC() are taken (no warnings).

Value

a list with the following elements (the values in parantheses [ ] are the defaults):

- alg.roi ["NEEDS_TO_BE_SET"] a data frame with columns lower, upper, type, row.names, each a vector with as many entries as there are parameter to be tuned
- opts ["NEEDS_TO_BE_SET"]
- sCName ["NEEDS_TO_BE_SET.conf"] a string ending on ".conf", the configuration name
- OCBA [FALSE] see spotControl
- plot [FALSE] TRUE: make a line plot showing progress
- seedSPOT [1] see spotControl
- funEvals [50] the budget, max number of algo evaluations
- design [designLHD] function that creates initial design, see spotControl
- designControl.size [10] number of initial design points (former init.design.size)
- designControl.replicates [2] number of initial repeats (former init.design.repeats)
- replicates [2] number of repeats for the same model design point
- noise [TRUE] whether the object function has noise or not (Note: TRUE is required if replicates>1 (!))
- seq.merge.func [mean] how to merge Y over replicates: mean or min
- model [buildKriging] function that builds the surrogate model, see spotControl
- optimizer [optimLHD] function that optimizes on surrogate, see spotControl
- optimizerControl.funEvals [100] optimizer budget (former seq.design.size)
- optimizerControl.retries [2] optimLHD retries (former seq.design.retries)
Author(s)

Wolfgang Konen (<wolfgang.konen@th-koeln.de>) THK, 2018

See Also

setparams, defaultOpts

---

dsetTest.TDMdata  
Return test data of TDMdata object

Description

Return the test part of a TDMdata object containing the task data.

Usage

```r
## S3 method for class 'TDMdata'
dsetTest(x, ...)
```

Arguments

- `x` return value from a prior call to `tdmReadAndSplit`, an object of class TDMdata.
- `...` may contain nExp, experiment number, needed only if `x$tdm$umode=="SP_T"`: add nExp to seed when randomly splitting in train and test data [default: nExp=0]

Value

tset, a data frame with all test records. If there are 0 test records, return NULL.

Author(s)

Wolfgang Konen, THK

See Also

unbiasedRun dsetTrnVa.TDMdata tdmReadAndSplit
**dsetTrnVa.TDMdata**

*Return train-validation data of TDMdata object*

---

**Description**

Return the train-validation part of a TDMdata object containing the task data.

**Usage**

```r
## S3 method for class 'TDMdata'
dsetTrnVa(x, ...)
```

**Arguments**

- `x` return value from a prior call to `tdmReadAndSplit`, an object of class TDMdata.
- `...` may contain nExp, experiment number, needed only if `xsdumodem="SP_T"`:
  add nExp to seed when randomly splitting in train and test data [default: nExp=0]

**Value**

- `dset`, a data frame with all train-validation records

**Author(s)**

Wolfgang Konen, THK

**See Also**

- `dsetTest.TDMdata`
- `tdmReadAndSplit`

---

**Opts**

*Return the list 'opts'.*

---

**Description**

Returns the list opts from objects of class TDMenvir, TDMclassifier, TDMregressor, tdmClass or tdmRegre.
Usage

```
Options(x, ...)
```

## S3 method for class 'TDMenvir'
Options(x, ...)

## S3 method for class 'TDMclassifier'
Options(x, ...)

## S3 method for class 'TDMregressor'
Options(x, ...)

## S3 method for class 'tdmClass'
Options(x, ...)

## S3 method for class 'tdmRegre'
Options(x, ...)

## Default S3 method:
Options(x, ...)

Arguments

- `x` is an object of class `TDMenvir, TDMclassifier, tdmClass, TDMregressor` or `tdmRegre`.
- `...` is currently not used.

Value

- the list `opts` with DM-specific settings contained in the specified object

---

**predict.TDMenvir**

_Make a prediction using the last model._

Description

Make a prediction with objects of class `TDMenvir, TDMclassifier, TDMregressor`. The prediction is based on the (last) model trained during `unbiasedRun`.

Usage

```
## S3 method for class 'TDMenvir'
predict(object, ...)
```

## S3 method for class 'TDMclassifier'
predict(object, ...)

---
print.TDMclassifier

## S3 method for class 'TDMregressor'
predict(object, ...)

### Arguments

- **object**: an object of class `TDMenvir, TDMclassifier, TDMregressor` containing in element `lastModel` the relevant model.
- **...**: arguments passed on to the model's predict function. Usually the first argument of ... should be `newdata`, a data frame for which new predictions are desired.

### Value

A vector with length `nrow(newdata)` containing the new predictions.

### Examples

```r
## Not run:
## This example requires that demo04cpu.r is executed first (it will write demo04cpu.RData)
path <- paste(find.package("TDMR"), "demo04cpu/", sep="/");
wd <- list( filenameEnvT="demo04cpu.RData" ); # file with environment envT
load(paste(path,wd$filenameEnvT,sep="/"));

# take only the first 15 records:
newdata=read.csv2(file=paste(path,"data/cpu.csv", sep=""), dec=".")[1:15,];
z=predict(envT,newdata);
print(z);

## End(Not run)
```

---

print.TDMclassifier  

*Print an overview for a TDMclassifier object.*

### Description

Print an overview for a `TDMclassifier` or `tdmClass` object.

### Usage

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

```r
## S3 method for class 'tdmClass'
print(x, ...)
```
Arguments

x an object of class `tdmClass`, as returned from a prior call to `tdmClassify`, or an object of class `TDMclassifier`, as returned from a prior call to `tdmClassifyLoop`.

... e.g. 'type' which information to print:
“overview” (default) relative gain on training/test set, number of records, see `tdmClassifySummary`
“cm.train” confusion matrix on train set
“cm.vali” confusion matrix on test set
“?” help on this method

Author(s)

Wolfgang Konen, THK

See Also

`tdmClassify`, `tdmClassifySummary`, `TDMclassifier`
print.TDMregressor

Print an overview for a TDMregressor object.

Description

Print an overview for a TDMregressor or tdmRegre object.

Usage

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

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

Arguments

- `x` an object of class `tdmRegre`, as returned from a prior call to `tdmRegres`, or an object of class `TDMregressor`, as returned from a prior call to `tdmRegressLoop`.
- `...` e.g. 'type' which information to print:
  - "overview" (def.) RMAE on training/test set, number of records, see `tdmRegressSummary`
  - "..." ... other choices, TODO ...
  - "?" help on this method

Author(s)

Wolfgang Konen, THK

See Also

`tdmRegres, tdmRegressSummary, TDMregressor`

setParams

Merge the parameters from a partial list and the default list

Description

Merge the parameters from a partial list and the default list

Usage

```r
setParams(opts, defaultOpt, keepNotMatching = FALSE)
```
Arguments

- opts: a partial list of parameters
- defaultOpt: a list with default values for every element
- keepNotMatching: [FALSE] if TRUE, copy the elements appearing in opts, but not in defaultOpt to the return value. If FALSE, do not copy them, but issue a warning.

Value

A list combined from opts and defaultOpt where every available element in opts overrides the default. For the rest of the elements the value from defaultOpt is taken.

A warning is issued for every element appearing in opts but not in defaultOpt (only if keepNotMatching=FALSE).

Author(s)

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), Samineh Bagheri

See Also

defaultSC, defaultOpts

tdmBigLoop  Tuning and unbiased evaluation in a big loop.

Description

For each configuration object .conf in tdm$runList call all tuning algorithms (SPOT, CMA-ES or other) specified in tdm$tuneMethod (via function tdmDispatchTuner). After each tuning process perform a run of tdm$unbiasedFunc (usually unbiasedRun). Each of these experiments is repeated tdm$nExperim times. Thus we have for each tripel

(confName,nExp,theTuner)

a tuning result. The ranges of the triple elements are:

confName in tdm$runList
nExp in 1,...,tdm$nExperim
theTuner in tdm$tuneMethod

Usage

tdmBigLoop(envT, dataObj = NULL)
Arguments

**envT**
an environment containing on input at least the element `tdm` (a list with general settings for TDMR, see `tdmDefaultsFill`), which has at least the elements `tdm$runList` vector of configuration names. 

**dataObj**
[NULL] optional object of class `TDMData` (the same for all runs in big loop). If it is NULL, it will be constructed here with the help of `tdmReadAndSplit`. Then it can be different for each configuration object in the big loop.

Details

`tdm` refers to `envT$tdm`.

The available tuning algorithms (tuners) are

- **spotTuner**: Call `spot`.
- **1hdTuner**: Perform a parameter tuning using a Latin hypercube design (LHD) for obtaining best design points. LHD is performed by configuring SPOT in such a way that all the budget is used for the initial design (usually LHD).
- **cma_jTuner**: Perform a parameter tuning by CMA-ES, using the *Java* implementation by Niko Hansen through the interface package `rcma`.
- **cmaesTuner**: Perform a parameter tuning by CMA-ES, using the *R*-implementation (package `cma_es` by Olaf Mersmann) (deprecated, use `cma_jTuner` instead).
- **bfgsTuner**: Perform a parameter tuning by Broyden, Fletcher, Goldfarb and Shanno (BFGS) method. The L-BFGS-B version allowing box constraints is used.
- **powellTuner**: Perform a parameter tuning by Powell’s UObyQA algorithm (unconstrained optimization by quadratic approximation), see package `powell`.

Value

environment `envT`, containing the results

**res**
data frame with results from last tuning (one line for each call of `tdmStart*`)

**bst**
data frame with the best-so-far results from last tuning (one line collected after each (SPO) step)

**resGrid**
list with data frames `res` from all tuning runs. Use `envT$getRes(envT,confFile,nExp,theTuner)` to retrieve a specific `res`.

**bstGrid**
list with data frames `bst` from all tuning runs. Use `envT$getBst(envT,confFile,nExp,theTuner)` to retrieve a specific `bst`.

**theFinals**
data frame with one line for each triple (`confFile`, `nExp`, `tuner`), each line contains summary information about the tuning run in the form: `confFile tuner nExp [params] NRUN NEVAL RGain bst RGain.* sdR.*` where `[params]` is written depending on `tdm$withParams`. `NRUN` is the number of unbiased evaluation runs. `NEVAL` is the number of function evaluations (model builds) during tuning.
RGain denotes the relative gain on a certain data set: the actual gain achieved with the model divided by the maximum gain possible for the current cost matrix and the current data set. This is for classification tasks, in the case of regression each RGain.* is replaced by RMAE.*, the relative mean absolute error. Each ‘sdR.’ denotes the standard deviation of the preceding RGain or RMAE. RGain.bst is the best result during tuning obtained on the training-validation data. RGain.avg is the average result during tuning. The following pairs RGain.* sdR.* are the results of one or several unbiased evaluations on the test data where '*' takes as many values as there are elements in tdmDumode (the possible values are explained in unbiasedRun).

result object of class TDMclassifier or TDMregressor. This is a list with results from tdm$mainFunc as called in the last unbiased evaluation using the best parameters found during tuning. Use print(envT$result) to get more info on such an object of class TDMclassifier.

tunerVal an object with the return value from the last tuning process. For every tuner, this is the list spotConfig, containing the SPOT settings plus the TDMR settings in elements opts and tdm. Every tuner extends this list by tunerVal$alg.currentResult and tunerVal$alg.currentBest, see tdmDispatchTuner. In addition, each tuning method might add specific elements to the list, see the description of each tuner.

Environment envT contains further elements, but they are only relevant for the internal operation of tdmBigLoop and its subfunctions.

Note

Side effects: A compressed version of envT is saved to file tdm$filenameEnvT (default: <runList[1]>.RData) in directory tdm$path. If tdm$path=NULL use the current directory. If tdm$saveModel==TRUE, then envT$result$lastRes$lastModel (the last trained model) will be saved to tdm$filenameEnvT. The default is tdm$saveModel==TRUE. If tdm$saveModel==FALSE then smaller .RData files will result.

Example usages of function tdmBigLoop are shown in

    demo(demo03sonar)
    demo(demo03sonar_B)
    demo(demo04cpu)

where the corresponding R-sources are in directory demo.

Author(s)

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), THK, Patrick Koch

See Also

   tdmDispatchTuner, unbiasedRun
Examples

```r
### This demo shows a complete tuned data mining process (level 3 of TDMR) where
### the data mining task is the classification task SONAR (from UCI repository,
### The data mining process is in main_sonar.r, which calls tdmClassifyLoop and tdmClassify
### with Random Forest as the prediction model.
### The three parameter to be tuned are CUTOFF1, CLASSWT2 and XPERC, as specified
### in controlSC() (control_sonar.r). The tuner used here is LHD.
### Tuning runs are rather short, to make the example run quickly.
### Do not expect good numeric results.
### See demo/demo03sonar_B.r for a somewhat longer tuning run, with two tuners SPOT and LHD.

### path is the dir with data and main_* file:
path <- paste(find.package("TDMR"), "demo02sonar", sep="/");
#path <- paste("./.../inst", "demo02sonar", sep="/");

### control settings for TDMR
tdm <- list(mainFunc="main_sonar"
  , runList = c("sonar_04.conf")
  , umode="CV"  # { "CV" | "RSUB" | "TST" | "SP_T" }
  , tuneMethod = c("lhd")
  , filenameEnvT="exBigLoop.RData"  # file to save environment envT
  , nrun=1, nfold=2  # repeats and CV-folds for the unbiased runs
  , nExperim=1
  , optsVerbosity = 0  # the verbosity for the unbiased runs
);
source(paste(path,"main_sonar.r",sep="/"));  # main_sonar, readTnrSonar

### This demo is for example and help {more meaningful, a bit higher budget}
source(paste(path,"control_sonar.r",sep="/"));  # controlDM, controlSC

ctr1SC <- controlSC();
ctr1SC$opts <- controlDM();

# construct envT from settings given in tdm & sList
envT <- tdmEnvMakeNew(tdm,sList=list(ctr1SC));
dataObj <- tdmReadTaskData(envT,envT$tdm);
envT <- tdmBigLoop(envT,dataObj=dataObj);  # start the big tuning loop
```

**tdmBindResponse**

*Bind a column to a data frame.*

**Description**

Bind the column with name `response` and contents `vec` as last column to data frame `d`
**tdmClassify**

Usage

`tdmBindResponse(d, response.predict, vec)`

Arguments

- `d`: data frame
- `response.predict`: name of new column
- `vec`: the contents for the last column bound to data frame `d`

Value

data frame `d` with column added

---

**tdmClassify**  
*Core classification function of TDMR.*

Description

tdmClassify is called by `tdmClassifyLoop` and returns an object of class `tdmClass`

It trains a model on training set `d_train` and evaluates it on test set `d_test`. If this function is used for tuning, the test set `d_test` plays the role of a validation set.

Usage

`tdmClassify(d_train, d_test, d_dis, d_preproc, response.variables, input.variables, opts, tsetStr = c("Validation", "validation"))`

Arguments

- `d_train`: training set
- `d_test`: validation set, same columns as training set
- `d_dis`: 'disregard set', i.e. everything what is neither train nor test. The model is applied to all records in `d_dis` (needed for active learning, see `ssl_methods.r`)
- `d_preproc`: data used for preprocessing. May be NULL, if no preprocessing is done (`opts$PRE.SFA=="none"` and `opts$PRE.PCA=="none"`). If preprocessing is done, then `d_preproc` is usually all non-validation data.
- `response.variables`: name of column which carries the target variable - or - vector of names specifying multiple target columns (these columns are not used during prediction, only for evaluation)
- `input.variables`: vector with names of input columns
- `opts`: additional parameters [defaults in brackets]
  - `SRF.*`: several parameters for `tdmModSortedRImport`
tdmClassify

RF. * several parameters for RF (Random Forest, defaults are set, if omitted)
SVM. * several parameters for SVM (Support Vector Machines, defaults are set, if omitted)
filename
data.title

MOD.method ["RF"] the main training method ["RF"|"MC.RF"|"SVM"|"NB"]: use [Random forest| MetaCost-RF| SVM| Naive Bayes] for the main model
MOD.SEED =NULL: get a new random number seed with tdmRandomSeed (different RF trainings).
=any value: set the random number seed to this value (+i) to get reproducible random numbers. In this way, the model training part (RF, NNET, ...) gets always a fixed seed (see also TST.SEED in tdmClassifyLoop)
CLASSWT class weights (NULL, if all classes should have the same weight) (currently used only by methods RF, MC.RF and by tdmModSortedRFinport)
fct.postproc [NULL] name of user-def'd function for postprocessing of predicted output
GO.DEVICE if !="non", then make a pairs-plot of the 5 most important variables and make a true-false bar plot
VERBOSE [2] =2: most printed output, =1: less, =0: no output
tsetStr [c("Validation", "validation")]

Details

Currently d_dis is allowed to be a 0-row data frame, but d_train and d_test must have at least one record.

Value

res, an object of class tdmClass, this is a list containing
d_train training set + predicted class column(s)
d_test test set + predicted class column(s)
d_dis disregard set + predicted class column(s)
avgEVAL list with evaluation measures, averaged over all response variables
allEVAL data frame with evaluation measures, one row for each response variable
lastCmTrain a list with evaluation info for training set (confusion matrix, gain, class errors, ...)
lastCmVali a list with evaluation info for validation set (confusion matrix, gain, class errors, ...)
lastModel the last model built (i.e. for the last response variable)
lastProbs a list with three probability matrices (row: records, col: classes) v_train, v_test, v_dis, if the model provides probabilities; NULL else.
lastPred name of the column where the prediction of the last model is appended to the datasets d_train, d_test and d_dis
predProb

A list with two data frames `Trn` and `Val`. They contain at least a column `IND.dset` (index of each train / validation record into data frame `dset`). If the model has probabilities, then they contain in addition a column for each response variable with the prediction probabilities.

`opts`

Parameter list from input, some default values might have been added.

The 9 evaluation measures in `avgEVAL` and `allEVAL` are `cerr.*` (misclassification error), `gain.*` (total gain) and `rgain.*` (relative gain, i.e. total gain divided by max. achievable gain in *) where * = [trn | tst | tst2 ] stands for [ training set | test set | test set with special treatment ] and the special treatment is either `opts$test2.string = "no postproc"` or `"default cutoff"`.

The five items `lastCmTrain`, `lastCmVali`, `lastModel`, `lastProbs`, `lastPred` are specific for the *last* model (the one built for the last response variable in the last run and last fold).

**Author(s)**

Wolfgang Konen, THK, 2013

**See Also**

`print.tdmClassify` `tdmClassifyLoop` `tdmRegressLoop`

**Examples**

```r
### This demo shows a simple data mining process (phase 1 of TDMR) for classification on
### dataset `iris`.
### The data mining process in `tdmClassify` calls `randomForest` as the prediction model.
### It is called `opts$NRun=1` time with one random train-validation set splits.
### Therefore data frame `res$allEval` has one row
###
### `opts=tdmOptDefaultsSet()` # set all defaults for data mining process
### `gdObj <- tdmGraAndLogInitialize(opts);` # init graphics and log file
###
### `data(iris)`
### `response.variables="Species"` # names, not data (!)
### `input.variables=setdiff(names(iris),"Species")`
### `opts$NRun=1`
###
### `idx_train = sample(nrow(iris))[1:110]`
### `d_train=iris[idx_train,]`
### `d_vali=iris[-idx_train,]`
### `d_dis=iris[numeric(0),]`
### `res <- tdmClassify(d_train,d_vali,d_dis,NULL,response.variables,input.variables,opts)`
###
### `cat("\n")`
### `print(res$allEval)`
```
tdmClassifyLoop Core classification double loop returning a TDMclassifier object.

Description

tdmClassifyLoop contains a double loop (opts$NRUN and CV-folds) and calls tdmClassify. It is called by all classification R-functions main_*.
It splits - if tset is NULL - the data in dset into training and validation data according to opts$TST.kind.
It returns an object of class TDMclassifier.

Usage

tdmClassifyLoop(dset, response.variables, input.variables, opts, tset = NULL)

Arguments

dset the data frame containing training and validation data.
response.variables name of column which carries the target variable - or - vector of names specifying multiple target columns (these columns are not used during prediction, only for evaluation)
input.variables vector with names of input columns
opts a list from which we need here the following entries
NRUN number of runs (outer loop)
TST.SEED =NULL: get a new random number seed with tdmRandomSeed. =any value: set the random number seed to this value to get reproducible random numbers and thus reproducible training-test-set-selection. (only relevant in case TST.kind="cv" or "rand") (see also MOD.SEED in tdmClassify)
TST.kind how to create cvi, handed over to tdmModCreateCVindex. If TST.kind="col", then cvi is taken from dset[,opts$TST.col].
GD.RESTART [TRUE] =TRUE/FALSE: do/don't restart graphic devices
GD.DEVICE ["non"|"win"|"pdf"|"png"]
tset [NULL] If not NULL, this is the test data set. If NULL, we are in tuning and the validation data set is build from dset according to the procedure prescribed in opts$TST.*.

Value

result, an object of class TDMclassifier, this is a list with results, containing
lastRes last run, last fold: result from tdmClassify
C_train classification error on training set
G_train gain on training set
R_train relative gain on training set (percentage of max. gain on this set)
tdmClassifyLoop

*_vali* — similar, with vali set instead of training set —

*_vali2* — similar, with vali2 set instead of training set —

Err a data frame with as many rows as opts$NRUN and 9 columns corresponding to
the nine variables described above

predictions last run: data frame with dimensions [nrow(dset),length(response.variable)]. In
case of CV, all CV predictions (for each record in dset), in other cases mixed
validation / train set predictions.

predictTest predictions on the test set tset (NULL if tset==NULL)

predProbList a list, predProbList[[i]] has the prediction probabilities of the ith run. See
info on predProb in tdmClassify.

Each performance measure C_*, G_* , R_* is a vector of length opts$NRUN. To be specific,
C_train[i] is the classification error on the training set from the i-th run. This error is mean(res$sallEVAL$cerr.trn),
i.e. the mean of the classification errors from all response variables when res is the return value of
tdmClassify. In the case of cross validation, for each performance measure an additional averaging
over all folds is done.

Author(s)

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), THK

See Also

print.TDMclassifier, tdmClassify, tdmRegress, tdmRegressLoop

Examples

```r
## This demo shows a simple data mining process (phase 1 of TDMR) for classification on
## dataset iris.
## The data mining process in tdmClassifyLoop calls randomForest as the prediction model.
## It is called opts$NRUN=2 times with different random train-validation set splits.
## Therefore data frame result$Err has two rows

opts=tdmOpt$DefaultsSet()               # set all defaults for data mining process
opts$TST.SEED <- opts$MOD.SEED <- 5     # reproducible results
#opts$VERBOSE <- opts$SRF.VERBOSE <- 0  # no printed output
gdObj <- tdmGraAndLogInitialize(opts);  # init graphics and log file

data(iris)
response.variables="Species"            # names, not data (!)
input.variables=setdiff(names(iris),"Species")

result = tdmClassifyLoop(iris,response.variables,input.variables,opts)

print(result$Err)
```
tdmClassifySummary

Print summary output for result from tdmClassifyLoop and add result$y.

Description

result$y is "minus OOB rgain" on training set for methods RF or MC.RF. result$y is "minus rgain" on test set (=validation set) for all other methods. result$y is the quantity which the tuner seeks to minimize.

Usage

tdmClassifySummary(result, opts, dset = NULL)

Arguments

result  
return value from a prior call to tdmClassifyLoop, an object of class TDMclassifier.

opts  
a list from which we need here the following entries

NRUN  number of runs (outer loop)

method

VERBOSE

dset [NULL] if !=NULL, attach it to result

Value

result, an object of class TDMclassifier, with result$y, result$sd.y (and optionally also result$dset) added

Author(s)

Wolfgang Konen, FHK, Sep’2010 - Oct’2011

See Also

tdmClassify, tdmClassifyLoop, print.TDMclassifier, tdmRegessummary
**tdmDefaultsFill**

Default values for list tdm.

**Description**

This list controls the tuning and unbiased evaluation phase. When called with `tdm = tdmDefaultsFill()`, a new list `tdm` is created and returned. When called with `tdm = tdmDefaultsFill(mainFile="my.r")`, a new list `tdm` is created and returned, with the element `mainFile` set to the specified value. When called with `tdm = tdmDefaultsFill(tdm)`, an existing list `tdm` is filled with further default values.

**Usage**

```r
tdmDefaultsFill(tdm = NULL, mainFile = NULL)
```

**Arguments**

- `tdm` (optional)
- `mainFile` (optional) if given, create or overwrite `tdm$mainFile` with this value

**Details**

If `tdm$mainFunc` is missing, but `tdm$mainFile` exists, then `tdmDefaultsFill` will set

```r
tdm$mainFunc = sub(".r","", basename(tdm$mainFile), fixed=TRUE)
```

**Value**

`tdm` the new / extended list, where additional elements, if they are not yet def’d, are set as:

- `mainFile` [NULL] if not NULL, source this file from the current dir. It should contain the definition of `tdm$mainFunc`.
- `mainFunc` `sub(".r","", basename(tdm$mainFile), fixed=TRUE)`, if `tdm$mainFile` is set and `tdm$mainFunc` is NULL, else "mainFunc" This is the name of the function called in `tdmStartSpot2` and `unbiasedRun`
- `unbiasedFunc` ["unbiasedRun"] which function to call for unbiased evaluation
- `tuneMethod` ["spot"] other choices: "cmaes", "bfgs", ..., see `tdmDispatchTuner`
- `nExperim` [1]
- `umode` ["RSUB"], one out of [ "RSUB" | "CV" | "TST" | "SP_T" ], see `unbiasedRun`
- `timeMode` [1] 1: proc time, 2: system time, 3: elapsed time (columns `timeTRN` in `envT$theFinals`
- `filenameEnvT` filename where `tdmBigLoop` will save a small version of environment `envT`. If NULL, it is set to `sub(".conf",".Rdata", tdm$runList[1])`
- `theSpotPath` [NA] use SPOT's package version
- `parallelCPUs` [1] 1: sequential, >1: parallel execution with this many CPUs (package `parallel`)
parallelFuncs  [NULL] in case tdm$parallelCPUs>1: a string vector with functions which are
clusterExport'ed in addition to tdm$mainFunc.

path        [NULL] from where to load and save envT resp. filenameEnvT. If it is NULL, tdm$path is set to the actual working directory at the time when tdmEnvTMak-

runList     [NULL] a list of configuration names .conf

stratified  [NULL] see tdmReadAndSplit

tdmPath     [NULL] from where to source the R sources. If NULL load library TDMR

run         ["default cutoff"]

withParams  [0] the verbosity for the unbiased runs

withParams  [TRUE] list the columns with tuned parameter in final results

nrun        [5] number of runs for unbiased run

U.saveModel [TRUE] if TRUE, save the last model, which is trained in unbiasedRun, onto filenameEnvT

tstCol      ["TST.COL"] opts$TST.COL for unbiased runs (only for umode="TST")
	nfold       [10] number of CV-folds for unbiased runs (only for umode="CV")

TST.trnFrac  [NULL] train set fraction (of all train-vali data), OVERWRITES opts$TST.trnFrac if not NULL.

TST.valiFrac [NULL] validation set fraction (of all train-vali data), OVERWRITES to opts$TST.valiFrac if not NULL.

TST.testFrac [0.2] test set fraction (of *all* data) for unbiased runs (only for umode="RSUB" or "SP_T")

CMA.propertyFile [NULL] (only for CMA-ES Java tuner) see cma_jTuner.

CMA.populationSize [NULL] (only for CMA-ES Java tuner) see cma_jTuner.

Note

The settings tdm$TST.trnFrac and tdm$TST.valiFrac allow to set programmatically certain values for opts$TST.trnFrac and opts$TST.valiFrac after opts has been constructed. So use tdm$TST.trnFrac and tdm$TST.valiFrac with CAUTION!

For tdm$timeMode, the ‘user time’ is the CPU time charged for the execution of user instructions of the calling process. The ‘system time’ is the CPU time charged for execution by the system on behalf of the calling process. The ‘elapsed time’ is the ‘real’ (wall-clock) time since the process was started.

Author(s)

Wolfgang Konen, THK, Patrick Koch
tdmEnvTAddBstRes

Add BST and RES data frames to an existing envT environment.

Description

Load an envT-type environment from file fileRData. Its elements bst, bstGrid res, and resGrid overwrite the elements in envT passed in as argument.

Usage

tdmEnvTAddBstRes(envT, fileRData)

Arguments

envT the TDMR environment
fileRData string with filename to load. This file is searched in envT$tdm$path.

Value

the augmented envT

tdmEnvTAddGetters

Add getter functions getBst and getRes to environment envT

Description

Add getter functions getBst and getRes to environment envT

Usage

tdmEnvTAddGetters(envT)

Arguments

envT the TDMR environment

Value

the augmented envT
**tdmEnvTGetOpts**

Return list opts from the k-th element of envT$sCList

**Usage**

tdmEnvTGetOpts(envT, k = 1)

**Arguments**

- **envT**: environment TDMR
- **k**: [1] index 1,...,length(envT$runList)

**Value**

- **opts**

---

**tdmEnvTLoad**

Load an envT-type environment from file fileRData.

**Description**

The loaded envT is augmented with getter functions, see tdmEnvTAddGetters.

**Usage**

tdmEnvTLoad(fileRData, path = NULL)

**Arguments**

- **fileRData**: string with filename to load.
- **path**: [NULL] dir where to search fileRData. If NULL, use current dir.

**Value**

- **envT**
Construct a new environment envT of class TDMenvir.

Description

Given the general TDMR settings in tdm, construct an appropriate environment envT. This is needed as input for tdmBigLoop.

Usage

tdmEnvTMakeNew(tdm = NULL, sList = defaultSList())

Arguments

dtm a list with general settings for TDMR, see tdmDefaultsFill

sList [defaultSCList()] a list of list with controls for SPOT or other tuners (one list for each element in tdm$runList)

Value

Environment envT, an object of class TDMenvir, containing (among others) the elements

runList = tdm$runList
tdm = tdmDefaultsFill(tdm)
getBst accessor function(confFile,nExp,theTuner) into envT$bstGrid
getRes accessor function(confFile,nExp,theTuner) into envT$resGrid
sList list of spotConfig-objects, as many as envT$runList has elements. Each spotConfig object sList[[k]] contains a list opts as element for the machine learning part.

Author(s)

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), THK, Patrick Koch

See Also

tdmBigLoop
**tdmEnvTReport**

Make a report plot based on envT

### Description

Given the results from a prior tuning run in envT, make a sensitivity plot for this run. If `envT$tdm$nrun > 0` then make additionally with the best-performing parameters from the tuning run a new unbiased run on the test data.

### Usage

```r
tdmEnvTReport(envT, ind)
```

### Arguments

- **envT** results from a prior tuning run.
- **ind** an integer from 1:length(envT$bstGrid): Take the tuning run with index ind.

### Value

`envT`, with data frame `finals` added, if `envT$tdm$nrun > 0`.

### See Also

- `tdmEnvTReportSens`

### Examples

```r
# The best results are read from demo02sonar/demoSonar.RData relative to the TDMR package directory.
path = paste(find.package("TDMR"), "demo02sonar", sep="/");
envT = tdmEnvTLoad("demoSonar.RData", path); # loads envT source(paste(path,"main_sonar.r",sep="/"));
envT$tdm$nrun=0; # =0: don't, >0: do unbiasedRun with opts$NRun=envT$tdm$nrun envT$slList[[1]]$opts$VERBOSE=1;
envT <- tdmEnvTReport(envT,1);
if (!is.null(envT$theFinals)) print(envT$theFinals);
```
tdmEnvTReportSens  
*Function to generate a report with sensitivity plot.*

**Description**

The sensitivity curves are based on a metamodel which is a random forest with 100 trees fitted to the result points from RES-file. The plot contains: x-axis: ROI for each parameter normalized to [-1,1] y-axis:

**Usage**

```
tdmEnvTReportSens(spotConfig)
```

**Arguments**

- `spotConfig`: the configuration list of all spot parameters

**See Also**

- `tdmEnvTReport`

---

### tdmEnvTSetOpts

Set list opts for the k-th element of `envT$sCList`

**Description**

Set list opts for the k-th element of `envT$sCList`

**Usage**

```
tdmEnvTSetOpts(envT, opts, k = 1)
```

**Arguments**

- `envT`: environment TDMR
- `opts`: list of options
- `k`: [1] index 1,...,length(envT$runList)

**Value**

`envT`
**tdmEnvTUpdate**

*Update env$tdm*

**Description**
Update env$tdm with the non-NULL elements of tdm

**Usage**
```r
tdmEnvTUpdate(envT, tdm)
```

**Arguments**

- `envT`  
  environment TDMR
- `tdm`  
  list for TDMR, see `tdmDefaultsFill`

**Value**

- `envT`

---

**tdmGraAndLogFinalize**

*Finalize graphics and log file*

**Description**
Finalize graphics and log file

**Usage**
```r
tdmGraAndLogFinalize(opts, gdObj = NULL)
```

**Arguments**

- `opts`  
  with opts$GD.DEVICE one out of [ "pdf" | "png" | "win" | "rstudio" | "non" ], see `tdmGraphicInit`
- `gdObj`  
  object of class TDMgdev, the return value from tdmGraAndLogInitialize, to ensure that tdmGraAndLogInitialize was called before (and the sink on opts$LOGFILE can be closed)
tdmGraAndLogInitialize

Initialize graphics and log file.

Description

The log file is opened in opts$dir.output/opts$LOGFILE, but only if opts$fileMode==TRUE.

Usage

tdmGraAndLogInitialize(opts)

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>opts</td>
<td>with opts$GD.DEVICE one out of [ &quot;pdf&quot;</td>
</tr>
</tbody>
</table>

Value

gdObj, an object of class TDMgdev. Pass this object on when calling tdmGraAndLogFinalize(opts,gdObj) (if not, a warning is issued before the sink-closing-error occurs)

---

tdmGraphicCloseDev

Close all open graphic devices.

Description

Close all open graphic devices.

Usage

tdmGraphicCloseDev(opts, ...)

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>opts</td>
<td>with opts$GD.DEVICE one out of [ &quot;pdf&quot;</td>
</tr>
<tr>
<td>...</td>
<td>optional arguments (currently not used)</td>
</tr>
</tbody>
</table>
tdmGraphicCloseWin  
*Close active file ("png").*

**Description**

Close active file ("png").

**Usage**

```r
tdmGraphicCloseWin(opts, ...)
```

**Arguments**

- `opts` with `opts$GD.DEVICE` one out of ["pdf" | "png" | "win" | "rstudio" | "non"], see `tdmGraphicInit`
- `...` optional arguments (currently not used)

---

**tdmGraphicInit**  
*Initialize graphic device.*

**Description**

Open multipage PDF or (create and) clear `opts$GD.PNGDIR`.

**Usage**

```r
tdmGraphicInit(opts, ...)
```

**Arguments**

- `opts` with `opts$GD.DEVICE` one out of ["pdf" | "png" | "win" | "rstudio" | "non"]
  - "pdf" plot everything in one multipage pdf file `opts$PDFFILE`
  - "png" each plot goes into a new png file in `opts$GD.PNGDIR`
  - "win" each plot goes into a new window (dev.new())
  - "rstudio" plot everything to the RStudio plot device (has a history)
  - "non" all plots are suppressed
- `...` optional arguments to hand over to `pdf` (the other devices require no further arguments)
**Description**

Initialize a new window ("win") / a new file ("png") for current graphic device.

**Usage**

```r
tdmGraphicNewWin(opts, ...)
```

**Arguments**

- `opts` with `opts$GD.DEVICE` one out of ["pdf" | "png" | "win" | "rstudio" | "non"], see `tdmGraphicInit`
- `...` optional arguments to hand over to `png` or `windows` or `x11` in package `grDevices` (the other devices require no further arguments)

---

**Description**

Only relevant for `opts$GD.DEVICE=="win"`.  

**Usage**

```r
tdmGraphicToTop(opts)
```

**Arguments**

- `opts` with `opts$GD.DEVICE` one out of ["pdf" | "png" | "win" | "rstudio" | "non"], see `tdmGraphicInit`
tdmMapDesApply 

Apply the mapping from des to opts.

Description

For each variable which appears in .roi (and thus in design point data frame des): set its counterpart in list opts to the values of the k-th row in des. For each variable not appearing: leave its counterpart in opts at its default value from defaultOpt.

Usage

tdmMapDesApply(des, opts, k, spotConfig, tdm)

Arguments

des design points data frame
opts list of options
k apply mapping for the k-th design point
spotConfig list, we needed here spotConfig$alg.ROI and envT$mapUser, see tdmMapDesLoad, and in addition envT$spotConfig$alg.ROI
tdm list, we need here tdm$map and tdm$mapUser

Value

opts, the modified list of options

See Also

tdmMapDesLoad

---

tdmMapDesLoad 

Load the mapping files.

Description

Load the map files "tdmMapDesign.csv" and optionally also "userMapDesign.csv" and store them in tdm$map and tdm$mapUser, resp. These maps are used by tdmMapDesApply.
"tdmMapDesign.csv" is searched in the TDMR library path find.package("TDMR"). (For the developer version: <studPath>/inst).
"userMapDesign.csv" is searched in tdm$path (which is getwd() if the user did not define tdm$path).

Usage

tdmMapDesLoad(tdm = list())
Arguments

dtm list, needed for dtm$tdmPath and dtm$path

Value

dtm, the modified list with new elements dtm$map and dtm$mapUser

See Also

tdmMapDesApply

tdmModConfmat

Calculate confusion matrix, gain and RGain measure.

Description

Calculate confusion matrix, gain and RGain measure.

Usage

tdmModConfmat(d, colreal, colpred, opts, predprob = NULL)

Arguments

d data frame

colreal name of column in d which contains the real class

colpred name of column in d which contains the predicted class

opts a list from which we use the elements:
  • gainmat: the gain matrix for each possible outcome, same size as cm$mat (see below).
  gainmat[R1,P2] is the gain associated with a record of real class R1 which we predict as class P2. (gain matrix = - cost matrix)
  • rgain.type: one out of {"rgain" | "meanCA" | "minCA" | "bYouden" | "arROC" | "arLIFT" | "arPRE" }, affects output cm$mat and cm$rgain, see below.

predProb if not NULL, a data frame with as many rows as data frame d, containing columns (index, true label, predicted label, prediction score). Is only needed for opts$rgain.type="ar*".
Value

- **cm**, a list containing:
  - **mat**: matrix with real class levels as rows, predicted class levels columns.  
    mat[R1,P2] is the number of records with real class R1 predicted as class P2, if opts$rgain.type=="rgain". If opts$rgain.type=="meanCA" or "minCA", then show this number as percentage of "records with real class R1" (percentage of each row). CAUTION: If there are NA's in column colpred, those cases are missing in mat (!) (but the class errors are correct as long as there are no NA's in column colreal).
  - **cerr**: class error rates, vector of size nlevels(colreal)+1. cerr[X] is the misclassification rate for real class X. cerr["Total"] is the total classification error rate.
  - **gain**: the total gain (sum of pointwise product opts$gainmat*cm$mat)
  - **gain.vector**: gain.vector[X] is the gain attributed to real class label X. gain.vector["Total"] is again the total gain.
  - **gainmax**: the maximum achievable gain, assuming perfect prediction
  - **rgain**: Depending on the value of opts$rgain.type:
    - "rgain": ratio gain/gainmax in percent,
    - "meanCA": mean class accuracy percentage (i.e. mean(diag(cm$mat)),
    - "minCA": min class accuracy percentage (i.e. min(diag(cm$mat)),
    - "bYouden": balanced Youden index: min(sensitivity,specificity),
    - "arROC": area under ROC curve (a number in [0,1]),
    - "arLIFT": area between lift curve and horizontal line 1.0,
    - "arPRE": area under precision-recall curve (a number in [0,1])

Note

For all measures rgain holds: The higher, the better.
The last four elements of opts$rgain.type = "bYouden", "arROC", "arLIFT", "arPre" are only available for binary classification.
For case "bYouden":
sensitivity = TP/(TP+FN)
specificity = TN/(TN+FP)

Author(s)

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), Patrick Koch

See Also

tdmClassify tdmROCRbase
Create and return a training-validation-set index vector.

Description

Depending on the value of member TST.kind in list opts, the returned index cvi is

1. TST.kind="cv": a random cross validation index P([111...222...333...]) - or -
2. TST.kind="rand": a random index with P([00...11...-1-1...]) for training (0), validation (1) and
disregard (-1) cases - or -
3. TST.kind="col": the column dset[,opts$TST.COL] contains the training (0), validation (1) and
disregard (-1) set division (and all records with a value <0 in column TST.COL are disre-
garded).

Here P(.) denotes random permutation of the sequence.
The disregard set is optional, i.e. cvi may contain only 0 and 1, if desired.
Special case TST.kind="cv" and TST.NFOLD=1: make *every* record a training record, i.e. index
[000...].
In case TST.kind="rand" and stratified=TRUE a stratified sample is drawn, where the strata in the
training case reflect the rel. frequency of each level of the *1st* response variable and are ensured
to be at least of size 1.
In summary, TST.kind="cv" means cross validation (TST.NFOLD models are built with TST.NFOLD
different train-validation data sets), while TST.kind="rand" or "col" means one model build with a
random ("rand") or user-defined ("col") training-validation split.

Usage

tdmModCreateCVindex(dset, response.variables, opts, stratified = FALSE)

Arguments

dset the data frame for which cvi is needed
response.variables issue a warning if length(response.variables)>1. Use the first response
variable for determining strata size.
opts a list from which we need here the following entries
  • TST.kind: ["cv","rand","col"]
  • TST.NFOLD: number of CV folds (only relevant in case TST.kind="cv")
  • TST.COL: column of dset containing the (0/1/<0) index (only relevant in
case TST.kind="col") or NULL if no such column exists
  • TST.valiFrac: fraction of records to set aside for validation (only relevant
in case TST.kind="rand")
  • TST.trnFrac: [1-opts$TST.valiFrac] fraction of records to use for training
  (only relevant in case TST.kind="rand")
stratified [F] do stratified sampling for TST.kind="rand" with at least one training record
for each response variable level (classification)
tdmModSortedRFimport

Value

cvi training-validation-set (0/>0) index vector (all records with cvi<0, e.g. from column TST.COL, are disregarded)

Note

Currently stratified sampling in case TST.KIND='rand' does only work correctly for one response variable. If there are more than one, the right fraction of validation records is taken, but the strata are drawn w.r.t. the first response variable. (For multiple response variables we would have to return a list of cvi’s or to call tdmModCreateCVindex for each response variable anew.)

tdmModSortedRFimport   Sort the input variables decreasingly by their RF-importance.

Description

Build a Random Forest using importance=TRUE. Usually the RF is smaller (50 trees), to speed up computation. Use na.roughfix for missing value replacement. Decide which input variables to keep and return them in SRF$input.variables

Usage

tdmModSortedRFimport(d_train, response.variable, input.variables, opts)

Arguments

d_train      training set
response.variable
             the target column from d_train to use for the RF-model
input.variables
             the input columns from d_train to use for the RF-model
opts         options, here we use the elements [defaults in brackets]:

  • SRF.kind:
    =$xperc$: keep a certain importance percentage, starting from the most important variable
    =$ndrop$: drop a certain number of least important variables
    =$nkeep$: keep a certain number of most important variables
    =$none$: do not call tdmModSortedRFimport at all (see tdmRegress.r and tdmClassify.r)
  • SRF.ndrop: [0] how many variables to drop (if SRF.kind=="ndrop")
  • SRF.XPerc: [0.95] if >=0, keep that importance percentage, starting with the most important variables (if SRF.kind=="xperc")
  • SRF.calc: [TRUE] =TRUE: calculate importance & save on SRF.file, =F: load from SRF.file (SRF.file = Output/<filename>.SRF.<response.variable>.Rdata)
  • SRF.ntree: [50] number of RF trees
• SRF.verbose: [2]
• SRF.maxS: [40] how many variables to show in plot
• SRF.minls: [1] a lower bound for the length of SRF$input.variables
• RF.sampsze: sampsze for RF, set prior to calling this func via tdmModAd-
  justSampsze(opts$SRF.samps,)
• GD.DEVICE: if !="non", then make a bar plot on current graphic device
• CLS.CLASSWT: class weight vector to use in random forest training

Value

SRF, a list with the following elements:

input.variales
the vector of input variables which remain after importance processing. These
are sorted by decreasing importance.
s_input
all input.variales sorted by decreasing (**NEW** importance)
s_imp1
the importance for s_input
s_dropped
vector with name of dropped variables
lsd
length of s_dropped
perc
the percentage of total importance which is in the dropped variables
opts
some defaults might have been added

Author(s)

Wolfgang Konen, Patrick Koch <wolfgang.konen@th-koeln.de>

---

**tdmModVote2Target**

*Analyze how the vote fraction corresponds to reliability of prediction.*

Description

This function analyzes whether in different vote bins the trained RF makes predictions with differ-
ent reliability. Only for RF-prediction in case of binary (0/1) classification.

Expected result: The larger the fraction of trees voting for class 0 is, the smaller is the percentage of
true class-1- cases in this vote bin. This function is somewhat specialized for the DMC2010-task.

Usage

`tdmModVote2Target(vote0, pred, target)`

Arguments

vote0 vector: which fraction of trees votes for class 0?
pred vector: the predicted class for each record (0/1)
target vector: the true class for each vector (0/1)
Value

a data frame with columns

vcut vote cut v

count number of cases with vote fraction in [v[i-1], v[i]]

pred0 fraction of 0-predictions

pCorr fraction of correct predictions

pR fraction of true 1-cases

Author(s)

Wolfgang Konen <wolfgang.konen@th-koeln.de>

---

tdmOptsDefaultsSet    Default values for list opts.

---

Description

Set up and return a list opts with default settings. The list opts contains all DM-related settings which are needed by main_<TASK>.

For better readability, most elements of opts are arranged in groups:

- dir.* path-related settings
- READ.* data-reading-related settings
- TST.* resampling-related settings (training, validation and test set, CV)
- PRE.* preprocessing parameters
- SRF.* several parameters for tdmModSortedRFimport
- MOD.* general settings for models and model building
- RF.* several parameters for model RF (Random Forest)
- SVM.* several parameters for model SVM (Support Vector Machines)
- ADA.* several parameters for model ADA (AdaBoost)
- CLS.* classification-related settings
- GD.* settings for the graphic devices

Usage

tdmOptsDefaultsSet(opts = NULL, path = ".")

Arguments

opts (optional) the options already set

path ["." where to find everything for the DM task.
Details

The path-related settings are relative to opts$path, if it is def’d, else relative to the current dir. Finally, the function `tdmOptDefaultsFill(opts)` is called to fill in further details, depending on the current settings of opts.

Value

a list opts, with defaults set for all options relevant for a DM task, containing the following elements

- **path** ["." ] where to find everything for the DM task
- **dir.txt** [data] where to find .txt/.csv files
- **dir.data** [data] where to find other data files, including .Rdata
- **dir.output** [Output] where to put output files
- **filename** ["default.txt"] the task data
- **filetest** [NULL] the test data, only relevant for READ.TstFn!=NULL
- **data.title** ["Default Data"] title for plots
- **READ.TXT** [T] =T: read data from .csv and save as .Rdata, =F: read from .Rdata
- **READ.NROW** [-1] read this amount of rows or -1 for 'read all rows'
- **READ.TrnFn** function to be passed into `tdmReadDataset`. Signature: function(opts) returning a data frame. It reads the train-validation data.
- **READ.TstFn** [NULL] function to be passed into `tdmReadDataset`. Signature: function(opts) returning a data frame. It reads a separate test data file. If NULL, this reading step is skipped.
- **READ.INI** [TRUE] read the task data initially, i.e. prior to tuning, using `tdmReadDataset`. If =FALSE, the data are read anew in each pass through main_TASK, i.e. in each tuning step (deprecated).
- **TST.kind** ["rand"] one of the choices from {"cv","rand","col"}, see `tdmModCreateCVindex` for details
- **TST.COL** ["TST.COL"] name of column with train/test/disregard-flag
- **TST.NFOLD** [3] number of CV-folds (only for TST.kind=="cv")
- **TST.valiFrac** [0.1] set this fraction of the train-validation data aside for validation (only for TST.kind=="rand")
- **TST.testFrac** [0.1] set prior to tuning this fraction of data aside for testing (if tdm$umode=="SP_T" and opts$READ.INI==TRUE) or set this fraction of data aside for testing after tuning (if tdm$umode=="RSUB" or =="CV")
- **TST.trnFrac** [NULL] train set fraction, if NULL then `tdmModCreateCVindex` will set it to 1 - opts$TST.valiFrac.
- **TST.SEED** [NULL] a seed for the random test set selection (`tdmRandomSeed`) and random validation set selection. (`tdmClassifyLoop`). If NULL, use `tdmRandomSeed`.
- **PRE.PCA** ["none" (default)|"linear"] PCA preprocessing: [don’t do normal PCA (prcomp)]
PRE.PCA REPLACE
[T] = T: replace with the PCA columns the original numerical columns, = F: add the PCA columns

PRE.PCA.npc
[0] if > 0: add monomials of degree 2 from the first PRE.PCA.npc columns (PCs) (only active, if opts$PRE.PCA!="none")

PRE.SFA
["none" (default)|"2nd"] SFA preprocessing (see package rsfa: [don’t | do normal SFA with 2nd degree expansion ]

PRE.SFA REPLACE
[F] = T: replace the original numerical columns with the SFA columns; = F: add the SFA columns

PRE.SFA.npc
[0] if > 0: add monomials of degree 2 from the first PRE.SFA.npc columns (only active, if opts$PRE.SFA!="none")

PRE.SFA.PPRANGE
[11] number of inputs after SFA preprocessing, only those inputs enter into SFA expansion

PRE.SFA.ODIM
[5] number of SFA output dimensions (slowest signals) to return

PRE.SFA.doPB
[T] = T: don’t | do parametric bootstrap for SFA in case of marginal training data

PRE.SFA.fctPB
[sfaPBootstrap] the function to call in case of parametric bootstrap, see sfaPBootstrap in package rsfa for its interface description

PRE.allNonVali
[F] if = T, then use all non-validation data in the training-validation set for PCA or SFA preprocessing. If = F, use only the training set for PCA or SFA processing (only relevant if opts$PRE.PCA!="none" or opts$PRE.SFA!="none").

PRE.Xpgroup
[0.99] bind the fraction 1-PRE.Xpgroup in column OTHER (see tdmPreGroupLevels)

PRE.MaxLevel
[32] bind the N-32+1 least frequent cases in column OTHER (see tdmPreGroupLevels)

SRF.kind
["xperc" (default)|"ndrop"|"nkeep"|"none"] the method used for feature selection, see tdmModSortedRfImport

SRF.ndrop
[0] how many variables to drop (only relevant if SRF.kind=="ndrop")

SRF.nkeep
[NULL] how many variables to keep, NULL="keep all" (only relevant if SRF.kind=="nkeep")

SRF.XPerc
[0.95] if >= 0, keep that importance percentage, starting with the most important variables (if SRF.kind=="xperc")

SRF.calc

SRF.ntree
[50] number of RF trees

SRF.samp
sampsize for RF in importance estimation. See RF.samp for further info on sampsize.

SRF.verbose
[2]

SRF.maxS
[40] how many variables to show in plot

SRF.minlsi
[1] a lower bound for the length of SRF$input.variables

SRF.method
["RFimp"]

SRF.scale
[TRUE] option ‘scale’ for call importance() in tdmModSortedRfImport
MOD. SEED [NULL] a seed for the random model initialization (if model is non-deterministic). If NULL, use `tdmRandomSeed`.

MOD.method ['"RF" (default) |"MC.RF" |"SVM" |"NB" ]: use [RF | MetaCost-RF | SVM | Naive Bayes] in `tdmClassify`
["RF" (default) |"SVM" |"LM" ]: use [RF | SVM | linear model] in `tdmRegress`

RF.ntree [500]

RF.samp [1000] sampsize for RF in model training. If RF.samp is a scalar, then it specifies the total size of the sample. For classification, it can also be a vector of length n.class (= # of levels in response variable), then it specifies the size of each strata. The sum of the vector is the total sample size. If NULL, RF.samp will be replaced by 3000 later in `tdmModAdjustSampsize`.

RF.mtry [NULL]

RF.nodesize [1]

RF.OOB [TRUE] if =T, return OOB-training set error as tuning measure; if =F, return validation set error

RF.p.all [FALSE]


SVM.epsilon [0.005] needed only for regression

SVM.gamma [0.005]

SVM.coef0 [0.0] (needed only for opts$SVM.kernel=="polynomial" or =="sigmoid")

SVM.degree [3] (needed only for opts$SVM.kernel=="polynomial")

SVM.tolerance [0.008]


ADA.mfinal [10] number of trees in AdaBoost = mfinal boosting(...,mfinal,...)

ADA.rpart.msplit [20] minimum number of observations in a node in order for a split to be attempted

CLS.cutoff [NULL] vote fractions for the classes (vector of length n.class = # of levels in response variable). The class i with maximum ratio (% votes)/CLS.cutoff[i] wins. If NULL, then each class gets the cutoff 1/n.class (i.e. majority vote wins). The smaller CLS.cutoff[i], the more likely class i will win.

CLS.CLASSWT [NULL] class weights for the n.class classes, e.g. c(A=10,B=20) for a 2-class problem with classes A and B (the higher, the more costly is a misclassification of that real class). It should be a named vector with the same length and names as the levels of the response variable. If no names are given, the levels of the response variables in lexicographical order will be attached in `tdmClassify`. CLS.CLASSWT=NULL for no weights.

CLS.gainmat [NULL] (n.class x n.class) gain matrix. If NULL, CLS.gainmat will be set to unit matrix in `tdmClassify`
rgain.type  ["rgain" (default) ] in case of tdmClassify: For classification, the measure Rgain returned from tdmClassifyLoop in result$R_* is [relative gain (i.e. gain/gainmax) | mean class accuracy | minimum class accuracy | minus Y]. The goal is to maximize Rgain. For binary classification there are the additional measures [ "arROC" | "arLIFT" | "arPRE" | "bYouden" ], see 'Value' in tdmModConfmat. For regression, the goal is to minimize result$R_* returned from tdmRegress. In this case, possible values are rgain.type = ["rmae" (default) |"rmse" |"made" ] which stands for [ relative mean absolute error | root mean squared error | mean absolute deviation ].

cmpcopies  [0] if >0, activate tdmParaBootstrap in tdmClassify
cfct.postproc [NULL] name of a function with signature (pred, dframe, opts) where pred is the prediction of the model on the data frame dframe and opts is this list. This function may do some postprocessing on pred and it returns a (potentially modified) pred. This function will be called in tdmClassify if it is not NULL.

GD.DEVICE  ["win"] ="win": all graphics to (several) windows (windows or X11 in package grDevices)
= "rstudio": same as "win", but all graphics go to the RStudio device
= "pdf": all graphics to one multi-page PDF
= "png": all graphics in separate PNG files in opts$GD.PNGDIR
= "non": no graphics at all
This concerns the TDMR graphics, not the SPOT (or other tuner) graphics. If running R from RStudio (if there is a device with name "RStudioGD") then the default "win" is changed to "rstudio" automatically.

GD.RESTART  [T] =T: restart the graphics device (i.e. close all 'old' windows or re-open multi-page pdf) in each call to tdmClassify or tdmRegress, resp.
= F: leave all windows open (suitable for calls from SPOT) or write more pages in same pdf.

GD.CLOSE  [T] =T: close graphics device "png", "pdf" at the end of main_*.r (suitable for main_*.r solo) or
= F: do not close (suitable for call from tdmStartSpot2, where all windows should remain open)

NRUN  [2] how many runs with different train & test samples - or - how many CV-runs, if opts$TST.kind="cv"

APPLY_TIME  [FALSE]
test2.show  [FALSE]
test2.string  ["default cutoff"]
VERBOSE  [2] =2: print much output, =1: less, =0: none

Note
The variables opts$PRE.PCA.numericV and opts$PRE.SFA.numericV (string vectors of numeric input columns to be used for PCA or SFA) are not set by tdmOptsDefaultsSet or tdmOptsDefaultsFill. Either they are supplied by the user or, if NULL, TDMR will set them to input.variables in tdmClassifyLoop, assuming that all columns are numeric.
tdmParaBootstrap

**Author(s)**

Wolfgang Konen, THK, 2013 - 2018

**See Also**

tdmOptsDefaultsFill tdmDefaultsFill

---

**tdmParaBootstrap**  
*Parametric bootstrap: add 'noisy copies' to a data frame (training data).*

**Description**

A normal distribution is approximated from the data given in `dset[,input.variables]` and new data are drawn from this distribution for the columns `input.variables`. The column `resp` is filled at random with levels with the same relative frequency as in `dset[,resp]`. Other columns of `dset` are filled by copying the entries from the first row of `dset`.

**Usage**

```r
tdmParaBootstrap(dset, resp, input.variables, opts)
```

**Arguments**

- `dset`  
  data frame with training set
- `resp`  
  name of column in `dset` which carries the target variable
- `input.variables`  
  vector with names of input columns
- `opts`  
  additional parameters [defaults in brackets]
  - `ncopies` how many noisy copies to add
  - `ncsigma` [1.0] multiplicative factor for each std.dev.
  - `ncmethod` [3] which method to use for parametric bootstrap
    - =1: each 'old' record from X in turn is the centroid for a new pattern;
    - =2: the centroid is the average of all records from the same class, the std.dev. is the same for all classes;
    - =3: centroid as in '2', the std.dev. is the std.dev. of all records from the same class (*recommended*)
  - `TST.COL` (optional) name of column in `dset` where each PB record is marked with a 0

**Value**

data frame `dset` with the new parametric bootstrap records added as last rows.
**tdmPreAddMonomials**

**Author(s)**
Wolfgang Konen, FHK, Nov’2011-Dec’2011

**See Also**
tdmclassify
tdmpreaddmonomials

Add monomials of degree 2 to a data frame.

**Description**
Given the data frame `dset` and a data frame `rx` with the same number of rows, add monomials of degree 2 to `dset` for all quadratic combinations of the first `PRE.npc` columns of `rx`. The naming of these new columns is "R1x2" for the combination of cols 1 and 2 and so on (if prefix="R").

**Usage**
tdmPreAddMonomials(dset, rx, PRE.npc, opts, degree = 2, prefix = "R")

**Arguments**
- `dset`: the target data frame
- `rx`: a data frame where to draw the monomials from
- `PRE.npc`: the number of columns from `rx` to use (clipped to `ncol(rx)` if necessary)
- `opts`: a list from which we need here the following entries:
  - `filename`
  - `VERBOSE`
- `degree`: [2] (currently only 2 is supported)
- `prefix`: ["R"] character prefix for the monomial column names

**Value**
data frame `dset` with the new monomial columns appended. If `PRE.npc==0`, the data frame is returned unchanged.

**Note**
CAVEAT: The double for-loop costs some time (e.g. 2-4 sec for `ncol(rx)=8` or `10`) How to fix: make a version w/o for-loop and w/o frequent assigns to `dset` (**TODO***)
**tdmPreFindConstVar**

*Find constant columns.*

**Description**

Find all those columns in data frame dset which are completely constant or completely NA and return a vector with their names.

**Usage**

```r
tdmPreFindConstVar(dset)
```

**Arguments**

- `dset` data frame

**Value**

name vector of constant columns

---

**tdmPreGroupLevels**

*Group the levels of factor variable in dset[,colname].*

**Description**

This function reduces the number of levels for factor variables with too many levels. It counts the cases in each level and orders them decreasingly. It binds the least frequent levels together in a new level "OTHER" such that the remaining untouched levels have more than opts$PRE.Xpgroup percent of all cases. OR it binds the levels with least cases together in "OTHER" such that the total number of new levels is opts$PRE.MaxLevel. From these two choices for "OTHER" take the one which binds more variables in column "OTHER".

**Usage**

```r
tdmPreGroupLevels(dset, colname, opts)
```

**Arguments**

- `dset` data frame
- `colname` name of column to be re-grouped
- `opts` list, here we need
  - `PRE.Xpgroup [0.99]`
  - `PRE.MaxLevel [32]` (32 is the maximum number of levels allowed for randomForest)
Value

dset, a data frame with \texttt{dset[,colname]} re-grouped

\section*{tdmPreLevel2Target}

\textit{Relate levels of a column with a target (column).}

\section*{Description}

Print for each level of factor variable \( f \) which ratio 0 / 1 of the binary target variable it contains and how many cases are in each level.

\section*{Usage}

\texttt{tdmPreLevel2Target(dset, target, f, opts)}

\section*{Arguments}

\begin{itemize}
  \item \texttt{dset} data frame
  \item \texttt{target} name of target column
  \item \texttt{f} number of column with factor variable
  \item \texttt{opts} list, here we need
    \begin{itemize}
      \item \texttt{opts$thresh_pR}
      \item \texttt{opts$verbose}
    \end{itemize}
\end{itemize}

\section*{Note}

SIDE EFFECTS: some printed output

\section*{tdmPreNAroughfix}

\textit{Replace <NA> values with suitable non <NA> values}

\section*{Description}

This function replaces <NA> values in a list entry or data frame column with the median (for numeric columns) or the most frequent mode (for factor columns). It does the same as \texttt{na.roughfix} in package \texttt{randomForest}, but does so faster.

\section*{Usage}

\texttt{tdmPreNAroughfix(object, \ldots)}

\section*{Arguments}

\begin{itemize}
  \item \texttt{object} list or data frame
  \item \texttt{\ldots} additional arguments
\end{itemize}
Value

object, the list or data frame with <NA> values replaced

---

**tdmPrePCA.apply**  
*Apply PCA (Principal Component Analysis) to new data.*

Description

The PCA rotation is taken from `pcalist`, a value returned from a prior call to `tdmPrePCA.train`.

Usage

```
tdmPrePCA.apply(dset, pcaList, opts, dtrain = NULL)
```

Arguments

- `dset`  
  the data frame with the new data
- `pcaList`  
  a value returned from a prior call to `tdmPrePCA.train`
- `opts`  
  a list from which we need here the following entries:
  - `PRE.knum`: if >0 and if `PRE.PCA`="kernel", take only a subset of `PRE.knum` records from `dset`
  - `PRE.PCA.npc`: if >0, then add for the first `PRE.PCA.npc` PCs the monomials of degree 2 (see `tdmPreAddMonomials`)
  - `PRE.PCA.numericV` vector with all column names in `dset` for which PCA is performed. These columns may contain *numeric* values only.
- `dtrain`  
  [NULL] optional, only needed in case that `dset` is a 0-row-data frame: then we 'borrow' the columns from `dtrain`, the data set returned from `tdmPrePCA.train` in `pca$dsset`.

Value

```
pca, a list with entries:

dset the input data frame `dset` with columns `numeric.variables` replaced by the PCs with names PC1, PC2, ... (in case `PRE.PCA`="linear") or with names KP1, KP2, ... (in case `PRE.PCA`="kernel") and optional with monomial columns added, if `PRE.PCA.npc`>0
numeric.variables the new column names for PCs and for the monomials
```

Author(s)

Wolfgang Konen, FHK, Mar'2011 - Jan'2012

See Also

`tdmPrePCA.train`
**tdmPrePCA.train**

PCA (Principal Component Analysis) for numeric columns in a data frame.

**Description**

tdmPrePCA.train is capable of linear PCA, based on prcomp (which uses SVD), and of kernel PCA (either KPCA, KHA or KFA).

**Usage**

tdmPrePCA.train(dset, opts)

**Arguments**

dset the data frame with training (and test) data.

opts a list from which we need here the following entries:

- PRE.PCA: ["linear" | "kernel" | "none" ]
- PRE.knum: if >0 and if PRE.PCA="kernel", take only a subset of PRE.knum records from dset
- PRE.PCA.REPLACE: [T] =T: replace the original numerical columns with the PCA columns; =F: add the PCA columns
- PRE.PCA.npc: if >0, then add for the first PRE.PCA.npc PCs the monomials of degree 2 (see tdmPreAddMonomials)
- PRE.PCA.numericV vector with all column names in dset for which PCA is performed. These columns may contain *numeric* values only.

**Value**

pca, a list with entries:

- dset the input data frame dset with columns numeric.variables replaced or extended (depending on opts$PRE.PCA.REPLACE) by the PCs with names PC1, PC2, ... (in case PRE.PCA=="linear") or with names KP1, KP2, ... (in case PRE.PCA=="kernel") and optional with monomial columns added, if PRE.PCA.npc>0. The number of PCs is min(nrows(dset),length(numeric.variables)).
- numeric.variables the new numeric column names (PCs, monomials, and optionally old numericV, if opts$PRE.PCA.REPLACE==F)
- pcaList a list with the items sdev, rotation, center, scale, x as returned from prcomp plus eigval, the eigenvalues for the PCs

**Note**

CAUTION: Kernel PCA (opts$PRE.PCA=="kernel") is currently disabled in code, it *crashes* for large number of records or large number of columns.
Apply SFA (Slow Feature Analysis) to new data.

**Description**

The SFA projection is taken from `sfalist`, a value returned from a prior call to `tdmPreSFA.train`.

**Usage**

```r
tdmPreSFA.apply(dset, sfalist, opts, dtrain = NULL)
```

**Arguments**

- **dset**: the data frame with the new data
- **sfalist**: a value returned from a prior call to `tdmPreSFA.train`
- **opts**: a list from which we need here the following entries:
  - **PRE.SFA.REPLACE**: [T] = T: replace the original numerical columns with the SFA columns; = F: add the SFA columns
  - **PRE.SFA.npc**: if > 0, then add for the first PRE.SFA.npc PCs the monomials of degree 2 (see `tdmPreAddMonomials`)
  - **PRE.SFA.ODIM**: [5] number of SFA output dimensions (slowest signals) to return
  - **PRE.SFA.numericV**: vector with all column names in `dset` for which SFA is performed. These columns may contain *numeric* values only.
- **dtrain**: [NULL] optional, only needed in case that `dset` is a 0-row-data frame: then we ‘borrow’ the columns from `dtrain`, the data set returned from `tdmPreSFA.train` in `sfa$dset`.

**Value**

- **sfa**: a list with entries:
  - **dset**: the input data frame `dset` with columns `numeric.variables` replaced by the PCs with names `PC1, PC2, ...` (in case `PRE.SFA=="linear"`) or with names `KP1, KP2, ...` (in case `PRE.SFA=="kernel"`) and optional with monomial columns added, if `PRE.SFA.npc>0`
  - **numeric.variables**: the new column names for PCs and for the monomials
Author(s)
Wolfgang Konen, Martin Zaefferer, FHK, Jan’2012 - Feb’2012

See Also
tdmPreSFA.train

SFA (Slow Feature Analysis) for numeric columns in a data frame.

Description
tdmPreSFA.train uses package rSFA. It is assumed that classification for the variable contained in
column response.var is done. SFA seeks features in an expanded function space for which the
intra-class variation w.r.t. response.var is as low as possible.

Usage
tdmPreSFA.train(dset, response.var, opts)

Arguments
dset the data frame with training (and test) data.
response.var the response variable for classification.
opts a list from which we need here the following entries:
  • PRE.SFA: [ "linear" l "2nd" l "none" ] which stands for [ 1st l 2nd degree
    monomial SFA l no SFA ]
  • PRE.SFA.REPLACE: [T] =T: replace the original numerical columns with
    the SFA columns; =F: add the SFA columns
  • PRE.SFA.npc: if >0, then add for the first PRE.SFA.npc PCs the monomials
    of degree 2 (see tdmPreAddMonomials)
  • PRE.SFA.PPRANGE: [11] number of inputs after preprocessing, they enter
    into expansion
  • PRE.SFA.ODIM: [5] number of SFA output dimensions (slowest signals)
    to return
  • PRE.SFA.numericV vector with all column names in dset which are input
    for SFA. These columns may contain *numeric* values only.

Value
sfa, a list with entries:
dset the input data frame dset with columns numeric.variables replaced or extended
  (depending on opts$PRE.SFA.REPLACE) by the SFA components with names
  SF1, SF2, ... and with optional monomial columns added, if PRE.SFA.npc>0
numeric.variables
the new numeric column names of dset, i.e. SFA components, monomials (and optionally PRE.SFA.numericV, if opts$PRE.SFA.REPLACE==F)
sfaList a list with the items opts (sfa0opts), matrices DSF and SF and many others, as returned from sfaStep

Author(s)
Wolfgang Konen, Martin Zaefferer, FHK, Jan’2012 - Feb’2012

See Also

tdmPreSFA.apply

---

`tdmRandomSeed`  
Generates pseudo-random random number seeds.

**Description**
To use this mechanism, create first an object `tdmRandomSeed` with a call to `makeTdmRandomSeed`.

**Usage**
`tdmRandomSeed()`

**Value**
In each call to this function a different integer in 0...100001+nCall is returned. This is true even if it is called many times within the same second (where Sys.time() will return the same integer). nCall is the number of calls to this function object.

**Author(s)**
Wolfgang Konen, Patrick Koch <wolfgang.konen@th-koeln.de>

**See Also**

`makeTdmRandomSeed`

**Examples**

```r
tdmRandomSeed = makeTdmRandomSeed();
for (i in 1:10) print(c(as.integer(Sys.time()), tdmRandomSeed()));
```
**tdmReadAndSplit**

*Read and split the task data.*

**Description**

Read the task data using `tdmReadDataset` and split them into a test part and a training/validation-part and return a `TMDdata` object.

**Usage**

```r
tdmReadAndSplit(opts, tdm, nExp = 0, dset = NULL)
```

**Arguments**

- `opts` a list from which we need here the elements
  - `readINI` \[T\] =T: do read and split, =F: return NULL
  - `read*:` other settings for `tdmReadDataset`
  - `filename`: needed for `tdmReadDataset`
  - `filetest`: needed for `tdmReadDataset`
  - `TST.testFrac`: \[0.1\] set this fraction of the data aside for testing
  - `TST.COL`: string with name for the partitioning column, if `tdm$umode` is not "SP_T". (If `tdm$umode`="SP_T", then `TST.COL`="tdmSplit" is used.)
- `tdm` a list from which we need here the elements
  - `mainfile`: if not NULL, set working dir to `dir(mainfile)` before executing `tdmReadDataset`
  - `umode`: \[ "RSUB" | "CV" | "TST" | "SP_T" \], how to divide in training/validation data for tuning and test data for the unbiased runs
  - `SPLIT.SEED`: if NULL, set random number generator (RNG) to `tdmRandomSeed` when constructing. `dataObj`. If not NULL, set RNG to `SPLIT.SEED + nExp` -> deterministic test set split
  - `stratified`: \[NULL\] string specifying the column with the response variable for classification. If not NULL, do the split by stratified sampling (at least one record of each class level found in `dset[,tdm$stratified]` shall appear in the train-vali-set). Recommended for classification
- `nExp` \[0\] experiment counter, used to select a reproducible different seed, if `tdm$SPLIT.SEED`!=NULL
- `dset` \[NULL\] if non-NULL, reading of `dset` is skipped and the given data frame `dset` is used.

**Details**

If `dset` is NULL, the files specified in `opts` are read into `dset`, see `tdmReadDataset` for details. Then, depending on the value of `tdm$umode`

- "SP_T": split the data randomly into training and test data with test set fraction according to `opts$TST.testFrac`. Make use of `tdm$SPLIT.SEED` and `tdm$stratified`, if given. Set `TST.COL` to "tdmSplit".
• "RSUB", "CV": use all data for training/validation. That is, the training-validation split is done later in `tdmClassifyLoop` or `tdmRegressLoop`.

• "TST": split the data into training and test data according to column. opts$TST.COL (usually "TST.COL"), which carries a 1 for each test record and a 0 else. If opts$filetest is specified, then all records from this file will carry a 1 in opts$TST.COL. All records from opts$filename carry a 0.

Value
dataObj, either NULL (if opts$READ.INI==FALSE) or an object of class `TDMdata` containing

dset a data frame with the complete data set
TST.COL string, the name of the column in dset which has a 1 for records belonging to the test set and a 0 for train/vali records. If tdm$umode=="SP_T", then TST.COL="tdmSplit", else TST.COL=opts$TST.COL.
filename opts$filename, from where the data were read

Use the accessor functions `dsetTrnVa.TDMdata` and `dsetTest.TDMdata` to extract the train/vali and the test data, resp., from dataObj.

Known caller: `tdmBigLoop`

Author(s)
Wolfgang Konen (<wolfgang.konen@th-koeln.de>), THK

See Also
`dsetTrnVa.TDMdata`, `dsetTest.TDMdata`, `tdmReadDataset`, `tdmBigLoop`

Description
Read the data according to the settings opts$READ.* and opts$TST.COL (see Details).

Usage
`tdmReadDataset(opts)`
Arguments

- `opts` list of options, we need here
  - `READ.TrnFn`: [tdmReadTrain] function with argument `opts` for reading the training data and returning them in a data frame
  - `READ.NROW`: `-1` read only that many rows from `opts$filename`. `-1` for 'read all rows'.
  - `READ.TstFn`: [NULL] function with argument `opts` for reading the test data and returning them in a data frame. If NULL then skip test file reading.
  - `TST.COL`: `"TST.COL"` string, create a column with the name of this string in `dset`, which has 0 for training and 1 for test data
  - `path`: used by `READ.TrnFn` to locate file
  - `dir.data`: used by `READ.TrnFn` to locate file

Details

When `opts$READ.TstFn==NULL`, then only `opts$READ.TrnFn` is used.
When `opts$READ.TstFn!=NULL`, the following things happen in `tdmReadDataset`: Data are read from `opts$filename` and from `opts$filetest`. Both data sets are bound together, with a new column `opts$TST.COL` having '0' for the data from `opts$filename` and having '1' for the data from `opts$filetest`. The branch using `opts$TST.COL` is invoked either with `umode="TST"` in `unbiasedRun` or with `opts$TST.kind="col"` in `tdmModCreateCvIndex`.

Value

dset, a data frame with all data read

See Also

- `tdmReadAndSplit`

---

**tdmReadTaskData**

Read task data.

**Description**

Read and split task data (wrapper for `tdmReadAndSplit`).

**Usage**

`tdmReadTaskData(envT, tdm)`

**Arguments**

- `envT` environment TDMR
- `tdm` list with TDMR controls
tdmRegress

**Value**

dataObj, see tdmReadAndSplit

**See Also**

dsetTrnVa.TMDdata, dsetTest.TMDdata, tdmReadAndSplit

tdmRegress

*Core regression function of TDMR.*

**Description**

tdmRegress is called by tdmRegressLoop and returns an object of class tdmRegre.

It trains a model on training set \( d_{\text{train}} \) and evaluates it on test set \( d_{\text{test}} \). If this function is used for tuning, the test set \( d_{\text{test}} \) plays the role of a validation set.

**Usage**

tdmRegress(d_train, d_test, d_preproc, response.variables, input.variables,
           opts, tsetStr = c("Validation", "validation", ".vali"))

**Arguments**

d_train training set
d_test test set, same columns as training set
d_preproc data used for preprocessing. May be NULL, if no preprocessing is done (opts$PRE.SFA="none" and opts$PRE.PCA="none"). If preprocessing is done, then d_preproc is usually all non-validation data.
response.variables name of column which carries the target variable - or - vector of names specifying multiple target columns (these columns are not used during prediction, only for evaluation)
input.variables vector with names of input columns
opts additional parameters [defaults in brackets]
  SRF:* several parameters for sorted_rf_importance (see tdmModelingUtils.r)
  RF:* several parameters for RF (Random Forest, defaults are set, if omitted)
  SVM:* several parameters for SVM (Support Vector Machines, defaults are set, if omitted)
  filename
data.title
mod.method ["RF"] the main training method ["RF","SVM","LM"]: use [Random forest| SVM| linear model] for the main model
### tdmRegress

- **MOD.SEED** = NULL: set the RNG to system time as seed (different RF trainings)
- = any value: set the random number seed to this value (+i) to get reproducible random numbers. In this way, the model training part (RF, NNET, ...) gets always a fixed seed. (see also TST.SEED in tdmRegressLoop)
- **OUTUTRAFO** [NULL] string, apply a transformation to the output variable
- **fct.postproc** [NULL] name of a user-def’d function for postprocessing of predicted output
- **gr.log** = FALSE (def): make scatter plot as-is, = TRUE: transform output x with log(x+1) (x should be nonnegative)
- **GD.DEVICE** if != "non", then make a pairs-plot of the 5 most important variables and make a true-false bar plot
- **VERBOSE** [2] = 2: most printed output, = 1: less, = 0: no output

#### Value
- res, an object of class tdmRegre, this is a list containing
  - **d_train** training set + predicted class column(s)
  - **d_test** test set + predicted target output
  - **allRMAE** data frame with columns = (rmae.train, rmae.test, theil.train, theil.test, ...) and rows = response variables. Here Theil’s U is based on RMAE (relative mean absolute error).
  - **allRMSE** data frame with columns = (rmse.train, rmse.test, theil.train, theil.test, ...) and rows = response variables. Here Theil’s U is based on RMSE (root mean square error).
  - **lastModel** the last model built (e.g. the last Random Forest in the case of MOD.method=="RF")
  - **opts** parameter list from input, some default values might have been added

The item lastModel is specific for the *last* model (the one built for the last response variable in the last run and last fold)

#### Author(s)
Wolfgang Konen, FHK, Sep’2009 - Jun’2012

#### See Also
print.tdmRegre tdmRegressLoop tdmClassifyLoop

#### Examples

```r
### This example shows a simple data mining process (phase 1 of TDMR) for regression on dataset iris.
### The data mining process in tdmRegress calls randomForest as the prediction model.
### It is called for 2 response variables. Therefore, the data frames allRMAE and allRMSE have 2 rows.
```
tdmRegressLoop

Core regression double loop returning a TDMregressor object.

Description

tdmRegressLoop contains a double loop (opts$NRUN and CV-folds) and calls \texttt{tdmRegress}. It is called by all R-functions \texttt{main\_\_\_*}. It returns an object of class \texttt{TDMregressor}.

Usage

tdmRegressLoop(dset, response\_variables, input\_variables, opts, tset = NULL)

Arguments

dset the data frame for which cvi is needed
response\_variables name of column which carries the target variable - or - vector of names specifying multiple target columns (these columns are not used during prediction, only for training and for evaluating the predicted result)
input\_variables vector with names of input columns
opts a list from which we need here the following entries
NRUN number of runs (outer loop)
TST\_SEED = NULL: leave the random number seed as it is. = any value: set the random number seed to this value to get reproducible random numbers and thus reproducible training-test-set-selection. (only relevant in case TST\_kind=="cv" or "rand") (see also MOD\_SEED in \texttt{tdmClassify})
TST\_kind how to create cvi, handed over to \texttt{tdmModCreateCVindex}. If TST\_kind=="col", then cvi is taken from dset[,opts$TST\_col].

doobj <- tdmGraAndLogInitialize(opts); # init graphics and log file
data(iris)
response\_variables=c("Petal\_Length","Petal\_Width") # names, not data (!)
input\_variables=setdiff(names(iris),response\_variables)
 opts$r\_gain\_type="rmae"
 opts$NRUN=1

idx_train = sample(nrow(iris))[1:110]
d\_train=iris[idx\_train,]
d\_vali=iris[-idx\_train,]
res <- tdmRegress(d\_train,d\_vali,NULL,response\_variables,input\_variables,opts)

print(res$all\_RMAE)
print(res$all\_RMSE)
tdmRegressLoop

G0.RESTART [TRUE] =TRUE/FALSE: do/don’t restart graphic devices
GRAPHDENV ["non"] other ]

tset [NULL] If not NULL, this is the test data set. If NULL, we are in tuning and
the validation data set is build from dset according to the procedure prescribed
in opts$TST.*.

Value

result, an object of class TDMregressor, this is a list with results, containing

opts the res$opts from tdmRegress
lastRes last run, last fold: result from tdmRegress
R_train RMAE / RMSE on training set (vector of length NRUN), depending on opts$rgain.type=="rmae"
or "rmse"
S_train RMSE on training set (vector of length NRUN)
T_train Theil’s U for RMAE on training set (vector of length NRUN)
*_test — similar, with test set instead of training set —
Err a data frame with as many rows as opts$NRUN and columns = (rmae.trn, rmse.trn
made.trn, rmae.theil.trn, ntrn, rmae.tst, rmse.tst, made.tst, rmae.theil.tst, ntst)
predictions last run: data frame with dimensions [nrow(dset),length(response.variable)]. In
case of CV, all validation set predictions (for each record in dset), in other cases
mixed validation / train set predictions.
predictTest predictions on the test set tset (NULL if tset==NULL )

Author(s)

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), THK

See Also

tdmRegress, tdmClassifyLoop, tdmClassify

Examples

```r
### -------- demo/demo00-lregress.r --------
### This demo shows a simple data mining process (phase 1 of TMR) for regression on
### dataset iris.
### The data mining process in tdmRegressLoop calls randomForest as the prediction model.
### It is called opts$NRUN=2 times with different random train-validation set splits.
### Therefore data frame result$Err has 2 rows.
###
### opts=tdmOptDefaultsSet()
### # set all defaults for data mining process
gdObj <- tdmGraAndLogInitialize(opts); # init graphics and log file

data(iris)
response.variables="Petal.Length" # names, not data (!)
input.variables=setdiff(names(iris),"Petal.Length")
opts$rgain.type="rmae"
```
tdmRegressSummary

result = tdmRegressLoop(iris,response.variables,input.variables,opts)

print(result$Err)

tdmRegressSummary  Print summary output for result from tdmRegressLoop and add result$y.

Description

result$y is "OOB RMAE" on training set for methods RF or MC.RF. result$y is "RMAE" on test set (=validation set) for all other methods. result$y is the quantity which the tuner seeks to minimize.

Usage

tdmRegressSummary(result, opts, dset = NULL)

Arguments

result  return value from a prior call to tdmRegressLoop, an object of class TDMregressor.
opts    a list from which we need here the following entries
        NRUN number of runs (outer loop)
        method
        VERBOSE
        dset  [NULL] if !=NULL, attach it to result
dset    [NULL] if not NULL, add this data frame to the return value (may cost a lot of memory!)

Value

result, an object of class TDMregressor, with result$y, result$sd.y (and optionally also result$dset) added

Author(s)

Wolfgang Konen, FHK, Sep’2010 - Oct’2011

See Also

tdmRegress, tdmRegressLoop, tdmClassifySummary
**tdmROCR.TDMclassifier**

*Interactive plot of ROC, lift or other charts for a TDMclassifier object.*

**Description**

Brings up a twiddle user interface, where the user may select a part of the dataset ("training" or "validation"), a run number (if $\text{Opts}(x)[NRUN>1]$) and a type-of-chart, see tdmROCRbase for details. Using tdmROCRbase, the appropriate chart is plotted on the current graphics device.

**Usage**

```r
## S3 method for class 'TDMclassifier'
tdmROCR(x, ...)
```

**Arguments**

- `x` return value from a prior call to tdmClassifyLoop, an object of class TDMclassifier.
- `...` – currently not used –

**Value**

The area under the curve plotted most recently.

**Note**

Side effect: Ror each chart, calculate and print the area between the curve and the bottom line (y=1.0 for typ="lift", y=0.0 else).

**See Also**

tdmClassifyLoop tdmROCRbase

**Examples**

```r
## Not run:
path <- paste(find.package("TDM"), "demo02sonar", sep="/");
source(paste(path,"main_sonar.r",sep="/"));
result = main_sonar();
tdmROCR(result);

## End(Not run)
```
Single plot of ROC, lift or other chart for a TDMclassifier object.

Description

Single plot of ROC, lift or other chart for a TDMclassifier object.

Usage

tdmROCRbase(x, dataset = "validation", nRun = 1, typ = "ROC",
          noPlot = FALSE, ...)

Arguments

- **x**: return value from a prior call to tdmclassifyLoop, an object of class TDMclassifier.
- **dataset**: ["validation"] which part of the data to use, either "training" or "validation"
- **nRun**: [1] if x contains multiple runs, which run to show (1, ..., opts(x)$NRUN)
- **typ**: ["ROC"] which chart type, one out of ("ROC", "lift", "precRec") for (ROC, lift, precision-recall)-chart (see performance in package ROCR for more details):
  - "ROC": receiver operating curve, TPR vs. FPR, with \( TPR = TP / (TP + FN) = TP / P \) and \( FPR = FP / (FP + TN) = FP / N \) (true and false positive rate).
  - "lift": lift chart, LIFT vs. RPP, with \( LIFT = TPR / RPR \) with random positive rate \( RPR = P / (P + N) \) and \( RPP = (TP + FP) / (P + N) \) (rate of pos. predictions).
  - "precRec": precision-recall-chart, PREC vs. RECALL, with \( PREC = TP / (TP + FP) \) and \( RECALL = TP / P \) (same as TPR).
- **noPlot**: [FALSE] if TRUE, suppress the plot, return only the area under curve
- ... currently not used

Value

The area between the curve and the bottom line \( y=0.0 \) in the case of \( \text{typ} = \text{"ROC"} \) or \( \text{typ} = \text{"precRec"} \) or the area between the curve and the bottom line \( y=1.0 \) in the case of \( \text{typ} = \text{"lift"} \).

If object x does not contain a prediction score, a warning is issued and the return value is NULL.

See Also

tdmclassifyLoop tdmROCR.TDMclassifier

Examples

```R
### Run task SONAR with "area under ROC curve" as performance measure (rgain.type="arROC").
### Other settings are similar to demo01-1sonar.r (level 1 of TDMR).
### Finally, plot ROC curve for validation data set and
### plot lift chart for training data set
###`
path <- paste(find.package("TDMR"), "demo02sonar", sep="/");
#path <- paste("./inst", "demo02sonar", sep="/");

source(paste(path,"main_sonar.r",sep="/"));  # defines readTrnSonar

controlDM <- function() {
  #
  # settings for the DM process (former sonar_00.apd file):
  # (see ?tdmOptDefaultsSet for a complete list of all default settings
  # and many explanatory comments)
  #
  opts = list(path = path,
             dir.data = "data",               # relative to path
             filename = "sonar.txt",
             READ.TrnFn = readTrnSonar,      # defined in main_sonar.r
             data.title = "Sonar Data",
             NRUN = 1,
             rgain.type = "arROC",
             VERBOSE = 2
          );
  opts <- setParams(opts, defaultOpts(), keepNotMatching = TRUE);
}

opts <- controlDM();
result <- main_sonar(opts);

tdmGraphicNewWin(opts);
cat("Area under ROC-curve for validation data set: ",
    tdmROCRbase(result),"\n");  # side effect: plot ROC-curve
tdmGraphicNewWin(opts);
cat("Area under lift curve for training data set: ",
    tdmROCRbase(result,dataset="training",typ="lift"),"\n");  # side effect: plot lift chart

tdmTuneIt

Tuning and unbiased evaluation (single tuning).

Description

For the first configuration name .conf in tdm$runList call the first tuning algorithm in tdm$tuneMethod (via function tdmDispatchTuner). After tuning perform with the best parameters a run of tdm$unbiasedFunc (usually unbiasedRun). This experiment is repeated tdm$nExperim times.

Usage

tdmTuneIt(envT, dataObj)
Arguments

envT an environment containing on input at least the element tdm (a list with general settings for TDMR, see tdmDefaultsFill), which has at least the elements
tdm$runList list of configuration names .conf
tdm$tuneMethod the tuner
dataObj object of class TDMdata (constructed here with the help of tdmReadAndSplit).

Details

tdmTuneIt differs from tdmBigLoop in that it processes only one configuration .conf and that it has dataObj as a mandatory calling parameter. This simplifies the data flow and is thus less error-prone.

tdm refers to envT$tdm.
See Details in tdmBigLoop for the list of available tuners.

Value

environment envT, containing the results

res data frame with results from last tuning (one line for each call of tdmStart*)
bst data frame with the best-so-far results from last tuning (one line collected after each (SPO) step)
resGrid list with data frames res from all tuning runs. Use envT:getRes(envT,confFile,nExp,theTuner) to retrieve a specific res.
bstGrid list with data frames bst from all tuning runs. Use envT:getBst(envT,confFile,nExp,theTuner) to retrieve a specific bst.
theFinals data frame with one line for each triple (confFile,nExp,tuner). Each line contains summary information about the tuning run in the form:
confFile tuner nExp [params] NRUN NEVAL RGain.bst RGain.* sdR.*
where [params] is written depending on tdm$withParams.
NRUN is the number of unbiased evaluation runs.
NEVAL is the number of function evaluations (model builds) during tuning.
RGain denotes the relative gain on a certain data set: the actual gain achieved with the model divided by the maximum gain possible for the current cost matrix and the current data set. This is for classification tasks, in the case of regression each RGain.* is replaced by RMAE.*, the relative mean absolute error.
Each ‘sdR.’ denotes the standard deviation of the preceding RGain or RMAE.
RGain.bst is the best result during tuning obtained on the training-validation data. RGain.avg is the average result during tuning. The following pairs RGain.* sdR.* are the results of one or several unbiased evaluations on the test data where ‘*’ takes as many values as there are elements in tdm$mode (the possible values are explained in unbiasedRun).

result object of class TDMclassifier or TDMregressor. This is a list with results from tdm$mainFunc as called in the last unbiased evaluation using the best parameters
found during tuning. Use `print(envT$result)` to get more info on such an object of class `TDMclassifier`.

`tunerVal` is an object with the return value from the last tuning process. For every tuner, this is the list `spotConfig`, containing the SPOT settings plus the TDMR settings in elements `opts` and `tdm`. Every tuner extends this list by `tunerVal$alg.currentResult` and `tunerVal$alg.currentBest`, see `tdmDispatchTuner`. In addition, each tuning method might add specific elements to the list, see the description of each tuner.

Environment `envT` contains further elements, but they are only relevant for the internal operation of `tdmBigLoop` and its subfunctions.

**Note**

Side effects:

- A compressed version of `envT` is saved to file `tdm$filenameEnvT` (default: `<runList[1]>RData`), in current directory.

If `tdm$U.saveModel==TRUE`, then `envT$result$lastRes$lastModel` (the last trained model) will be saved to `tdm$filenameEnvT`. The default is `tdm$U.saveModel==TRUE` (with `tdm$U.saveModel==FALSE` smaller `.RData` files).

Example usages of function `tdmBigLoop` are shown in

```r
demo(demo03sonar)
demo(demo03sonar_B)
demo(demo04cpu)
```

where the corresponding R-sources are in directory `demo`.

**Author(s)**

Wolfgang Konen (<wolfgang.konen@th-koeln.de>), THK

**See Also**

`tdmBigLoop`, `tdmDispatchTuner`, `unbiasedRun`

**Examples**

```r
### This demo shows a complete tuned data mining process (level 3 of TDMR) where
### the data mining task is the classification task SONAR (from UCI repository,
### The data mining process is in main_sonar.R, which calls tdmClassifyLoop and tdmClassify
### with Random Forest as the prediction model.
### The three parameter to be tuned are CUTOFF1, CLASSWT2 and XPERC, as specified
### in file sonar_04.roi. The tuner used here is LHD.
### Tuning runs are rather short, to make the example run quickly.
```
**unbiasedRun**

Perform unbiased runs with best-solution parameters.

**Description**
Read the best solution of a parameter-tuning run from envT$bst, execute with these best parameters the function tdm$mainFunc (usually a classification or regression machine learning task), to see whether the result quality is reproducible on independent test data or on independently trained models.

**Usage**

```r
unbiasedRun(confFile, envT, dataObj = NULL, umode = "RSUB", withParams = FALSE, tdm = NULL)
```
Arguments

conffile  the configuration name, e.g. "appAcid_02.conf"

envT  environment, from which we need the objects

bst  data frame containing best results (merged over repeats)

res  data frame containing all results

theTuner  ["spot"] string

spotConfig  [NULL] a list with SPOT settings. If NULL, try to read spotConfig from conffFile.

finals  [NULL] a one-row data frame to which new columns with final results are added

dataObj  [NULL] contains the pre-fetched data with training-set and test-set part. If NULL, set it to tdmReadAndSplit(opts, tdm).

It is now deprecated to have dataObj==NULL.

umode  — deprecated as argument to unbiasedRun — , use the division provided in dataObj = tdmReadAndSplit(opts, tdm) which makes use of tdm$umode.

For downward compatibility only (if dataObj==NULL):

[ "RSUB" (default) | "CV" | "TST" | "SP_T" ], how to divide in training and test data for the unbiased runs:

"RSUB"  random subsampling into (1-tdm$TST.testFrac)% training and tdm$TST.testFrac% test data

"CV"  cross validation (CV) with tdm$nrun folds

"TST"  all data in opts$filename (or dsetTrnVa(dataObj)) are used for training, all data in opts$filetest (or dsetTest(dataObj)) are used for testing

"SP_T"  'split_test': prior to tuning, the data set was split by random subsampling into tdm$TST.testFrac% test and (1-tdm$TST.testFrac)% training-vali data, tagged via column "tdmSplit". Tuning was done on training-vali data. Now we use column "tdmSplit" to select the test data for unbiased evaluation. Training during unbiased evaluation is done on a fraction tdm$TST.trnFrac of the training-vali data

withParams  [FALSE] if =TRUE, add columns with best parameters to data frame finals (should be FALSE, if different runs have different parameters)

tdm  a list with TDM settings from which we use here the elements

mainFunc  the function to be called for unbiased evaluations

mainFile  change to the directory of mainFile before starting mainFunc

nrun  [5] how often to call the unbiased evaluation

nfold  [10] how many folds in CV (only relevant for umode="CV")

TST.testFrac  [0.2] test set fraction (only relevant for umode="RSUB" or ="SP_T")

The defaults in ' [...] ' are set by tdmDefaultsFill, if they are not defined on input.

Value

envT the augmented environment envT, with the following items updated
unbiasedRun

finals the final results
tdm the updated list with TDM settings
results last results (from last unbiased training)

Note

Side Effects: The list result, an object of class TDMClassifier or TDMRegressor as returned from tdm$mainFunc is written onto envT$result.
If envT$spotConfig is NULL, it is constructed from confFile.
spotConfig$opts (list with all parameter settings for the DM task) has to be non-NULL.

Author(s)

Wolfgang Konen, THK, 2013 - 2018

If envT$bst or envT$res is NULL, try to read it from the file (the filename is inferred envT$spotConfig. If this is NULL, it is constructed from confFile). We try to find the files for envT$bst or envT$res in dir envT$theTuner).

See Also

tdmBigLoop, TDMClassifier, TDMRegressor

Examples

```r
## Load the best results obtained in a prior tuning for the configuration "sonar_04.conf"
## with tuning method "spot". The result envT from a prior run of tdmBigLoop with this .conf
## is read from demo02sonar/demoSonar.RData.
## Run task main_sonar again with these best parameters, using the default settings from
## tdmDefaultsFill: umode="RSUB", tdm$nrun=5 and tdm$TST.testFrac=0.2.
path = paste(find.package("TDMR"), "demo02sonar", sep="/"
envT = tdmEnvTLoad("demoSonar.RData",path);     # loads envT
source(paste(path,"main_sonar.r",sep="/"));
envT$tdm$opts$Verbosity=1;
envT$sCList[[1]]$opts$path=path;                # overwrite a possibly older stored path
envT$spotConfig <- envT$sCList[[1]];
dataObj <- tdmReadTaskData(envT,envT$tdm);
envT <- unbiasedRun("sonar_04.conf",envT,dataObj,tdm=envT$tdm);
print(envT$finals);
```
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