Package ‘RSSL’

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LinkingTo Rcpp, RcppArmadillo

Suggests testthat, rmarkdown, SparseM, numDeriv, LiblineaR

Description A collection of implementations of semi-supervised classifiers and methods to evaluate their performance. The package includes implementations of, among others, Implicitly Constrained Learning, Moment Constrained Learning, the Transductive SVM, Manifold regularization, Maximum Contrastive Pessimistic Likelihood estimation, S4VM and WellSVM.

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URL https://github.com/jkrijthe/RSSL

BugReports https://github.com/jkrijthe/RSSL

Collate 'Generics.R' 'Classifier.R' 'CrossValidation.R'
'LeastSquaresClassifier.R' 'EMLeastSquaresClassifier.R'
'NormalBasedClassifier.R' 'LinearDiscriminantClassifier.R'
'ELinearDiscriminantClassifier.R' 'NearestMeanClassifier.R'
'EMNearestMeanClassifier.R' 'LogisticRegression.R'
'EntropyRegularizedLogisticRegression.R' 'Evaluate.R'
'GRFClassifier.R' 'GenerateSSLData.R' 'HelperFunctions.R'
'ICLeastSquaresClassifier.R' 'ICLinearDiscriminantClassifier.R'
'KernelLeastSquaresClassifier.R'
'KernelICLeastSquaresClassifier.R'
'LaplacianKernelLeastSquaresClassifier.R' 'LaplacianSVM.R'
'LearningCurve.R' 'LinearSVM.R' 'LogisticLossClassifier.R'
'MCLinearDiscriminantClassifier.R' 'MCNearestMeanClassifier.R'
'MCPLDA.R' 'MajorityClassClassifier.R' 'Measures.R'
'Plotting.R' 'QuadraticDiscriminantClassifier.R' 'RSSL.R'
'RcppExports.R' 'S4VM.R' 'SVM.R' 'SelfLearning.R' 'TSVM.R'
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**add_missinglabels_mar**

Throw out labels at random

**Description**

Original labels are saved in attribute `y_true`

**Usage**

```r
add_missinglabels_mar(df, formula = NULL, prob = 0.1)
```

**Arguments**

- `df` : data.frame; Data frame of interest
- `formula` : formula; Formula to indicate the outputs
- `prob` : numeric; Probability of removing the label

**See Also**

Other RSSL utilities: `LearningCurveSSL()`, `SSLDataFrameToMatrices()`, `df_to_matrices()`, `measure_accuracy()`, `missing_labels()`, `split_dataset_ssl()`, `split_random()`, `true_labels()`
adjacency_knn

Calculate knn adjacency matrix

Description

Calculates symmetric adjacency: objects are neighbours if either one of them is in the set of nearest neighbours of the other.

Usage

adjacency_knn(X, distance = "euclidean", k = 6)

Arguments

X matrix; input matrix
distance character; distance metric used in the dist function
k integer; Number of neighbours

Value

Symmetric binary adjacency matrix

BaseClassifier

Classifier used for enabling shared documenting of parameters

Description

Classifier used for enabling shared documenting of parameters

Usage

BaseClassifier(X, y, X_u, verbose, scale, eps, x_center, intercept, lambda, y_scale, kernel, use_Xu_for_scaling, ...)

Arguments

X matrix; Design matrix for labeled data
y factor or integer vector; Label vector
X_u matrix; Design matrix for unlabeled data
verbose logical; Controls the verbosity of the output
scale logical; Should the features be normalized? (default: FALSE)
eps numeric; Stopping criterion for the maximinimization
x_center logical; Should the features be centered?
### c.CrossValidation

Merge result of cross-validation runs on single datasets into a the same object

#### Description

Merge result of cross-validation runs on single datasets into a the same object

#### Usage

```r
## S3 method for class 'CrossValidation'
c(...)
```

#### Arguments

- `...` Named arguments for the different objects, where the name reflects the dataset name

### clapply

Use mclapply conditional on not being in RStudio

#### Description

Use mclapply conditional on not being in RStudio

#### Usage

```r
clapply(X, FUN, ..., mc.cores = getOption("mc.cores", 2L))
```

#### Arguments

- `X` vector
- `FUN` function to be applied to the elements of X
- `...` optional arguments passed to FUN
- `mc.cores` number of cores to use
### cov_ml

**Biased (maximum likelihood) estimate of the covariance matrix**

**Description**

Biased (maximum likelihood) estimate of the covariance matrix

**Usage**

`cov_ml(X)`

**Arguments**

- `X` matrix with observations

---

### CrossValidationSSL

**Cross-validation in semi-supervised setting**

**Description**

Cross-validation for semi-supervised learning, in which the dataset is split into three parts: labeled training object, unlabeled training object and validation objects. This can be used to evaluate different approaches to semi-supervised classification under the assumption the labels are missing at random. Different cross-validation schemes are implemented. See below for details.

**Usage**

`CrossValidationSSL(X, y, ..., verbose = FALSE, mc.cores = 1)`

#### S3 method for class 'list'

```r
CrossValidationSSL(X, y, ..., verbose = FALSE, mc.cores = 1)
```

#### S3 method for class 'matrix'

```r
CrossValidationSSL(X, y, classifiers, measures = list(Error = measure_error, k = 10, repeats = 1, verbose = FALSE, leaveout = "test", n_labeled = 10, prop_unlabeled = 0.5, time = TRUE, pre_scale = FALSE, pre_pca = FALSE, n_min = 1, low_level_cores = 1, ...))
```

**Arguments**

- `X` design matrix of the labeled objects
- `y` vector with labels
- `...` arguments passed to underlying functions
- `verbose` logical; Controls the verbosity of the output
mc.cores integer; Number of cores to be used
classifiers list; Classifiers to crossvalidate
measures named list of functions giving the measures to be used
k integer; Number of folds in the cross-validation
repeats integer; Number of repeated assignments to folds
leaveout either "labeled" or "test", see details
n_labeled Number of labeled examples, used in both leaveout modes
prop_unlabeled numeric; proportion of unlabeled objects
time logical; Whether execution time should be saved.
pre_scale logical; Whether the features should be scaled before the dataset is used
pre_pca logical; Whether the features should be preprocessed using a PCA step
n_min integer; Minimum number of labeled objects per class
low_level_cores integer; Number of cores to use compute repeats of the learning curve

Details

The input to this function can be either: a dataset in the form of a feature matrix and factor containing the labels, a dataset in the form of a formula and data.frame or a named list of these two options. There are two main modes in which the cross-validation can be carried out, controlled by the leaveout parameter. When leaveout is "labeled", the folds are formed by non-overlapping labeled training sets of a user specified size. Each of these folds is used as a labeled set, while the rest of the objects are split into the an unlabeled and the test set, controlled by prop_unlabeled parameter. Note that objects can be used multiple times for testing, when training on a different fold, while other objects may never used for testing.

The "test" option of leaveout, on the other hand, uses the folds as the test sets. This means every object will be used as a test object exactly once. The remaining objects in each training iteration are split randomly into a labeled and an unlabeled part, where the number of the labeled objects is controlled by the user through the n_labeled parameter.

Examples

```r
X <- model.matrix(Species~.-1,data=iris)
y <- iris$Species

classifiers <- list("LS"=function(X,y,X_u,y_u) {
   LeastSquaresClassifier(X,y,lambda=0)},
   "EM"=function(X,y,X_u,y_u) {
      SelfLearning(X,y,X_u,
      method=LeastSquaresClassifier})
)

measures <- list("Accuracy" = measure_accuracy,
    "Loss" = measure_losstest,
    "Loss labeled" = measure_losslab,
    "Loss Lab+Unlab" = measure_losstrain
)```
decisionvalues

) # Cross-validation making sure test folds are non-overlapping cvresults1 <- CrossValidationSSL(X,y,
classifiers=classifiers,
measures=measures,
leaveout="test", k=10,
repeats = 2,n_labeled = 10)
print(cvresults1)
plot(cvresults1)

# Cross-validation making sure labeled sets are non-overlapping cvresults2 <- CrossValidationSSL(X,y,
classifiers=classifiers,
measures=measures,
leaveout="labeled", k=10,
repeats = 2,n_labeled = 10,
prop_unlabeled=0.5)
print(cvresults2)
plot(cvresults2)

decisionvalues

Decision values returned by a classifier for a set of objects

Description

Returns decision values of a classifier

Usage

decisionvalues(object, newdata)

## S4 method for signature 'LeastSquaresClassifier'
decisionvalues(object, newdata)

## S4 method for signature 'KernelLeastSquaresClassifier'
decisionvalues(object, newdata)

## S4 method for signature 'LinearSVM'
decisionvalues(object, newdata)

## S4 method for signature 'SVM'
decisionvalues(object, newdata)

## S4 method for signature 'TSVM'
decisionvalues(object, newdata)

## S4 method for signature 'svmlinClassifier'
decisionvalues(object, newdata)
df_to_matrices

Convert data.frame with missing labels to matrices

Arguments

df: data.frame; Data

formula: formula; Description of problem

Description

Convert data.frame with missing labels to matrices

Usage

df_to_matrices(df, formula = NULL)

Arguments

df: data.frame; Data

formula: formula; Description of problem

See Also

Other RSSL utilities: LearningCurveSSL(), SSLDataFrameToMatrices(), add_missinglabels_mar(), measure_accuracy(), missing_labels(), split_dataset_ssl(), split_random(), true_labels()

diabetes

Useful for testing the WellSVM implementation
EMLeastSquaresClassifier

An Expectation Maximization like approach to Semi-Supervised Least Squares Classification

Description

As studied in Krijthe & Loog (2016), minimizes the total loss of the labeled and unlabeled objects by finding the weight vector and labels that minimize the total loss. The algorithm proceeds similar to EM, by subsequently applying a weight update and a soft labeling of the unlabeled objects. This is repeated until convergence.

Usage

EMLeastSquaresClassifier(X, y, X_u, x_center = FALSE, scale = FALSE, verbose = FALSE, intercept = TRUE, lambda = 0, eps = 1e-09, y_scale = FALSE, alpha = 1, beta = 1, init = "supervised", method = "block", objective = "label", save_all = FALSE, max_iter = 1000)

Arguments

X          matrix; Design matrix for labeled data
y          factor or integer vector; Label vector
X_u        matrix; Design matrix for unlabeled data
x_center   logical; Should the features be centered?
scale      Should the features be normalized? (default: FALSE)
verbose    logical; Controls the verbosity of the output
intercept  logical; Whether an intercept should be included
lambda     numeric; L2 regularization parameter
eps        Stopping criterion for the minimization
y_scale    logical; whether the target vector should be centered
alpha      numeric; the mixture of the new responsibilities and the old in each iteration of the algorithm (default: 1)
beta       numeric; value between 0 and 1 that determines how much to move to the new solution from the old solution at each step of the block gradient descent
init       objective character; "random" for random initialization of labels, "supervised" to use supervised solution as initialization or a numeric vector with a coefficient vector to use to calculate the initialization
method     character; one of "block", for block gradient descent or "simple" for LBFGS optimization (default="block")
objective  character; "responsibility" for hard label self-learning or "label" for soft-label self-learning
save_all   logical; saves all classifiers trained during block gradient descent
max_iter   integer; maximum number of iterations
EMLeastSquaresClassifier

Details

By default (method="block") the weights of the classifier are updated, after which the unknown labels are updated. method="simple" uses LBFGS to do this update simultaneously. Objective="responsibility" corresponds to the responsibility based, instead of the label based, objective function in Krijthe & Loog (2016), which is equivalent to hard-label self-learning.

References


See Also

Other RSSL classifiers: EMLinearDiscriminantClassifier, GRCFClassifier, ICLeastSquaresClassifier, ILinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

Examples

library(dplyr)
library(ggplot2)
set.seed(1)

df <- generate2ClassGaussian(200,d=2,var=0.2) %>%
  add_missinglabels_mar(Class~.,prob = 0.96)

# Soft-label vs. hard-label self-learning
classifiers <- list(
  "Supervised"=LeastSquaresClassifier(Class~.,df),
  "EM-Soft"=EMLeastSquaresClassifier(Class~.,df,objective="label"),
  "EM-Hard"=EMLeastSquaresClassifier(Class~.,df,objective="responsibility"))

df %>%
  ggplot(aes(x=X1,y=X2,color=Class)) +
  geom_point() +
  coord_equal() +
  scale_y_continuous(limits=c(-2,2)) +
  stat_classifier(aes(linetype=..classifier..),
                  classifiers=classifiers)
Description

Expectation Maximization applied to the linear discriminant classifier assuming Gaussian classes with a shared covariance matrix.

Usage

EMLinearDiscriminantClassifier(X, y, X_u, method = "EM", scale = FALSE, eps = 1e-08, verbose = FALSE, max_iter = 100)

Arguments

- **X**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **X_u**: matrix; Design matrix for unlabeled data
- **method**: character; Currently only "EM"
- **scale**: logical; Should the features be normalized? (default: FALSE)
- **eps**: Stopping criterion for the maximinimization
- **verbose**: logical; Controls the verbosity of the output
- **max_iter**: integer; Maximum number of iterations

Details

Starting from the supervised solution, uses the Expectation Maximization algorithm (see Dempster et al. (1977)) to iteratively update the means and shared covariance of the classes (Maximization step) and updates the responsibilities for the unlabeled objects (Expectation step).

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSM, LogisticLossClassifier, LogisticRegression, MLinearDiscriminantClassifier, MClNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVMe, USMLeastSquaresClassifier, WellSVM, svmlin()
EMNearestMeanClassifier

Semi-Supervised Nearest Mean Classifier using Expectation Maximization

Description

Expectation Maximization applied to the nearest mean classifier assuming Gaussian classes with a spherical covariance matrix.

Usage

EMNearestMeanClassifier(X, y, X_u, method = "EM", scale = FALSE, eps = 1e-04)

Arguments

- X: matrix; Design matrix for labeled data
- y: factor or integer vector; Label vector
- X_u: matrix; Design matrix for unlabeled data
- method: character; Currently only "EM"
- scale: Should the features be normalized? (default: FALSE)
- eps: Stopping criterion for the maximinimization

Details

Starting from the supervised solution, uses the Expectation Maximization algorithm (see Dempster et al. (1977)) to iteratively update the means and shared covariance of the classes (Maximization step) and updates the responsibilities for the unlabeled objects (Expectation step).

References


EntropyRegularizedLogisticRegression

Entropy Regularized Logistic Regression

Description

R Implementation of entropy regularized logistic regression implementation as proposed by Grandvalet & Bengio (2005). An extra term is added to the objective function of logistic regression that penalizes the entropy of the posterior measured on the unlabeled examples.
EntropyRegularizedLogisticRegression

Usage

EntropyRegularizedLogisticRegression(X, y, X_u = NULL, lambda = 0,
        lambda_entropy = 1, intercept = TRUE, init = NA, scale = FALSE,
        x_center = FALSE)

Arguments

X          matrix; Design matrix for labeled data
y          factor or integer vector; Label vector
X_u        matrix; Design matrix for unlabeled data
lambda     l2 Regularization
lambda_entropy Weight of the labeled observations compared to the unlabeled observations
intercept  logical; Whether an intercept should be included
init       Initial parameters for the gradient descent
scale      logical; Should the features be normalized? (default: FALSE)
x_center   logical; Should the features be centered?

Value

S4 object of class EntropyRegularizedLogisticRegression with the following slots:

w          weight vector
classnames the names of the classes

References


Examples

library(RSSL)
library(ggplot2)
library(dplyr)

# An example where ERLR finds a low-density separator, which is not
# the correct solution.
set.seed(1)
df <- generateSlicedCookie(1000,expected=FALSE) %>%
  add_missinglabels_mar(Class~,0.98)

class_lr <- LogisticRegression(Class~,df,lambd = 0.01)
class_erlr <- EntropyRegularizedLogisticRegression(Class~,df,
  lambda=0.01,lambda_entropy = 100)
ggplot(df, aes(x=X1, y=X2, color=Class)) + geom_point() +
stat_classifier(aes(linetype=.classifier..),
classifiers = list("LR"=class_lr, "ERLR"=class_erlr)) +
scale_y_continuous(limits=c(-2,2)) +
scale_x_continuous(limits=c(-2,2))

df_test <- generateSlicedCookie(1000, expected=FALSE)
mean(predict(class_lr, df_test)==df_test$Class)
mean(predict(class_erlr, df_test)==df_test$Class)

find_a_violated_label  Find a violated label

Description
Find a violated label

Usage
find_a_violated_label(alpha, K, y, ind_y, lr, y_init)

Arguments
alpha  classifier weights
K  kernel matrix
y  label vector
ind_y  Labeled/Unlabeled indicator
lr  positive ratio
y_init  label initialization

gaussian_kernel  calculated the gaussian kernel matrix

description

calculated the gaussian kernel matrix

Usage
gaussian_kernel(x, gamma, x_test = NULL)
**generate2ClassGaussian**

**Arguments**

- **x**: A \(d \times n\) training data matrix
- **gamma**: kernel parameter
- **x_test**: A \(d \times m\) testing data matrix

**Value**

- \(k\) - A \(n \times m\) kernel matrix and \(\text{dis\_mat}\) - A \(n \times m\) distance matrix

---

**generate2ClassGaussian**

*Generate data from 2 Gaussian distributed classes*

**Description**

Generate data from 2 Gaussian distributed classes

**Usage**

```r
generate2ClassGaussian(n = 10000, d = 100, var = 1, expected = TRUE)
```

**Arguments**

- **n**: integer; Number of examples to generate
- **d**: integer; dimensionality of the problem
- **var**: numeric; size of the variance parameter
- **expected**: logical; whether the decision boundary should be the expected or perpendicular

**See Also**

Other RSSL datasets: `generateABA()`, `generateCrescentMoon()`, `generateFourClusters()`, `generateParallelPlanes()`, `generateSlicedCookie()`, `generateSpirals()`, `generateTwoCircles()`

**Examples**

```r
data <- generate2ClassGaussian(n=1000,d=2,expected=FALSE)
plot(data[,1],data[,2],col=data$Class,asp=1)
```
generateABA

*Generate data from 2 alternating classes*

**Description**

Two clusters belonging to three classes: the cluster in the middle belongs to one class and the two on the outside to the others.

**Usage**

```r
generateABA(n = 100, d = 2, var = 1)
```

**Arguments**

- **n**: integer; Number of examples to generate
- **d**: integer; Dimensionality of the problem
- **var**: numeric; size of the variance parameter

**See Also**

Other RSSL datasets: `generate2ClassGaussian()`, `generateCrescentMoon()`, `generateFourClusters()`, `generateParallelPlanes()`, `generateSlicedCookie()`, `generateSpirals()`, `generateTwoCircles()`

**Examples**

```r
data <- generateABA(n=1000, d=2, var=1)
plot(data[,1], data[,2], col=data$Class, asp=1)
```

generateCrescentMoon

*Generate Crescent Moon dataset*

**Description**

Generate a "crescent moon"/"banana" dataset

**Usage**

```r
generateCrescentMoon(n = 100, d = 2, sigma = 1)
```

**Arguments**

- **n**: integer; Number of objects to generate
- **d**: integer; Dimensionality of the dataset
- **sigma**: numeric; Noise added
**generateFourClusters**

**See Also**

Other RSSL datasets: `generate2ClassGaussian()`, `generateABA()`, `generateFourClusters()`, `generateParallelPlanes()`, `generateSlicedCookie()`, `generateSpirals()`, `generateTwoCircles()`

**Examples**

```r
data <- generateCrescentMoon(150, 2, 1)
plot(data$X1, data$X2, col = data$Class, asp = 1)
```

---

**generateFourClusters  Generate Four Clusters dataset**

**Description**

Generate a four clusters dataset

**Usage**

```r
generateFourClusters(n = 100, distance = 6, expected = FALSE)
```

**Arguments**

- `n` integer; Number of observations to generate
- `distance` numeric; Distance between clusters (default: 6)
- `expected` logical; TRUE if the large margin equals the class boundary, FALSE if the class boundary is perpendicular to the large margin

**See Also**

Other RSSL datasets: `generate2ClassGaussian()`, `generateABA()`, `generateCrescentMoon()`, `generateParallelPlanes()`, `generateSlicedCookie()`, `generateSpirals()`, `generateTwoCircles()`

**Examples**

```r
data <- generateFourClusters(1000, distance = 6, expected = TRUE)
plot(data[, 1], data[, 2], col = data$Class, asp = 1)
```
generateParallelPlanes

Generate Parallel planes

Description
Generate Parallel planes

Usage
generateParallelPlanes(n = 100, classes = 3, sigma = 0.1)

Arguments
n integer; Number of objects to generate
classes integer; Number of classes
sigma double; Noise added

See Also
Other RSSL datasets: generate2ClassGaussian(), generateABA(), generateCrescentMoon(), generateFourClusters(), generateSlicedCookie(), generateSpirals(), generateTwoCircles()

Examples
library(ggplot2)
df <- generateParallelPlanes(100,3)
ggplot(df, aes(x=x,y=y,color=Class,shape=Class)) + geom_point()

generateSlicedCookie

Generate Sliced Cookie dataset

Description
Generate a sliced cookie dataset: a circle with a large margin in the middle.

Usage
generateSlicedCookie(n = 100, expected = FALSE, gap = 1)
Arguments

- **n**: integer; number of observations to generate
- **expected**: logical; TRUE if the large margin equals the class boundary, FALSE if the class boundary is perpendicular to the large margin
- **gap**: numeric; Size of the gap

Value

A data.frame with n objects from the sliced cookie example

See Also

Other RSSL datasets: `generate2ClassGaussian()`, `generateABA()`, `generateCrescentMoon()`, `generateFourClusters()`, `generateParallelPlanes()`, `generateSpirals()`, `generateTwoCircles()`

Examples

```r
data <- generateSlicedCookie(1000, expected=FALSE)
plot(data[,1],data[,2],col=data$Class,asp=1)
```

---

**generateSpirals**  
*Generate Intersecting Spirals*

Description

Generate Intersecting Spirals

Usage

```r
generateSpirals(n = 100, sigma = 0.1)
```

Arguments

- **n**: integer; Number of objects to generate per class
- **sigma**: numeric; Noise added

See Also

Other RSSL datasets: `generate2ClassGaussian()`, `generateABA()`, `generateCrescentMoon()`, `generateFourClusters()`, `generateParallelPlanes()`, `generateSlicedCookie()`, `generateTwoCircles()`

Examples

```r
data <- generateSpirals(100,sigma=0.1)
#plot3D::scatter3D(data$x,data$y,data$z,col="black")
```
generateTwoCircles  Generate data from 2 circles

Description
One circle circumscribes the other

Usage

```r
generateTwoCircles(n = 100, noise_var = 0.2)
```

Arguments

- `n`: integer; Number of examples to generate
- `noise_var`: numeric; size of the variance parameter

See Also

Other RSSL datasets: `generate2ClassGaussian()`, `generateABA()`, `generateCrescentMoon()`, `generateFourClusters()`, `generateParallelPlanes()`, `generateSlicedCookie()`, `generateSpirals()`

geom_classifier  Plot RSSL classifier boundary (deprecated)

Description
Deprecated: Use geom_linearclassifier or stat_classifier to plot classification boundaries

Usage

```r
geom_classifier(..., show_guide = TRUE)
```

Arguments

- `...`: List of trained classifiers
- `show_guide`: logical (default: TRUE); Show legend
geom_linearclassifier  Plot linear RSSL classifier boundary

Description

Plot linear RSSL classifier boundary

Usage

gem_linearclassifier(..., show_guide = TRUE)

Arguments

... List of trained classifiers
show_guide logical (default: TRUE); Show legend

Examples

library(ggplot2)
library(dplyr)

df <- generate2ClassGaussian(100,d=2,var=0.2) %>%
  add_missinglabels_mar(Class~., 0.8)

df %>%
ggplot(aes(x=X1,y=X2,color=Class)) +
geom_point() +
geom_linearclassifier("Supervised"=LinearDiscriminantClassifier(Class~.,df),
  "EM"=EMLinearDiscriminantClassifier(Class~.,df))

GRFClassifier  Label propagation using Gaussian Random Fields and Harmonic functions

Description

Implements the approach proposed in Zhu et al. (2003) to label propagation over an affinity graph. Note, as in the original paper, we consider the transductive scenario, so the implementation does not generalize to out of sample predictions. The approach minimizes the squared difference in labels assigned to different objects, where the contribution of each difference to the loss is weighted by the affinity between the objects. The default in this implementation is to use a knn adjacency matrix based on euclidean distance to determine this weight. Setting adjacency="heat" will use an RBF kernel over euclidean distances between objects to determine the weights.
GRFClassifier

Usage

```
GRFClassifier(X, y, X_u, adjacency = "nn",
adjacency_distance = "euclidean", adjacency_k = 6,
adjacency_sigma = 0.1, class_mass_normalization = TRUE, scale = FALSE,
x_center = FALSE)
```

Arguments

- **X**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **X_u**: matrix; Design matrix for unlabeled data
- **adjacency**: character; "nn" for nearest neighbour graph or "heat" for radial basis adjacency matrix
- **adjacency_distance**: character; distance metric for nearest neighbour adjacency matrix
- **adjacency_k**: integer; number of neighbours for the nearest neighbour adjacency matrix
- **adjacency_sigma**: double; width of the rbf adjacency matrix
- **class_mass_normalization**: logical; Should the Class Mass Normalization heuristic be applied? (default: TRUE)
- **scale**: logical; Should the features be normalized? (default: FALSE)
- **x_center**: logical; Should the features be centered?

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM, LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

Examples

```
library(RSSL)
library(ggplot2)
library(dplyr)

set.seed(1)
df_circles <- generateTwoCircles(400, noise=0.1) %>%
```
```
add_missinglabels_mar(Class~,0.99)

# Visualize the problem
df_circles %>%
  ggplot(aes(x=X1,y=X2,color=Class)) +
  geom_point() +
  coord_equal()

# Visualize the solution
class_grf <- GRFClassifier(Class~,df_circles,
                          adjacency="heat",
                          adjacency_sigma = 0.1)
df_circles %>%
  filter(is.na(Class)) %>%
  mutate(Responsibility=responsibilities(class_grf)[,1]) %>%
  ggplot(aes(x=X1,y=X2,color=Responsibility)) +
  geom_point() +
  coord_equal()

# Generate problem
df_para <- generateParallelPlanes()
df_para$Class <- NA
df_para$Class[1] <- "a"
df_para$Class[101] <- "b"
df_para$Class[201] <- "c"
df_para$Class <- factor(df_para$Class)

# Visualize problem
df_para %>%
  ggplot(aes(x=x,y=y,color=Class)) +
  geom_point() +
  coord_equal()

# Estimate GRF classifier with knn adjacency matrix (default)
class_grf <- GRFClassifier(Class~,df_para)

df_para %>%
  filter(is.na(Class)) %>%
  mutate(Assignment=factor(apply(responsibilities(class_grf),1,which.max))) %>%
  ggplot(aes(x=x,y=y,color=Assignment)) +
  geom_point()
```

### harmonic_function

Direct R Translation of Xiaojin Zhu’s Matlab code to determine harmonic solution

### Description

Direct R Translation of Xiaojin Zhu’s Matlab code to determine harmonic solution
Usage

harmonic_function(W, Y)

Arguments

W  
matrix; weight matrix where the first L rows/columns correspond to the labeled examples.

Y  
matrix; l by c 0,1 matrix encoding class assignments for the labeled objects

Value

The harmonic solution, i.e. eq (5) in the ICML paper, with or without class mass normalization

ICLeastSquaresClassifier

*Implicitly Constrained Least Squares Classifier*

Description


Usage

ICLeastSquaresClassifier(X, y, X_u = NULL, lambda1 = 0, lambda2 = 0, 
intercept = TRUE, x_center = FALSE, scale = FALSE, method = "LBFGS", 
projection = "supervised", lambda_prior = 0, trueprob = NULL, 
eps = 1e-09, y_scale = FALSE, use_Xu_for_scaling = TRUE)

Arguments

X  
Design matrix, intercept term is added within the function

y  
Vector or factor with class assignments

X_u  
Design matrix of the unlabeled data, intercept term is added within the function

lambda1  
Regularization parameter in the unlabeled+labeled data regularized least squares

lambda2  
Regularization parameter in the labeled data only regularized least squares

intercept  
TRUE if an intercept should be added to the model

x_center  
logical; Whether the feature vectors should be centered

scale  
logical; If TRUE, apply a z-transform to all observations in X and X_u before running the regression

method  
Either "LBFGS" for solving using L-BFGS-B gradient descent or "QP" for a quadratic programming based solution

projection  
One of "supervised", "semisupervised" or "euclidean"

lambda_prior  
numeric; prior on the deviation from the supervised mean y
**ICLeastSquaresClassifier**

trueprob numeric; true mean y for all data
eps numeric; Stopping criterion for the maximinimization
y_scale logical; whether the target vector should be centered
use_Xu_for_scaling logical; whether the unlabeled objects should be used to determine the mean and scaling for the normalization

**Details**

In Implicitly Constrained semi-supervised Least Squares (ICLS) of Krijthe & Loog (2015), we minimize the quadratic loss on the labeled objects, while enforcing that the solution has to be a solution that minimizes the quadratic loss for all objects for some (fractional) labeling of the data (the implicit constraints). The goal of this classifier is to use the unlabeled data to update the classifier, while making sure it still works well on the labeled data.

The Projected estimator of Krijthe & Loog (2016) builds on this by finding a classifier within the space of classifiers that minimize the quadratic loss on all objects for some labeling (the implicit constrained), that minimizes the distance to the supervised solution for some appropriately chosen distance measure. Using the projection="semisupervised", we get certain guarantees that this solution is always better than the supervised solution (see Krijthe & Loog (2016)), while setting projection="supervised" is equivalent to ICLS.

Both methods (ICLS and the projection) can be formulated as a quadratic programming problem and solved using either a quadratic programming solver (method="QP") or using a gradient descent approach that takes into account certain bounds on the labelings (method="LBFGS"). The latter is the preferred method.

**Value**

S4 object of class ICLeastSquaresClassifier with the following slots:

theta weight vector
classnames the names of the classes
modelform formula object of the model used in regression
scaling a scaling object containing the parameters of the z-transforms applied to the data
optimization the object returned by the optim function
unlabels the labels assigned to the unlabeled objects

**References**


ICLinearDiscriminantClassifier

**Implicitly Constrained Semi-supervised Linear Discriminant Classifier**

**Description**

Semi-supervised version of Linear Discriminant Analysis using implicit constraints as described in (Krijthe & Loog 2014). This method finds the soft labeling of the unlabeled objects, whose resulting LDA solution gives the highest log-likelihood when evaluated on the labeled objects only. See also ICLeastSquaresClassifier.

**Usage**

ICLinearDiscriminantClassifier(X, y, X_u, prior = NULL, scale = FALSE, init = NULL, sup_prior = FALSE, x_center = FALSE, ...)

**Arguments**

- **x** design matrix of the labeled objects
- **y** vector with labels
- **X_u** design matrix of the labeled objects
- **prior** set a fixed class prior

**Examples**

```r
data(testdata)
w1 <- LeastSquaresClassifier(testdata$X, testdata$y, 
                              intercept = TRUE, x_center = FALSE, scale=FALSE)
w2 <- ICLeastSquaresClassifier(testdata$X, testdata$y, 
                              testdata$X_u, intercept = TRUE, x_center = FALSE, scale=FALSE)
plot(testdata$X[,1],testdata$X[,2],col=factor(testdata$y),asp=1)
points(testdata$X_u[,1],testdata$X_u[,2],col="darkgrey",pch=16,cex=0.5)
abline(line_coefficients(w1)$intercept, 
       line_coefficients(w1)$slope,lty=2)
abline(line_coefficients(w2)$intercept, 
       line_coefficients(w2)$slope,lty=1)
```

**See Also**

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()
KernelICLeastSquaresClassifier

scale logical; Should the features be normalized? (default: FALSE)
init not currently used
sup_prior logical; use the prior estimates based only on the labeled data, not the imputed labels (default: FALSE)
x_center logical; Whether the data should be centered

... Additional Parameters, Not used

References


See Also

Other RSSL classifiers: EMLEastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

KernelICLeastSquaresClassifier

Kernelized Implicitly Constrained Least Squares Classification

Description

A kernel version of the implicitly constrained least squares classifier, see ICLeastSquaresClassifier.

Usage

KernelICLeastSquaresClassifier(X, y, X_u, lambda = 0, kernel = vanilladot(), x_center = TRUE, scale = TRUE, y_scale = TRUE, lambda_prior = 0, classprior = 0, method = "LBFGS", projection = "semisupervised")

Arguments

X matrix; Design matrix for labeled data
y factor or integer vector; Label vector
X_u matrix; Design matrix for unlabeled data
lambda numeric; L2 regularization parameter
kernel kernlab::kernel to use
x_center logical; Should the features be centered?
KernelLeastSquaresClassifier

Kernelized Least Squares Classifier

Description

Use least squares regression as a classification technique using a numeric encoding of classes as targets. Note this method minimizes quadratic loss, not the truncated quadratic loss.

Usage

KernelLeastSquaresClassifier(X, y, lambda = 0, kernel = vanilladot(), x_center = TRUE, scale = TRUE, y_scale = TRUE)

Arguments

X     Design matrix, intercept term is added within the function
y     Vector or factor with class assignments
lambda Regularization parameter of the l2 penalty in regularized least squares
kernel kernlab kernel function
x_center TRUE, whether the dependent variables (features) should be centered
scale If TRUE, apply a z-transform to the design matrix X before running the regression
y_scale TRUE center the target vector

Value

S4 object of class LeastSquaresClassifier with the following slots:

theta weight vector
classnames the names of the classes
modelform formula object of the model used in regression
scaling a scaling object containing the parameters of the z-transforms applied to the data
See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, LaplacianKernelLeastSquaresClassifier().
LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

Examples

library(RSSL)
library(ggplot2)
library(dplyr)

# Two class problem
df <- generateCrescentMoon(200)

class_lin <- KernelLeastSquaresClassifier(Class~.,df,
kernel=kernlab::vanilladot(), lambda=1)
class_rbf1 <- KernelLeastSquaresClassifier(Class~.,df,
kernel=kernlab::rbfdot(), lambda=1)
class_rbf5 <- KernelLeastSquaresClassifier(Class~.,df,
kernel=kernlab::rbfdot(5), lambda=1)
class_rbf10 <- KernelLeastSquaresClassifier(Class~.,df,
kernel=kernlab::rbfdot(10), lambda=1)

df %>%
  ggplot(aes(x=X1,y=X2,color=Class,shape=Class)) +
geom_point() +
coord_equal() +
stat_classifier(aes(linetype=..classifier..),
classifiers = list("Linear"=class_lin,
"RBF sigma=1"=class_rbf1,
"RBF sigma=5"=class_rbf5,
"RBF sigma=10"=class_rbf10),
color="black")

# Second Example
dmat<-model.matrix(Species~.-1,iris[51:150,])
tvec<-droplevels(iris$Species[51:150])
testdata <- data.frame(tvec,dmat[,1:2])
colnames(testdata)<-c("Class","X1","X2")

precision<-100
xgrid<-seq(min(dmat[,1]),max(dmat[,1]),length.out=precision)
ygrid<-seq(min(dmat[,2]),max(dmat[,2]),length.out=precision)
gridmat <- expand.grid(xgrid,ygrid)

g_kernel<-KernelLeastSquaresClassifier(dmat[,1:2],tvec,
kernel=kernlab::rbfdot(0.01),
lambda=0.000001, scale = TRUE)

plotframe <- cbind(gridmat, decisionvalues(g_kernel,gridmat))
LaplacianKernelLeastSquaresClassifier

Description

Implements manifold regularization through the graph Laplacian as proposed by Belkin et al. 2006. As an adjacency matrix, we use the k nearest neighbour graph based on a chosen distance (default: euclidean).

Usage

LaplacianKernelLeastSquaresClassifier(X, y, X_u, lambda = 0, gamma = 0, 
kernel = kernlab::vanilladot(), adjacency_distance = "euclidean", 
adjacency_k = 6, x_center = TRUE, scale = TRUE, y_scale = TRUE, 
normalized_laplacian = FALSE)

Arguments

x  matrix; Design matrix for labeled data
y  factor or integer vector; Label vector
LaplacianKernelLeastSquaresClassifier

X_u  matrix; Design matrix for unlabeled data
lambda numeric; L2 regularization parameter
gamma numeric; Weight of the unlabeled data
kernel kernlab::kernel to use
adjacency_distance character; distance metric used to construct adjacency graph from the dist function. Default: "euclidean"
adjacency_k integer; Number of of neighbours used to construct adjacency graph.
x_center logical; Should the features be centered?
scale logical; Should the features be normalized? (default: FALSE)
y_scale logical; whether the target vector should be centered
normalized_laplacian logical; If TRUE use the normalized Laplacian, otherwise, the Laplacian is used

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

Examples

library(RSSL)
library(ggplot2)
library(dplyr)

## Example 1: Half moons

# Generate a dataset
set.seed(2)
df_orig <- generateCrescentMoon(100,sigma = 0.3)
df <- df_orig %>%
  add_missinglabels_mar(Class~.,0.98)
lambda <- 0.01
gamma <- 10000
rbf_param <- 0.125

# Train classifiers
## Not run:
class_sup <- KernelLeastSquaresClassifier(
  Class~.,df,
  kernel=kernlab::rbfdot(rbf_param),
  lambda=lambda,scale=FALSE)

class_lap <- LaplacianKernelLeastSquaresClassifier(
  Class~.,df,
  kernel=kernlab::rbfdot(rbf_param),
  lambda=lambda,gamma=gamma,
  normalized_laplacian = TRUE,
  scale=FALSE)

classifiers <- list("Lap"=class_lap,"Sup"=class_sup)

# Plot classifiers (can take a couple of seconds)

df %>%
  ggplot(aes(x=X1,y=X2,color=Class)) +
  geom_point() +
  coord_equal() +
  stat_classifier(aes(linetype=..classifier..),
    classifiers = classifiers ,
    color="black")

# Calculate the loss
lapply(classifiers,function(c) mean(loss(c,df_orig)))

## End(Not run)

## Example 2: Two circles
set.seed(1)
df_orig <- generateTwoCircles(1000,noise=0.05)
df <- df_orig %>%
  add_missinglabels_mar(Class~.,0.994)

lambda <- 10e-12
gamma <- 100
rbf_param <- 0.1

# Train classifiers
## Not run:
class_sup <- KernelLeastSquaresClassifier(
  Class~.,df,
  kernel=kernlab::rbfdot(rbf_param),
  lambda=lambda,scale=TRUE)

class_lap <- LaplacianKernelLeastSquaresClassifier(
  Class~.,df,
  kernel=kernlab::rbfdot(rbf_param),
  adjacency_k = 30,
  lambda=lambda,gamma=gamma,
LaplacianSVM

```r
normalized_laplacian = TRUE,
scale=TRUE)

classifiers <- list("Lap"=class_lap,"Sup"=class_sup)

# Plot classifiers (Can take a couple of seconds)
df %>%
ggplot(aes(x=X1,y=X2,color=Class,size=Class)) +
scale_size_manual(values=c("1"=3,"2"=3),na.value=1) +
geom_point() +
coord_equal() +
stat_classifier(aes(linetype=..classifier..),
classifiers = classifiers ,
color="black",size=1)

## End(Not run)
```

---

**LaplacianSVM**

**Laplacian SVM classifier**

**Description**

Manifold regularization applied to the support vector machine as proposed in Belkin et al. (2006). As an adjacency matrix, we use the k nearest neighbour graph based on a chosen distance (default: euclidean).

**Usage**

```
LaplacianSVM(X, y, X_u = NULL, lambda = 1, gamma = 1, scale = TRUE,
             kernel = vanilladot(), adjacency_distance = "euclidean",
             adjacency_k = 6, normalized_laplacian = FALSE, eps = 1e-09)
```

**Arguments**

- `X` matrix; Design matrix for labeled data
- `y` factor or integer vector; Label vector
- `X_u` matrix; Design matrix for unlabeled data
- `lambda` numeric; L2 regularization parameter
- `gamma` numeric; Weight of the unlabeled data
- `scale` logical; Should the features be normalized? (default: FALSE)
- `kernel` kernlab::kernel to use
- `adjacency_distance` character; distance metric used to construct adjacency graph from the dist function. Default: "euclidean"
- `adjacency_k` integer; Number of of neighbours used to construct adjacency graph.
normalized_laplacian
   logical; If TRUE use the normalized Laplacian, otherwise, the Laplacian is used

eps
   numeric; Small value to ensure positive definiteness of the matrix in the QP formulation

Value

S4 object of type LaplacianSVM

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

Examples

library(RSSL)
library(ggplot2)
library(dplyr)

## Example 1: Half moons

# Generate a dataset
set.seed(2)
df_orig <- generateCrescentMoon(100, sigma = 0.3)
df <- df_orig %>%
   add_missinglabels_mar(Class~., 0.98)

lambda <- 0.001
C <- 1/(lambda*2*sum(!is.na(df$Class)))
gamma <- 10000
rbf_param <- 0.125

# Train classifiers
class_sup <- SVM(
   Class~., df,
   kernel=kernlab::rbfdot(rbf_param),
   C=C, scale=FALSE)

class_lap <- LaplacianSVM(
   Class~., df,
LaplacianSVM

```r
kernel=kernlab::rbfdot(rbf_param),
lambda=lambda, gamma=gamma,
normalized_laplacian = TRUE,
scale=FALSE)

classifiers <- list("Lap"=class_lap,"Sup"=class_sup)

# This takes a little longer to run:
# class_tsvm <- TSVM(
# Class~.,df,
# kernel=kernlab::rbfdot(rbf_param),
# C=C,Cstar=0,s=-0.8,
# scale=FALSE,balancing_constraint=TRUE)
# classifiers <- list("Lap"=class_lap,"Sup"=class_sup,"TSVM"=class_tsvm)

# Plot classifiers (Can take a couple of seconds)
## Not run:
df %>%
  ggplot(aes(x=X1,y=X2,color=Class)) +
  geom_point() +
  coord_equal() +
  stat_classifier(aes(linetype=..classifier..),
                  classifiers = classifiers ,
                  color="black")

## End(Not run)

# Calculate the loss
lapply(classifiers,function(c) mean(loss(c,df_orig)))

## Example 2: Two circles
set.seed(3)
df_orig <- generateTwoCircles(1000,noise=0.05)
df <- df_orig %>%
  add_missinglabels_mar(Class~.,0.994)

lambda <- 0.000001
C <- 1/(lambda*2*sum(!is.na(df$Class)))
gamma <- 100
rbf_param <- 0.1

# Train classifiers (Takes a couple of seconds)
## Not run:
class_sup <- SVM(
  Class~.,df,
  kernel=kernlab::rbfdot(rbf_param),
  C=C,scale=FALSE)

class_lap <- LaplacianSVM(
  Class~.,df,
  kernel=kernlab::rbfdot(rbf_param),
  adjacency_k=50, lambda=lambda, gamma=gamma,
  normalized_laplacian = TRUE,
```
LearningCurveSSL

Description

Evaluate semi-supervised classifiers for different amounts of unlabeled training examples or different fractions of unlabeled vs. labeled examples.

Usage

LearningCurveSSL(X, y, ...)  

## S3 method for class 'matrix'
LearningCurveSSL(X, y, classifiers, measures = list(Accuracy = measure_accuracy), type = "unlabeled", n_l = NULL, with_replacement = FALSE, sizes = 2^(1:8), n_test = 1000, repeats = 100, verbose = FALSE, n_min = 1, dataset_name = NULL, test_fraction = NULL, fracs = seq(0.1, 0.9, 0.1), time = TRUE, pre_scale = FALSE, pre_pca = FALSE, low_level_cores = 1, ...)

Arguments

X      design matrix
y      vector of labels
...    arguments passed to underlying function
classifiers list; Classifiers to crossvalidate
measures named list of functions giving the measures to be used
LearningCurveSSL

- **type**: Type of learning curve, either "unlabeled" or "fraction"
- **n_l**: Number of labeled objects to be used in the experiments (see details)
- **with_replacement**: Indicated whether the subsampling is done with replacement or not (default: FALSE)
- **sizes**: vector with number of unlabeled objects for which to evaluate performance
- **n_test**: Number of test points if with_replacement is TRUE
- **repeats**: Number of learning curves to draw
- **verbose**: Print progressbar during execution (default: FALSE)
- **n_min**: Minimum number of labeled objects per class in
- **dataset_name**: character; Name of the dataset
- **test_fraction**: numeric; If not NULL a fraction of the object will be left out to serve as the test set
- **fracs**: list; fractions of labeled data to use
- **time**: logical; Whether execution time should be saved.
- **pre_scale**: logical; Whether the features should be scaled before the dataset is used
- **pre_pca**: logical; Whether the features should be preprocessed using a PCA step
- **low_level_cores**: integer; Number of cores to use compute repeats of the learning curve

**Details**

- **classifiers**: is a named list of classifiers, where each classifier should be a function that accepts
  4 arguments: a numeric design matrix of the labeled objects, a factor of labels, a numeric design
  matrix of unlabeled objects and a factor of labels for the unlabeled objects.
- **measures**: is a named list of performance measures. These are functions that accept seven argu-
  ments: a trained classifier, a numeric design matrix of the labeled objects, a factor of labels, a nu-
  meric design matrix of unlabeled objects and a factor of labels for the unlabeled objects, a numeric
  design matrix of the test objects and a factor of labels of the test objects. See measure_accuracy
  for an example.

This function allows for two different types of learning curves to be generated. If type="unlabeled",
the number of labeled objects remains fixed at the value of n_l, where sizes controls the number of
unlabeled objects. n_test controls the number of objects used for the test set, while all remaining
objects are used if with_replacement=FALSE in which case objects are drawn without replacement
from the input dataset. We make sure each class is represented by at least n_min labeled objects of
each class. For n_l, additional options include: "enough" which takes the max of the number of
features and 20, max(ncol(X)+5,20), "d" which takes the number of features or "2d" which takes 2
times the number of features.

If type="fraction" the total number of objects remains fixed, while the fraction of labeled objects
is changed. frac sets the fractions of labeled objects that should be considered, while test_fraction
determines the fraction of the total number of objects left out to serve as the test set.

**Value**

LearningCurve object
Least Squares Classifier

See Also

Other RSSL utilities: SSLDataFrameToMatrices(), add_missinglabels_mar(), df_to_matrices(), measure_accuracy(), missing_labels(), split_dataset_ssl(), split_random(), true_labels()

Examples

```r
set.seed(1)
df <- generate2ClassGaussian(2000,d=2,var=0.6)

classifiers <- list("LS"=function(X,y,X_u,y_u) {
  LeastSquaresClassifier(X,y,lambda=0),
  "Self"=function(X,y,X_u,y_u) {
    SelfLearning(X,y,X_u,LeastSquaresClassifier))
})

measures <- list("Accuracy" = measure_accuracy,
  "Loss Test" = measure_lossTest,
  "Loss labeled" = measure_losslab,
  "Loss Lab+Unlab" = measure_losstrain)

# These take a couple of seconds to run
## Not run:
# Increase the number of unlabeled objects
lc1 <- LearningCurveSSL(as.matrix(df[,1:2]),df$Class,
  classifiers=classifiers,
  measures=measures, n_test=1800,
  n_l=10,repeats=3)

plot(lc1)

# Increase the fraction of labeled objects, example with 2 datasets
lc2 <- LearningCurveSSL(X=list("Dataset 1"=as.matrix(df[,1:2]),
  "Dataset 2"=as.matrix(df[,1:2])),
  y=list("Dataset 1"=df$Class,
  "Dataset 2"=df$Class),
  classifiers=classifiers,
  measures=measures,
  type = "fraction",repeats=3,
  test_fraction=0.9)

plot(lc2)

## End(Not run)
```
**LeastSquaresClassifier**

**Description**

Classifier that minimizes the quadratic loss or, equivalently, least squares regression applied to a numeric encoding of the class labels as target. Note this method minimizes quadratic loss, not the truncated quadratic loss. Optionally, L2 regularization can be applied by setting the lambda parameter.

**Usage**

```
LeastSquaresClassifier(X, y, lambda = 0, intercept = TRUE,
  x_center = FALSE, scale = FALSE, method = "inverse", y_scale = FALSE)
```

**Arguments**

- **X**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **lambda**: Regularization parameter of the l2 penalty
- **intercept**: TRUE if an intercept should be added to the model
- **x_center**: TRUE, whether the dependent variables (features) should be centered
- **scale**: If TRUE, apply a z-transform to the design matrix X before running the regression
- **method**: Method to use for fitting. One of c("inverse","Normal","QR","BFGS")
- **y_scale**: If True scale the target vector

**Value**

S4 object of class LeastSquaresClassifier with the following slots:

- **theta**: weight vector
- **classnames**: the names of the classes
- **modelform**: formula object of the model used in regression
- **scaling**: a scaling object containing the parameters of the z-transforms applied to the data

**See Also**

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()
LinearDiscriminantClassifier

Linear Discriminant Classifier

Description

Implementation of the linear discriminant classifier. Classes are modeled as Gaussians with different means but equal covariance matrices. The optimal covariance matrix and means for the classes are found using maximum likelihood, which, in this case, has a closed form solution.

Usage

LinearDiscriminantClassifier(X, y, method = "closedform", prior = NULL, scale = FALSE, x_center = FALSE)

Arguments

X  Design matrix, intercept term is added within the function
y  Vector or factor with class assignments
method  the method to use. Either "closedform" for the fast closed form solution or "ml" for explicit maximum likelihood maximization
prior  A matrix with class prior probabilities. If NULL, this will be estimated from the data
scale  logical; If TRUE, apply a z-transform to the design matrix X before running the regression
x_center  logical; Whether the feature vectors should be centered

Value

S4 object of class LeastSquaresClassifier with the following slots:

modelform  weight vector
prior  the prior probabilities of the classes
mean  the estimates means of the classes
sigma  The estimated covariance matrix
classnames  a vector with the classnames for each of the classes
scaling  scaling object used to transform new observations

See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()
**LinearSVM**

**Description**

Implementation of the Linear Support Vector Classifier. Can be solved in the Dual formulation, which is equivalent to SVM or the Primal formulation.

**Usage**

```r
LinearSVM(X, y, C = 1, method = "Dual", scale = TRUE, eps = 1e-09,
    reltol = 1e-13, maxit = 100)
```

**Arguments**

- `X`: matrix; Design matrix for labeled data
- `y`: factor or integer vector; Label vector
- `C`: Cost variable
- `method`: Estimation procedure c("Dual","Primal","BGD")
- `scale`: Whether a z-transform should be applied (default: TRUE)
- `eps`: Small value to ensure positive definiteness of the matrix in QP formulation
- `reltol`: relative tolerance using during BFGS optimization
- `maxit`: Maximum number of iterations for BFGS optimization

**Value**

S4 object of type LinearSVM

**See Also**

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()
LinearTSVM

Linear CCCP Transductive SVM classifier

Description

Implementation for the Linear TSVM. This method is mostly for debugging purposes and does not allow for the balancing constraint or kernels, like the TSVM function.

Usage

LinearTSVM(X, y, X_u, C, Cstar, s = 0, x_center = FALSE, scale = FALSE, eps = 1e-06, verbose = FALSE, init = NULL)

Arguments

- **X**: matrix; Design matrix, intercept term is added within the function
- **y**: vector; Vector or factor with class assignments
- **X_u**: matrix; Design matrix of the unlabeled data, intercept term is added within the function
- **C**: numeric; Cost parameter of the SVM
- **Cstar**: numeric; Cost parameter of the unlabeled objects
- **s**: numeric; Parameter controlling the loss function of the unlabeled objects
- **x_center**: logical; Should the features be centered?
- **scale**: logical; If TRUE, apply a z-transform to all observations in X and X_u before running the regression
- **eps**: numeric; Convergence criterion
- **verbose**: logical; Print debugging messages (default: FALSE)
- **init**: numeric; Initial classifier parameters to start the convex concave procedure

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()
line_coefficients

Description

Loss of a classifier or regression function

Usage

line_coefficients(object, ...)

## S4 method for signature 'LeastSquaresClassifier'
line_coefficients(object)

## S4 method for signature 'NormalBasedClassifier'
line_coefficients(object)

## S4 method for signature 'LogisticRegression'
line_coefficients(object)

## S4 method for signature 'LinearSVM'
line_coefficients(object)

## S4 method for signature 'LogisticLossClassifier'
line_coefficients(object)

## S4 method for signature 'QuadraticDiscriminantClassifier'
line_coefficients(object)

## S4 method for signature 'SelfLearning'
line_coefficients(object)

Arguments

object  Classifier; Trained Classifier object

...  Not used

Value

numeric of the total loss on the test data
localDescent  
*Local descent*

**Description**

Local descent used in S4VM

**Usage**

```r
localDescent(instance, label, labelNum, unlabelNum, gamma, C, beta, alpha)
```

**Arguments**

- `instance`: Design matrix
- `label`: label vector
- `labelNum`: Number of labeled objects
- `unlabelNum`: Number of unlabeled objects
- `gamma`: Parameter for RBF kernel
- `C`: cost parameter for SVM
- `beta`: Controls fraction of objects assigned to positive class
- `alpha`: Controls fraction of objects assigned to positive class

**Value**

```r
list(predictLabel=predictLabel, acc=acc, values=values, model=model)
```

---

**LogisticLossClassifier**

*Logistic Loss Classifier*

**Description**

Find the linear classifier which minimizing the logistic loss on the training set, optionally using L2 regularization.

**Usage**

```r
LogisticLossClassifier(X, y, lambda = 0, intercept = TRUE, scale = FALSE, init = NA, x_center = FALSE, ...)
```
LogisticLossClassifier-class

Arguments

- **x**: Design matrix, intercept term is added within the function
- **y**: Vector with class assignments
- **lambda**: Regularization parameter used for l2 regularization
- **intercept**: TRUE if an intercept should be added to the model
- **scale**: If TRUE, apply a z-transform to all observations in X and X_u before running the regression
- **init**: Starting parameter vector for gradient descent
- **x_center**: logical; Whether the feature vectors should be centered
- **...**: additional arguments

Value

A S4 object with the following slots:

- **w**: the weight vector of the linear classifier
- **classnames**: vector with names of the classes

See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()
LogisticRegressionFast

(Regularized) Logistic Regression implementation

Description

Implementation of Logistic Regression that is useful for comparisons with semi-supervised logistic regression implementations, such as EntropyRegularizedLogisticRegression.

Usage

LogisticRegressionFast(X, y, lambda = 0, intercept = TRUE, scale = FALSE, init = NA, x_center = FALSE)

Arguments

- **X**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **lambda**: numeric; L2 regularization parameter
- **intercept**: logical; Whether an intercept should be included
- **scale**: logical; Should the features be normalized? (default: FALSE)
- **init**: numeric; Initialization of parameters for the optimization
- **x_center**: logical; Should the features be centered?

See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICleastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM, LogisticLossClassifier, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()
Arguments

- **x**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **lambda**: numeric; not used
- **intercept**: logical; Whether an intercept should be included
- **scale**: logical; Should the features be normalized? (default: FALSE)
- **init**: numeric; not used
- **x_center**: logical; Should the features be centered?

---

**Description**

Numerically more stable way to calculate log sum exp

**Usage**

```r
logsumexp(M)
```

**Arguments**

- **M**: matrix; m by n input matrix, sum with be over the rows

**Value**

matrix; m by 1 matrix

---

**loss**

*Loss of a classifier or regression function*

**Description**

Hinge loss on new objects of a trained LinearSVM

Hinge loss on new objects of a trained SVM
Usage

```r
loss(object, ...) 
## S4 method for signature 'LeastSquaresClassifier'
loss(object, newdata, y = NULL, ...)
## S4 method for signature 'NormalBasedClassifier'
loss(object, newdata, y = NULL)
## S4 method for signature 'LogisticRegression'
loss(object, newdata, y = NULL)
## S4 method for signature 'KernelLeastSquaresClassifier'
loss(object, newdata, y = NULL, ...)
## S4 method for signature 'LinearSVM'
loss(object, newdata, y = NULL)
## S4 method for signature 'LogisticLossClassifier'
loss(object, newdata, y = NULL, ...)
## S4 method for signature 'MajorityClassClassifier'
loss(object, newdata, y = NULL)
## S4 method for signature 'SVM'
loss(object, newdata, y = NULL)
## S4 method for signature 'SelfLearning'
loss(object, newdata, y = NULL, ...)
## S4 method for signature 'USMLeastSquaresClassifier'
loss(object, newdata, y = NULL, ...)
## S4 method for signature 'svmlinClassifier'
loss(object, newdata, y = NULL)
```

Arguments

- `object` Classifier; Trained Classifier
- `...` additional parameters
- `newdata` data.frame; object with test data
- `y` factor; True classes of the test data

Value

numeric; the total loss on the test data
**losslogsum**  
*LogsumLoss of a classifier or regression function*

**Description**

LogsumLoss of a classifier or regression function

**Usage**

```r
losslogsum(object, ...)  
```

```r
## S4 method for signature 'NormalBasedClassifier'
losslogsum(object, newdata, Y, X_u, Y_u)
```

**Arguments**

- `object`: Classifier or Regression object
- `...`: Additional parameters
- `newdata`: Design matrix of labeled objects
- `Y`: label matrix of labeled objects
- `X_u`: Design matrix of unlabeled objects
- `Y_u`: label matrix of unlabeled objects

**losspart**  
*Loss of a classifier or regression function evaluated on partial labels*

**Description**

Loss of a classifier or regression function evaluated on partial labels

**Usage**

```r
losspart(object, ...)  
```

```r
## S4 method for signature 'NormalBasedClassifier'
losspart(object, newdata, Y)
```

**Arguments**

- `object`: Classifier; Trained Classifier
- `...`: additional parameters
- `newdata`: design matrix
- `Y`: class responsibility matrix
**MajorityClassClassifier**

*Majority Class Classifier*

**Description**

Classifier that returns the majority class in the training set as the prediction for new objects.

**Usage**

```
MajorityClassClassifier(X, y, ...)
```

**Arguments**

- `X` : matrix; Design matrix for labeled data
- `y` : factor or integer vector; Label vector
- `...` : Not used

**See Also**

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM, LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

---

**MCLinearDiscriminantClassifier**

*Moment Constrained Semi-supervised Linear Discriminant Analysis.*

**Description**

A linear discriminant classifier that updates the estimates of the means and covariance matrix based on unlabeled examples.

**Usage**

```
MCLinearDiscriminantClassifier(X, y, X_u, method = "invariant",
                               prior = NULL, x_center = TRUE, scale = FALSE)
```
**MCNearestMeanClassifier**

**Arguments**

- `X` matrix; Design matrix for labeled data
- `y` factor or integer vector; Label vector
- `X_u` matrix; Design matrix for unlabeled data
- `method` character; One of c("invariant","closedform")
- `prior` Matrix (k by 1); Class prior probabilities. If NULL, estimated from data
- `x_center` logical; Should the features be centered?
- `scale` logical; Should the features be normalized? (default: FALSE)

**Details**

This method uses the parameter updates of the estimated means and covariance proposed in (Loog 2010). Using the method="invariant" option, uses the scale invariant parameter update proposed in (Loog 2014), while method="closedform" using the non-scale invariant version from (Loog 2012).

**References**


**See Also**

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCNearestMeanClassifier, MCPLDA, MajorityClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

---

**Description**

Update the means based on the moment constraints as defined in Loog (2010). The means estimated using the labeled data are updated by making sure their weighted mean corresponds to the overall mean on all (labeled and unlabeled) data. Optionally, the estimated variance of the classes can be re-estimated after this update is applied by setting `update_sigma` to TRUE. To get the true nearest mean classifier, rather than estimate the class priors, set them to equal priors using, for instance `prior=matrix(0.5,2).`
Usage

MCNearestMeanClassifier(X, y, X_u, update_sigma = FALSE, prior = NULL,
                        x_center = FALSE, scale = FALSE)

Arguments

- **X**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **X_u**: matrix; Design matrix for unlabeled data
- **update_sigma**: logical; Whether the estimate of the variance should be updated after the means have been updated using the unlabeled data
- **prior**: matrix; Class priors for the classes
- **x_center**: logical; Should the features be centered?
- **scale**: logical; Should the features be normalized? (default: FALSE)

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GLRClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM, LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

---

MCPLDA  
Maximum Contrastive Pessimistic Likelihood Estimation for Linear Discriminant Analysis

Description

Maximum Contrastive Pessimistic Likelihood (MCPL) estimation (Loog 2016) attempts to find a semi-supervised solution that has a higher likelihood compared to the supervised solution on the labeled and unlabeled data even for the worst possible labeling of the data. This is done by attempting to find a saddle point of the maximin problem, where the max is over the parameters of the semi-supervised solution and the min is over the labeling, while the objective is the difference in likelihood between the semi-supervised and the supervised solution measured on the labeled and unlabeled data. The implementation is a translation of the Matlab code of Loog (2016).
Usage
MCPLDA(X, y, X_u, x_center = FALSE, scale = FALSE, max_iter = 1000)

Arguments
- **X**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **X_u**: matrix; Design matrix for unlabeled data
- **x_center**: logical; Should the features be centered?
- **scale**: logical; Should the features be normalized? (default: FALSE)
- **max_iter**: integer; Maximum number of iterations

References

See Also
Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM, LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

measure_accuracy  Performance measures used in classifier evaluation

Description
Classification accuracy on test set and other performance measure that can be used in CrossValidationSSL and LearningCurveSSL

Usage
measure_accuracy(trained_classifier, X_l = NULL, y_l = NULL, X_u = NULL, y_u = NULL, X_test = NULL, y_test = NULL)

measure_error(trained_classifier, X_l = NULL, y_l = NULL, X_u = NULL, y_u = NULL, X_test = NULL, y_test = NULL)

measure_losstest(trained_classifier, X_l = NULL, y_l = NULL, X_u = NULL, y_u = NULL, X_test = NULL, y_test = NULL)
minimaxlda

Implements weighted likelihood estimation for LDA

Description

Implements weighted likelihood estimation for LDA

Usage

minimaxlda(a, w, u, iter)

Arguments

a is the data set
w is an indicator matrix for the K classes of a or, potentially, a weight matrix in which the fraction with which a sample belongs to a particular class is indicated
u is a bunch of unlabeled data
iter decides on the amount of time we spend on minimaxing the stuff
### missing_labels

**Value**

- m contains the means, p contains the class priors, iW contains the INVERTED within covariance matrix, uw returns the weights for the unlabeled data, i returns the number of iterations used

**Description**

Access the true labels for the objects with missing labels when they are stored as an attribute in a data frame

**Usage**

```r
missing_labels(df)
```

**Arguments**

- `df` data.frame; data.frame with y_true attribute

**See Also**

Other RSSL utilities: `LearningCurveSSL()`, `SSLDataFrameToMatrices()`, `add_missinglabels_mar()`, `df_to_matrices()`, `measure_accuracy()`, `split_dataset_ssl()`, `split_random()`, `true_labels()`

### NearestMeanClassifier  Nearest Mean Classifier

**Description**

Implementation of the nearest mean classifier modeled. Classes are modeled as gaussians with equal, spherical covariance matrices. The optimal covariance matrix and means for the classes are found using maximum likelihood, which, in this case, has a closed form solution. To get true nearest mean classification, set prior as a matrix with equal probability for all classes, i.e. matrix(0.5, 2).

**Usage**

```r
NearestMeanClassifier(X, y, prior = NULL, x_center = FALSE, scale = FALSE)
```

**Arguments**

- `X` matrix; Design matrix for labeled data
- `y` factor or integer vector; Label vector
- `prior` matrix; Class prior probabilities. If NULL, this will be estimated from the data
- `x_center` logical; Should the features be centered?
- `scale` logical; Should the features be normalized? (default: FALSE)
**Value**

S4 object of class LeastSquaresClassifier with the following slots:

- **modelform**: weight vector
- **prior**: the prior probabilities of the classes
- **mean**: the estimates means of the classes
- **sigma**: The estimated covariance matrix
- **classnames**: a vector with the classnames for each of the classes
- **scaling**: scaling object used to transform new observations

**See Also**

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

---

**plot.CrossValidation**

*Plot CrossValidation object*

**Description**

Plot CrossValidation object

**Usage**

```r
## S3 method for class 'CrossValidation'
plot(x, y, ...)
```

**Arguments**

- **x**: CrossValidation object
- **y**: Not used
- **...**: Not used
plot.LearningCurve

Plot LearningCurve object

Description

Plot LearningCurve object

Usage

```r
## S3 method for class 'LearningCurve'
plot(x, y, ...)
```

Arguments

- `x` LearningCurve object
- `y` Not used
- `...` Not used

posterior

Class Posteriors of a classifier

Description

Class Posteriors of a classifier

Usage

```r
posterior(object, ...)
```

Arguments

- `object` Classifier or Regression object
- `...` Additional parameters
- `newdata` matrix of dataframe of objects to be classified
predict,scaleMatrix-method

Predict for matrix scaling inspired by stdize from the PLS package

Description

Predict for matrix scaling inspired by stdize from the PLS package

Usage

## S4 method for signature 'scaleMatrix'
predict(object, newdata, ...)

Arguments

object scaleMatrix object
newdata data to be scaled
... Not used

PreProcessing

Preprocess the input to a classification function

Description

The following actions are carried out: 1. data.frames are converted to matrix form and labels converted to an indicator matrix 2. An intercept column is added if requested 3. centering and scaling is applied if requested.

Usage

PreProcessing(X, y, X_u = NULL, scale = FALSE, intercept = FALSE,
x_center = FALSE, use_Xu_for_scaling = TRUE)

Arguments

X Design matrix, intercept term is added within the function
y Vector or factor with class assignments
X_u Design matrix of the unlabeled observations
scale If TRUE, apply a z-transform to the design matrix X
intercept Whether to include an intercept in the design matrices
x_center logical (default: TRUE); Whether the feature vectors should be centered
use_Xu_for_scaling logical (default: TRUE); Should the unlabeled data be used to determine scaling?
PreProcessingPredict

Description

The following actions are carried out: 1. data.frames are converted to matrix form and labels converted to integers 2. An intercept column is added if requested 3. centering and scaling is applied if requested.

Usage

PreProcessingPredict(modelform, newdata, y = NULL, classnames = NULL, scaling = NULL, intercept = FALSE)

Arguments

modelform Formula object with model
newdata data.frame object with objects
y Vector or factor with class assignments (default: NULL)
classnames Vector with class names
scaling Apply a given z-transform to the design matrix X (default: NULL)
intercept Whether to include an intercept in the design matrices

Value

list object with the following objects:

X design matrix of the labeled data
y integer vector indicating the labels of the labeled data
X_u design matrix of the unlabeled data
classnames names of the classes corresponding to the integers in y
scaling a scaling object used to scale the test observations in the same way as the training set
modelform a formula object containing the used model
print.CrossValidation  Print CrossValidation object

Description

Print CrossValidation object

Usage

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

Arguments

x CrossValidation object
...
Not used

print.LearningCurve  Print LearningCurve object

Description

Print LearningCurve object

Usage

## S3 method for class 'LearningCurve'
print(x, ...)
**projection_simplex**

**Description**

\[ D_n = \{ x : x \text{ n-dim, } 1 \geq x \geq 0, \sum(x) = 1 \} \]

R translation of Loog’s version of Xiaojing Ye’s initial implementation. The algorithm works row-wise.

**Usage**

```
projection_simplex(y)
```

**Arguments**

- `y` matrix with vectors to be projected onto the simplex

**Value**

projection of \( y \) onto the simplex

**References**

Algorithm is explained as in [http://arxiv.org/abs/1101.6081](http://arxiv.org/abs/1101.6081)

---

**QuadraticDiscriminantClassifier**

**Quadratic Discriminant Classifier**

**Description**

Implementation of the quadratic discriminant classifier. Classes are modeled as Gaussians with different covariance matrices. The optimal covariance matrix and means for the classes are found using maximum likelihood, which, in this case, has a closed form solution.

**Usage**

```
QuadraticDiscriminantClassifier(X, y, prior = NULL, scale = FALSE, ...)
```

**Arguments**

- `X` matrix; Design matrix for labeled data
- `y` factor or integer vector; Label vector
- `prior` A matrix with class prior probabilities. If NULL, this will be estimated from the data
- `scale` logical; Should the features be normalized? (default: FALSE)
- `...` Not used
Value

S4 object of class LeastSquaresClassifier with the following slots:

- `modelform`: weight vector
- `prior`: the prior probabilities of the classes
- `mean`: the estimates means of the classes
- `sigma`: The estimated covariance matrix
- `classnames`: a vector with the classnames for each of the classes
- `scaling`: scaling object used to transform new observations

See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

---

**responsibilities**  
**Responsibilities assigned to the unlabeled objects**

Description

Responsibilities assigned to the unlabeled objects

Usage

```r
responsibilities(object, ...)  
```

Arguments

- `object`: Classifier; Trained Classifier
- `...`: additional parameters

Value

numeric; responsibilities on the unlabeled objects
Description

RSSL provides implementations for semi-supervised classifiers, as well as some functions to aid in the evaluation of these procedures.

Details

Most functions take a formula and data.frame or a matrix and factor as input and output a trained Classifier object, whose class is the class of a specific type of classifier model. predict can then be used to generate predictions for new objects, decisionvalues returns the decision values for new objects and loss outputs the loss used by the classifier evaluated on a set of new objects.

For a complete list of functions, use library(help = "RSSL").

Description

Show RSSL classifier

Show the contents of a classifier

Usage

```r
## S4 method for signature 'Classifier'
show(object)

## S4 method for signature 'NormalBasedClassifier'
show(object)

## S4 method for signature 'scaleMatrix'
show(object)
```

Arguments

- `object`: classifier
Description

Predict using RSSL classifier

For the SelfLearning Classifier the Predict Method delegates prediction to the specific model object

Usage

## S4 method for signature 'LeastSquaresClassifier'
predict(object, newdata, ...)

## S4 method for signature 'NormalBasedClassifier'
predict(object, newdata)

## S4 method for signature 'LogisticRegression'
predict(object, newdata)

## S4 method for signature 'GRFClassifier'
responsibilities(object, newdata, ...)

## S4 method for signature 'GRFClassifier'
predict(object, newdata = NULL, ...)

## S4 method for signature 'KernelLeastSquaresClassifier'
predict(object, newdata, ...)

## S4 method for signature 'LinearSVM'
predict(object, newdata)

## S4 method for signature 'LogisticLossClassifier'
predict(object, newdata)

## S4 method for signature 'MajorityClassClassifier'
predict(object, newdata)

## S4 method for signature 'SVM'
predict(object, newdata)

## S4 method for signature 'SelfLearning'
predict(object, newdata, ...)

## S4 method for signature 'USMLEastSquaresClassifier'
predict(object, newdata, ...)
## S4 method for signature 'WellSVM'
predict(object, newdata, ...)

## S4 method for signature 'WellSVM'
decisionvalues(object, newdata)

## S4 method for signature 'svmlinClassifier'
predict(object, newdata, ...)

### Arguments

- **object**: classifier
- **newdata**: objects to generate predictions for
- **...**: Other arguments

---

**S4VM**

*Safe Semi-supervised Support Vector Machine (S4VM)*

---

**Description**


**Usage**

S4VM(X, y, X_u = NULL, C1 = 100, C2 = 0.1, sample_time = 100, gamma = 0, x_center = FALSE, scale = FALSE, lambda_tradeoff = 3)

**Arguments**

- **X**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **X_u**: matrix; Design matrix for unlabeled data
- **C1**: double; Regularization parameter for labeled data
- **C2**: double; Regularization parameter for unlabeled data
- **sample_time**: integer; Number of low-density separators that are generated
- **gamma**: double; Width of RBF kernel
- **x_center**: logical; Should the features be centered?
- **scale**: logical; Should the features be normalized? (default: FALSE)
- **lambda_tradeoff**: numeric; Parameter that determines the amount of “risk” in obtaining a worse solution than the supervised solution, see Li & Zhou (2011)
Details

The method randomly generates multiple low-density separators (controlled by the sample_time parameter) and merges their predictions by solving a linear programming problem meant to penalize the cost of decreasing the performance of the classifier, compared to the supervised SVM. S4VM is a bit of a misnomer, since it is a transductive method that only returns predicted labels for the unlabeled objects. The main difference in this implementation compared to the original implementation is the clustering of the low-density separators: in our implementation empty clusters are not dropped during the k-means procedure. In the paper by Li (2011) the features are first normalized to [0,1], which is not automatically done by this function. Note that the solution may not correspond to a linear classifier even if the linear kernel is used.

Value

S4VM object with slots:

predictions Predictions on the unlabeled objects
labelings Labelings for the different clusters

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM, LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, SVM, SelfLearning, TSVM, USMLEastSquaresClassifier, WellSVM, svmlin()

Examples

library(RSSL)
library(dplyr)
library(ggplot2)
library(tidyr)
set.seed(1)
df_orig <- generateSlicedCookie(100, expected=TRUE)
df <- df_orig %>% add_missinglabels_mar(Class~., 0.95)
g_s <- SVM(Class~., df, C=1, scale=TRUE, x_center=TRUE)
g_s4 <- S4VM(Class~., df, C1=1, C2=0.1, lambda_tradeoff = 3, scale=TRUE, x_center=TRUE)
labs <- g_s4@labelings[,-c(1:5),]
colnames(labs) <- paste("Class", seq_len(ncol(g_s4@labelings)), sep="-")
# Show the labelings that the algorithm is considering
df %>%
filter(is.na(Class)) %>%
bind_cols(data.frame(labs,check.names = FALSE)) %>%
select(-Class) %>%
gather(Classifier,Label,-X1,-X2) %>%
ggplot(aes(x=X1,y=X2,color=Label)) +
geom_point() +
facet_wrap(~Classifier,ncol=5)

# Plot the final labeling that was selected
# Note that this may not correspond to a linear classifier
# even if the linear kernel is used.
# The solution does not seem to make a lot of sense,
# but this is what the current implementation returns
df %>%
filter(is.na(Class)) %>%
mutate(prediction=g_s4@predictions) %>%
ggplot(aes(x=X1,y=X2,color=prediction)) +
geom_point() +
stat_classifier(color="black", classifiers=list(g_s))

---

**S4VM-class** *LinearSVM Class*

**Description**

LinearSVM Class

**sample_k_per_level** *Sample k indices per levels from a factor*

**Description**

Sample k indices per levels from a factor

**Usage**

```
sample_k_per_level(y, k)
```

**Arguments**

- `y` factor; factor with levels
- `k` integer; number of indices to sample per level

**Value**

vector with indices for sample
scaleMatrix  
Matrix centering and scaling

Description

This function returns an object with a predict method to center and scale new data. Inspired by `stdize` from the PLS package.

Usage

scaleMatrix(x, center = TRUE, scale = TRUE)

Arguments

- `x`: matrix to be standardized
- `center`: TRUE if `x` should be centered
- `scale`: logical; TRUE if `x` should be scaled by the standard deviation

SelfLearning  
Self-Learning approach to Semi-supervised Learning

Description

Use self-learning (also known as Yarowsky’s algorithm or pseudo-labeling) to turn any supervised classifier into a semi-supervised method by iteratively labeling the unlabeled objects and adding these predictions to the set of labeled objects until the classifier converges.

Usage

SelfLearning(X, y, X_u = NULL, method, prob = FALSE, cautious = FALSE, max_iter = 100, ...)

Arguments

- `X`: matrix; Design matrix for labeled data
- `y`: factor or integer vector; Label vector
- `X_u`: matrix; Design matrix for unlabeled data
- `method`: Supervised classifier to use. Any function that accepts as its first argument a design matrix `X` and as its second argument a vector of labels `y` and that has a predict method.
- `prob`: Not used
- `cautious`: Not used
- `max_iter`: integer; Maximum number of iterations
- `...`: additional arguments to be passed to method
References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, TSVM, USMLastSquaresClassifier, WellSVM, svmlin()

Examples

data(testdata)
t_self <- SelfLearning(testdata$X, testdata$y, testdata$X_u, method=NearestMeanClassifier)
t_sup <- NearestMeanClassifier(testdata$X, testdata$y)
# Classification Error
1-mean(predict(t_self, testdata$X_test)==testdata$y_test)
1-mean(predict(t_sup, testdata$X_test)==testdata$y_test)
loss(t_self, testdata$X_test, testdata$y_test)

solve_svm

---

SVM solve.QP implementation

Description

SVM solve.QP implementation

Usage

solve_svm(K, y, C = 1)

Arguments

K Kernel matrix
y Output vector
C Cost parameter
split_dataset_ssl Create Train, Test and Unlabeled Set

Description
Create Train, Test and Unlabeled Set

Usage
split_dataset_ssl(X, y, frac_train = 0.8, frac_ssl = 0.8)

Arguments
- X: matrix; Design matrix
- y: factor; Label vector
- frac_train: numeric; Fraction of all objects to be used as training objects
- frac_ssl: numeric; Fraction of training objects to used as unlabeled objects

See Also
Other RSSL utilities: LearningCurveSSL(), SSLDataFrameToMatrices(), add_missinglabels_mar(), df_to_matrices(), measure_accuracy(), missing_labels(), split_random(), true_labels()

split_random Randomly split dataset in multiple parts

Description
The data.frame should start with a vector containing labels, or formula should be defined.

Usage
split_random(df, formula = NULL, splits = c(0.5, 0.5), min_class = 0)

Arguments
- df: data.frame; Data frame of interest
- formula: formula; Formula to indicate the outputs
- splits: numeric; Probability of of assigning to each part, automatically normalized, should be >1
- min_class: integer; minimum number of objects per class in each part

Value
list of data.frames
SSLDataFrameToMatrices

Convert data.frame to matrices for semi-supervised learners

Description

Given a formula object and a data.frame, extract the design matrix $X$ for the labeled observations, $X_u$ for the unlabeled observations and $y$ for the labels of the labeled observations. Note: always removes the intercept

Usage

SSLDataFrameToMatrices(model, D)

Arguments

- `model`: Formula object with model
- `D`: data.frame object with objects

Value

list object with the following objects:
- $X$: design matrix of the labeled data
- $X_u$: design matrix of the unlabeled data
- $y$: integer vector indicating the labels of the labeled data
- `classnames`: names of the classes corresponding to the integers in $y$

See Also

Other RSSL utilities: `LearningCurveSSL()`, `add_missinglabels_mar()`, `df_to_matrices()`, `measure_accuracy()`, `missing_labels()`, `split_dataset_ssl()`, `true_labels()`

Examples

```r
library(dplyr)

df <- generate2ClassGaussian(200,d=2)
dfs <- df %>% split_random(Class~.,split=c(0.5,0.3,0.2),min_class=1)
names(dfs) <- c("Train","Validation","Test")
lapply(dfs,summary)
```
stat_classifier

Plot RSSL classifier boundaries

Description

Plot RSSL classifier boundaries

Usage

stat_classifier(mapping = NULL, data = NULL, show.legend = NA,
breaks = 0, precision = 50, brute_force = FALSE,
classifiers = classifiers, ...)

Arguments

mapping aes; aesthetic mapping
data data.frame; data to be displayed
show.legend logical; Whether this layer should be included in the legend
inherit.aes logical; If FALSE, overrides the default aesthetics
breaks double; decision value for which to plot the boundary
precision integer; grid size to sketch classification boundary
brute_force logical; If TRUE, uses numerical estimation even for linear classifiers
classifiers List of Classifier objects to plot
... Additional parameters passed to geom

Examples

library(RSSL)
library(ggplot2)
library(dplyr)

df <- generateCrescentMoon(200)

# This takes a couple of seconds to run
## Not run:
g_svm <- SVM(Class~.,df,kernel = kernlab::rbfdot(sigma = 1))
g_ls <- LeastSquaresClassifier(Class~.,df)
g_nm <- NearestMeanClassifier(Class~.,df)

df %>%
ggplot(aes(x=X1,y=X2,color=Class,shape=Class)) +
  geom_point(size=3) +
  coord_equal() +
  scale_x_continuous(limits=c(-20,20), expand=c(0,0)) +
  scale_y_continuous(limits=c(-20,20), expand=c(0,0)) +
stderror

```
stat_classifier(aes(linetype=..classifier..),
               color="black", precision=50,
               classifiers=list("SVM"=g_svm,"NM"=g_nm,"LS"=g_ls))

## End(Not run)
```

---

**stderror**

*Calculate the standard error of the mean from a vector of numbers*

**Description**

Calculate the standard error of the mean from a vector of numbers

**Usage**

```
stderror(x)
```

**Arguments**

- `x` numeric; vector for which to calculate standard error

---

**summary.CrossValidation**

*Summary of Crossvalidation results*

**Description**

Summary of Crossvalidation results

**Usage**

```
## S3 method for class 'CrossValidation'
summary(object, measure = NULL, ...)
```

**Arguments**

- `object` CrossValidation object
- `measure` Measure of interest
- `...` Not used
svdinv

Inverse of a matrix using the singular value decomposition

Description
Inverse of a matrix using the singular value decomposition

Usage
svdinv(X)

Arguments
X matrix; square input matrix

Value
Y matrix; inverse of the input matrix

svdinvsqrtm

Taking the inverse of the square root of the matrix using the singular value decomposition

Description
Taking the inverse of the square root of the matrix using the singular value decomposition

Usage
svdinvsqrtm(X)

Arguments
X matrix; square input matrix

Value
Y matrix; inverse of the square root of the input matrix
svdsqrtm

Taking the square root of a matrix using the singular value decomposition

Description

Taking the square root of a matrix using the singular value decomposition

Usage

svdsqrtm(X)

Arguments

X matrix; square input matrix

Value

Y matrix; square root of the input matrix

SVM

SVM Classifier

Description

Support Vector Machine implementation using the quadprog solver.

Usage

SVM(X, y, C = 1, kernel = NULL, scale = TRUE, intercept = FALSE, x_center = TRUE, eps = 1e-09)

Arguments

X matrix; Design matrix for labeled data
y factor or integer vector; Label vector
C numeric; Cost variable
kernel kernlab::kernel to use
scale logical; Should the features be normalized? (default: FALSE)
intercept logical; Whether an intercept should be included
x_center logical; Should the features be centered?
eps numeric; Small value to ensure positive definiteness of the matrix in the QP formulation
Details

This implementation will typically be slower and use more memory than the svmlib implementation in the e1071 package. It is, however, useful for comparisons with the TSVM implementation.

Value

S4 object of type SVM

See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SelfLearning, TSVM, USMLeastSquaresClassifier, WellSVM, svmlin()

svmlin

svmlin implementation by Sindhwani & Keerthi (2006)

Description

R interface to the svmlin code by Vikas Sindhwani and S. Sathiya Keerthi for fast linear transductive SVMs.

Usage

svmlin(X, y, X_u = NULL, algorithm = 1, lambda = 1, lambda_u = 1,
       max_switch = 10000, pos_frac = 0.5, Cp = 1, Cn = 1,
       verbose = FALSE, intercept = TRUE, scale = FALSE, x_center = FALSE)

Arguments

X Matrix or sparseMatrix containing the labeled feature vectors, without intercept
y factor containing class assignments
X_u Matrix or sparseMatrix containing the unlabeled feature vectors, without intercept
algorithm integer; Algorithm choice, see details (default:1)
lambda double; Regularization parameter lambda (default 1)
lambda_u double; Regularization parameter lambda_u (default 1)
max_switch integer; Maximum number of switches in TSVM (default 10000)
pos_frac double; Positive class fraction of unlabeled data (default 0.5)
Cp double; Relative cost for positive examples (only available with algorithm 1)
Cn double; Relative cost for positive examples (only available with algorithm 1)
verbose logical; Controls the verbosity of the output
intercept logical; Whether an intercept should be included
scale logical; Should the features be normalized? (default: FALSE)
x_center logical; Should the features be centered?

Details
The codes to select the algorithm are the following: 0. Regularized Least Squares Classification 1. SVM (L2-SVM-MFN) 2. Multi-switch Transductive SVM (using L2-SVM-MFN) 3. Deterministic Annealing Semi-supervised SVM (using L2-SVM-MFN).

References

See Also
Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLEastSquaresClassifier, WellSVM

Examples
data(svmlin_example)
t_svmlin_1 <- svmlin(svmlin_example$X_train[1:50,],
                     svmlin_example$y_train, X_u=NULL, lambda = 0.001)
t_svmlin_2 <- svmlin(svmlin_example$X_train[1:50,],
                     svmlin_example$y_train, X_u=svmlin_example$X_train[-c(1:50),],
                     lambda = 10, lambda_u=100, algorithm = 2)

# Calculate Accuracy
mean(predict(t_svmlin_1,svmlin_example$X_test)==svmlin_example$y_test)
mean(predict(t_svmlin_2,svmlin_example$X_test)==svmlin_example$y_test)

data(testdata)
g_svm <- SVM(testdata$X,testdata$y)
g_sup <- svmlin(testdata$X,testdata$y,testdata$X_u,algorithm = 3)
g_semi <- svmlin(testdata$X,testdata$y,testdata$X_u,algorithm = 2)

mean(predict(g_svm,testdata$X_test)==testdata$y_test)
mean(predict(g_sup,testdata$X_test)==testdata$y_test)
mean(predict(g_semi,testdata$X_test)==testdata$y_test)
svmlin_example  
*Test data from the svmlin implementation*

**Description**
Useful for testing the svmlin interface and to serve as an example.

---

svmproblem  
*Train SVM*

**Description**
Train SVM

**Usage**

```
svmproblem(K)
```

**Arguments**

- $K$: kernel

**Value**

- alpha, b, obj

---

testdata  
*Example semi-supervised problem*

**Description**
A list containing a sample from the GenerateSlicedCookie dataset for unit testing and examples.
**threshold**

Refine the prediction to satisfy the balance constraint

**Description**
Refine the prediction to satisfy the balance constraint

**Usage**
\[
\text{threshold}(y_1, \text{options})
\]

**Arguments**
- \(y_1\): predictions
- \(\text{options}\): options passed

**Value**
\(y_2\)

**true_labels**
Access the true labels when they are stored as an attribute in a data frame

**Description**
Access the true labels when they are stored as an attribute in a data frame

**Usage**
\[
\text{true_labels}(df)
\]

**Arguments**
- \(df\): data.frame; data.frame with y_true attribute

**See Also**
Other RSSL utilities: LearningCurveSSL(), SSLDataFrameToMatrices(), add_missinglabels_mar(), df_to_matrices(), measure_accuracy(), missing_labels(), split_dataset_ssl(), split_random()
Transductive SVM classifier using the convex concave procedure

Description

Transductive SVM using the CCCP algorithm as proposed by Collobert et al. (2006) implemented in R using the quadprog package. The implementation does not handle large datasets very well, but can be useful for smaller datasets and visualization purposes.

Usage

```r
TSVM(X, y, X_u, C, Cstar, kernel = kernlab::vanilladot(),
   balancing_constraint = TRUE, s = 0, x_center = TRUE, scale = FALSE,
   eps = 1e-09, max_iter = 20, verbose = FALSE)
```

Arguments

- **X**: matrix; Design matrix for labeled data
- **y**: factor or integer vector; Label vector
- **X_u**: matrix; Design matrix for unlabeled data
- **C**: numeric; Cost parameter of the SVM
- **Cstar**: numeric; Cost parameter of the unlabeled objects
- **kernel**: kernlab::kernel to use
- **balancing_constraint**: logical; Whether a balancing constraint should be enforced that causes the fraction of objects assigned to each label in the unlabeled data to be similar to the label fraction in the labeled data.
- **s**: numeric; parameter controlling the loss function of the unlabeled objects (generally values between -1 and 0)
- **x_center**: logical; Should the features be centered?
- **scale**: If TRUE, apply a z-transform to all observations in X and X_u before running the regression
- **eps**: numeric; Stopping criterion for the maximinimization
- **max_iter**: integer; Maximum number of iterations
- **verbose**: logical; print debugging messages, only works for vanilladot() kernel (default: FALSE)

Details

C is the cost associated with labeled objects, while Cstar is the cost for the unlabeled objects. s control the loss function used for the unlabeled objects: it controls the size of the plateau for the symmetric ramp loss function. The balancing constraint makes sure the label assignments of the unlabeled objects are similar to the prior on the classes that was observed on the labeled data.
References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, USMLeastSquaresClassifier, WellSVM, svmLin()

Examples

library(RSSL)

# Simple example with a few objects
X <- matrix(c(0,0.001,1,-1),nrow=2)
X_u <- matrix(c(-1,-1,0,0,-0.4,-0.5,-0.6,1.2,1.3,1.25),ncol=2)
y <- factor(c(-1,1))
g_sup <- SVM(X,y, scale=FALSE)
g_constraint <- TSVM(X=X,y=y, X_u=X_u, C=1,Cstar=0.1,balancing_constraint = TRUE)
g_noconstraint <- TSVM(X=X,y=y, X_u=X_u, C=1,Cstar=0.1,balancing_constraint = FALSE)
g_lin <- LinearTSVM(X=X,y=y, X_u=X_u,C=1, Cstar=0.1)
w1 <- g_sup@alpha %*% X
w2 <- g_constraint@alpha %*% rbind(X,X_u,X_u,colMeans(X_u))
w3 <- g_noconstraint@alpha %*% rbind(X,X_u,X_u)
w4 <- g_lin@w

plot(X[,1],X[,2],col=factor(y),asp=1,ylim=c(-3,3))
points(X_u[,1],X_u[,2],col="darkgrey",pch=16,cex=1)
abline(-g_sup@bias/w1[2],-w1[1]/w1[2],lty=2)
abline((-1-g_sup@bias)/w1[2],-w1[1]/w1[2],lty=2) # +1 Margin
abline((-1-g_constraint@bias)/w[2][2],-w[1][1]/w[2][2],lty=2) # -1 Margin
abline(-g_constraint@bias/w[2][2],-w[2][1]/w[2][2],lty=1,col="green")
abline(-g_noconstraint@bias/w[3][2],-w[3][1]/w[3][2],lty=1,col="red")
abline(-w4[1]/w4[3],-w4[2]/w4[3],lty=1,lwd=3,col="blue")

# An example
set.seed(42)
data <- generateSlicedCookie(200,expected=TRUE,gap=1)
X <- model.matrix(Class~.-1,data)
y <- factor(data$Class)
problem <- split_dataset_ssl(X,y,frac_ssl=0.98)

X <- problem$X
y <- problem$y
X_u <- problem$X_u
y_e <- unlist(list(problem$y,problem$y_u))
X_e <- rbind(X,X_u)

g_sup <- SVM(X,y,x_center=FALSE,scale=FALSE,C = 10)
g_constraint <- TSVM(X=X,y=y,X_u=X_u,
                        C=10,Cstar=10,balancing_constraint = TRUE,
                        x_center = FALSE,verbose=TRUE)

g_noconstraint <- TSVM(X=X,y=y,X_u=X_u,
                        C=10,Cstar=10,balancing_constraint = FALSE,
                        x_center = FALSE,verbose=TRUE)

g_lin <- LinearTSVM(X=X,y=y,X_u=X_u,C=10,Cstar=10,
                        verbose=TRUE,x_center = FALSE)

g_oracle <- SVM(X_e,y_e,scale=FALSE)

w1 <- c(g_sup@bias,g_sup@alpha %*% X)
w2 <- c(g_constraint@bias,g_constraint@alpha %*% rbind(X,X_u,X_u,colMeans(X_u)))
w3 <- c(g_noconstraint@bias,g_noconstraint@alpha %*% rbind(X,X_u,X_u))
w4 <- g_lin@w
w5 <- c(g_oracle@bias, g_oracle@alpha %*% Xe)

print(sum(abs(w4-w3)))

plot(X[,1],X[,2],col=factor(y),asp=1,ylim=c(-3,3))
points(X_u[,1],X_u[,2],col="darkgrey",pch=16,cex=1)
abline(-w1[1]/w1[3],-w1[2]/w1[3],lty=2)
abline(((1-w1[1])/w1[3]),-w1[2]/w1[3],lty=2)  # +1 Margin
abline(((1-w1[1])/w1[3]),-w1[2]/w1[3],lty=2)  # -1 Margin

# Oracle:
abline(-w5[1]/w5[3],-w5[2]/w5[3],lty=1,col="purple")

# With balancing constraint:
abline(-w2[1]/w2[3],-w2[2]/w2[3],lty=1,col="green")

# Linear TSVM implementation (no constraint):
abline(-w4[1]/w4[3],-w4[2]/w4[3],lty=1,lwd=3,col="blue")

# Without balancing constraint:
abline(-w3[1]/w3[3],-w3[2]/w3[3],lty=1,col="red")
USMLeastSquaresClassifier

Description

This methods uses the closed form solution of the supervised least squares problem, except that the second moment matrix \((X'X)\) is exchanged with a second moment matrix that is estimated based on all data. See for instance Shaffer1991, where in this implementation we use all data to estimate \(E(X'X)\), instead of just the labeled data. This method seems to work best when the data is first centered \(x\_center=TRUE\) and the outputs are scaled using \(y\_scale=TRUE\).

Usage

USMLeastSquaresClassifier(X, y, X_u, lambda = 0, intercept = TRUE, 
        x_center = FALSE, scale = FALSE, y_scale = FALSE, ..., 
        use_Xu_for_scaling = TRUE)

Arguments

- **X** matrix; Design matrix for labeled data
- **y** factor or integer vector; Label vector
- **X_u** matrix; Design matrix for unlabeled data
- **lambda** numeric; L2 regularization parameter
- **intercept** logical; Whether an intercept should be included
- **x_center** logical; Should the features be centered?
- **scale** logical; Should the features be normalized? (default: FALSE)
- **y_scale** logical; whether the target vector should be centered
- **...** Not used
- **use_Xu_for_scaling** logical; whether the unlabeled objects should be used to determine the mean and scaling for the normalization

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier(), LaplacianSVM, LeastSquaresClassifier, LinearDiscriminantClassifierLinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, WellSVM, svmlin()
USMLeastSquaresClassifier-class

USMLeastSquaresClassifier

Description

USMLeastSquaresClassifier

wdbc

wdbc data for unit testing

Description

Useful for testing the S4VM and WellSVM implementations

WellSVM

WellSVM for Semi-supervised Learning

Description

WellSVM is a minimax relaxation of the mixed integer programming problem of finding the optimal labels for the unlabeled data in the SVM objective function. This implementation is a translation of the Matlab implementation of Li (2013) into R.

Usage

WellSVM(X, y, X_u, C1 = 1, C2 = 0.1, gamma = 1, x_center = TRUE, scale = FALSE, use_Xu_for_scaling = FALSE, max_iter = 20)

Arguments

X matrix; Design matrix for labeled data
y factor or integer vector; Label vector
X_u matrix; Design matrix for unlabeled data
C1 double; A regularization parameter for labeled data, default 1;
C2 double; A regularization parameter for unlabeled data, default 0.1;
gamma double; Gaussian kernel parameter, i.e., k(x,y) = exp(-gamma^2||x-y||^2/avg) where avg is the average distance among instances; when gamma = 0, linear kernel is used. default gamma = 1;
x_center logical; Should the features be centered?
scale logical; Should the features be normalized? (default: FALSE)
use_Xu_for_scaling logical; whether the unlabeled objects should be used to determine the mean and scaling for the normalization
max_iter integer; Maximum number of iterations
wellsvm_direct

References


See Also

Other RSSL classifiers: EMLeastSquaresClassifier, EMLinearDiscriminantClassifier, GRFClassifier, ICLeastSquaresClassifier, ICLinearDiscriminantClassifier, KernelLeastSquaresClassifier, LaplacianKernelLeastSquaresClassifier, LaplacianSVM, LeastSquaresClassifier, LLinearDiscriminantClassifier, LinearSVM, LinearTSVM(), LogisticLossClassifier, LogisticRegression, MCLinearDiscriminantClassifier, MCNearestMeanClassifier, MCPLDA, MajorityClassClassifier, NearestMeanClassifier, QuadraticDiscriminantClassifier, S4VM, SVM, SelfLearning, TSVM, USMLeastSquaresClassifier, svmlin()

Examples

```r
library(RSSL)
library(ggplot2)
library(dplyr)

set.seed(1)
df_orig <- generateSlicedCookie(200, expected=TRUE)
df <- df_orig %>%
    add_missinglabels_mar(Class~.,0.98)

classifiers <- list("Well"=WellSVM(Class~.,df,C1 = 1, C2=0.1,
    gamma = 0,x_center=TRUE,scale=TRUE),
    "Sup"=SVM(Class~.,df,C=1,x_center=TRUE,scale=TRUE))

df %>%
    ggplot(aes(x=X1,y=X2,color=Class)) +
    geom_point() +
    coord_equal() +
    stat_classifier(aes(color=..classifier..),
        classifiers = classifiers)
```

wellsvm_direct  wellsvm implements the wellsvm algorithm as shown in [1].

Description

wellsvm implements the wellsvm algorithm as shown in [1].

Usage

```r
wellsvm_direct(x, y, testx, testy, C1 = 1, C2 = 0.1, gamma = 1)
```
Arguments

\( x \) A Nx\(d \) training data matrix, where \( N \) is the number of training instances and \( d \) is the dimension of instance;

\( y \) A Nx1 training label vector, where \( y = 1/-1 \) means positive/negative, and \( y = 0 \) means unlabeled;

\( testx \) A Mx\(d \) testing data matrix, where \( M \) is the number of testing instances;

\( testy \) A Mx1 testing label vector

\( C1 \) A regularization parameter for labeled data, default 1;

\( C2 \) A regularization parameter for unlabeled data, default 0.1;

\( gamma \) Gaussian kernel parameter, i.e., \( k(x,y) = \exp(-\gamma^2\|x-y\|^2/\text{avg}) \) where \( \text{avg} \) is the average distance among instances; when \( \gamma = 0 \), linear kernel is used. default \( \gamma = 1 \);

Value

prediction - A Mx1 predicted testing label vector; accuracy - The accuracy of prediction; cputime - cpu running time;

References


WellSVM\_supervised

A degenerated version of WellSVM where the labels are complete, that is, supervised learning

**Description**

A degenerated version of WellSVM where the labels are complete, that is, supervised learning

**Usage**

WellSVM\_supervised(K0, y, opt, ind\_y)

**Arguments**

- **K0** kernel matrix  
- **y** labels  
- **opt** options  
- **ind\_y** Labeled/Unlabeled indicator

---

wlda

Implements weighted likelihood estimation for LDA

**Description**

Implements weighted likelihood estimation for LDA

**Usage**

wlda(a, w)

**Arguments**

- **a** is the data set  
- **w** is an indicator matrix for the K classes or, potentially, a weight matrix in which the fraction with which a sample belongs to a particular class is indicated

**Value**

m contains the means, p contains the class priors, iW contains the INVERTED within covariance matrix
**wlda_error**

*Measures the expected error of the LDA model defined by m, p, and iW on the data set a, where weights w are potentially taken into account.*

**Description**

Measures the expected error of the LDA model defined by m, p, and iW on the data set a, where weights w are potentially taken into account.

**Usage**

```
wlda_error(m, p, iW, a, w)
```

**Arguments**

- **m**: means
- **p**: class prior
- **iW**: is the inverse of the within covariance matrix
- **a**: design matrix
- **w**: weights

**wlda_loglik**

*Measures the expected log-likelihood of the LDA model defined by m, p, and iW on the data set a, where weights w are potentially taken into account.*

**Description**

Measures the expected log-likelihood of the LDA model defined by m, p, and iW on the data set a, where weights w are potentially taken into account.

**Usage**

```
wlda_loglik(m, p, iW, a, w)
```

**Arguments**

- **m**: means
- **p**: class prior
- **iW**: is the inverse of the within covariance matrix
- **a**: design matrix
- **w**: weights

**Value**

Average log likelihood
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