

Package ‘pros’

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Title Penalized Regression on Steroids

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Description This is a project for STAT8053 at the University of Minnesota. Please see the README on github for compiler flags to improve performance.

Depends R (>= 3.5.0)

SystemRequirements C++11

License GPL-2

Encoding UTF-8

LazyData true

RoxygenNote 6.1.1

R topics documented:

cv.pros	1
predict.cv_pros	2
predict.pros	3
pros	3

Index	5
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cv.pros	<i>Cross-validation</i>
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Description

The cv.pros function is used for K-fold cross-validation.

Usage

```
cv.pros(X, y, K_fold = 10, alpha = c(1, 0, 0, 0, 0, 0),  
        lambdas = c(), step_size, algorithm = "proximal_gradient_cd",  
        max_iter = 10000, tolerance = 10^(-8), random_seed = 0)
```

Arguments

<code>X</code>	is an $n \times m$ -dimensional matrix of the data.
<code>y</code>	is an $n \times m$ -dimensional matrix of the data.
<code>K_fold</code>	is the number of folds in cross-validation.
<code>alpha</code>	is a 6-dimensional vector of the convex combination corresponding to the penalization: <ul style="list-style-type: none"> • α_1 is the l^1 penalty. • α_2 is the l^2 penalty. • α_3 is the l^4 penalty. • α_4 is the l^6 penalty. • α_5 is the l^8 penalty. • α_6 is the l^{10} penalty.
<code>lambdas</code>	is a vector of dual penalization values to be evaluated.
<code>step_size</code>	is a tuning parameter defining the step size. Larger values are more aggressive and smaller values are less aggressive.
<code>algorithm</code>	is the optimization algorithm <ul style="list-style-type: none"> • proximal_gradient_cd uses proximal gradient coordinate descent. • subgradient_cd uses subgradient coordinate descent.
<code>max_iter</code>	is the maximum iterations the algorithm will run regardless of convergence.
<code>tolerance</code>	is the accuracy of the stopping criterion.
<code>random_seed</code>	is the random seed used in the algorithms.

Value

A class `cv_pros`

<code>predict.cv_pros</code>	<i>Cross-validation Prediction</i>
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Description

The prediction function for `cv.pros`.

Usage

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

Arguments

<code>object</code>	an object of class <code>cv_pros</code>
<code>X_new</code>	is an $n \times m$ -dimensional matrix of the data.
<code>...</code>	Other parameters (this is required by R)

Value

A vector of prediction values.

Examples

```

n = 1000
X1 <- rnorm(n)
X2 <- rnorm(n)
y = 3 + 5 * X1 + 0 * X2
X = matrix(c(X1, X2), ncol = 2)
cv = cv.pros(X, y, step_size = .001)
predict(cv, X)

```

predict.pros	<i>Pros Prediction</i>
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Description

The prediction function for pros.

Usage

```

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

```

Arguments

object	an object of class pros
X	is an $n \times m$ -dimensional matrix of the data.
...	Other parameters (this is required by R)

Value

A vector of prediction values.

pros	<i>Pros</i>
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Description

The pros function is used to fit a single regression model with a specified penalization.

Usage

```

pros(X, y, alpha = c(1, 0, 0, 0, 0, 0), lambda, step_size,
     algorithm = "proximal_gradient_cd", max_iter = 10000,
     tolerance = 10^(-8), random_seed = 0)

```

Arguments

<code>X</code>	is an $n \times m$ -dimensional matrix of the data.
<code>y</code>	is an $n \times m$ -dimensional matrix of the data.
<code>alpha</code>	is a 6-dimensional vector of the convex combination corresponding to the penalization: <ul style="list-style-type: none"> • α_1 is the l^1 penalty. • α_2 is the l^2 penalty. • α_3 is the l^4 penalty. • α_4 is the l^6 penalty. • α_5 is the l^8 penalty. • α_6 is the l^{10} penalty.
<code>lambda</code>	is the Lagrangian dual penalization parameter.
<code>step_size</code>	is a tuning parameter defining the step size. Larger values are more aggressive and smaller values are less aggressive.
<code>algorithm</code>	is the optimization algorithm <ul style="list-style-type: none"> • <code>proximal_gradient_cd</code> uses proximal gradient coordinate descent. • <code>subgradient_cd</code> uses subgradient coordinate descent.
<code>max_iter</code>	is the maximum iterations the algorithm will run regardless of convergence.
<code>tolerance</code>	is the accuracy of the stopping criterion.
<code>random_seed</code>	is the random seed used in the algorithms.

Value

A class `pros`

Examples

```
n = 1000
X1 <- rnorm(n)
X2 <- rnorm(n)
y = 3 + 5 * X1 + 0 * X2
X = matrix(c(X1, X2), ncol = 2)
fit = pros(X, y, lambda = 2, step_size = .001)
predict(fit, X)
```

Index

`cv.pros`, [1](#)

`predict.cv_pros`, [2](#)

`predict.pros`, [3](#)

`pros`, [3](#)