Package 'CPGLIB'

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coef.CPGLIB

Coefficients for CPGLIB Object

Description

coef. CPGLIB returns the coefficients for a CPGLIB object.

Usage

```
## S3 method for class 'CPGLIB'
coef(object, groups = NULL, ensemble_average = FALSE, ...)
```

Arguments

object An object of class CPGLIB.

groups The groups in the ensemble for the coefficients. Default is all of the groups in

the ensemble.

ensemble_average

Option to return the average of the coefficients over all the groups in the ensem-

ble. Default is FALSE.

... Additional arguments for compatibility.

Value

The coefficients for the CPGLIB object.

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

cpg

coef.cv.CPGLIB 3

Examples

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 300
beta.active \leftarrow c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 150
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5</pre>
diag(Sigma) <- 1</pre>
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
# CPGLIB - Multiple Groups
cpg.out <- cpg(x.train, y.train,</pre>
                glm_type="Logistic",
                G=5, include_intercept=TRUE,
                alpha_s=3/4, alpha_d=1,
                lambda_sparsity=0.01, lambda_diversity=1,
                tolerance=1e-5, max_iter=1e5)
# Coefficients for each group
cpg.coef <- coef(cpg.out, ensemble_average = FALSE)</pre>
```

coef.cv.CPGLIB

Coefficients for cv.CPGLIB Object

Description

coef.cv.CPGLIB returns the coefficients for a cv.CPGLIB object.

Usage

```
## S3 method for class 'cv.CPGLIB'
coef(object, groups = NULL, ensemble_average = FALSE, ...)
```

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Arguments

object An object of class cv.CPGLIB.

groups The groups in the ensemble for the coefficients. Default is all of the groups in

the ensemble.

ensemble_average

Option to return the average of the coefficients over all the groups in the ensem-

ble. Default is FALSE.

... Additional arguments for compatibility.

Value

The coefficients for the cv.CPGLIB object. Default is FALSE.

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

cv.cpg

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 300
beta.active <- c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 150
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5</pre>
diag(Sigma) <- 1
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
mean(y.test)
# CV CPGLIB - Multiple Groups
```

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coef.cv.ProxGrad

Coefficients for cv.ProxGrad Object

Description

coef.cv.ProxGrad returns the coefficients for a cv.ProxGrad object.

Usage

```
## S3 method for class 'cv.ProxGrad'
coef(object, ...)
```

Arguments

object An object of class cv.ProxGrad.

. . . Additional arguments for compatibility.

Value

The coefficients for the cv.ProxGrad object.

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

```
cv.ProxGrad
```

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Examples

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 1000
beta.active <- c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 100
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5
diag(Sigma) <- 1</pre>
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
# CV ProxGrad - Single Group
proxgrad.out <- cv.ProxGrad(x.train, y.train,</pre>
                              glm_type = "Logistic",
                              include_intercept = TRUE,
                              alpha_s = 3/4,
                              n_lambda_sparsity = 100,
                              tolerance = 1e-5, max_iter = 1e5)
# Coefficients
coef(proxgrad.out)
```

coef.ProxGrad

Coefficients for ProxGrad Object

Description

coef.ProxGrad returns the coefficients for a ProxGrad object.

Usage

```
## S3 method for class 'ProxGrad'
coef(object, ...)
```

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Arguments

object An object of class ProxGrad.... Additional arguments for compatibility.

Value

The coefficients for the ProxGrad object.

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

ProxGrad

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 1000
beta.active < c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 100
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5</pre>
diag(Sigma) <- 1
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test \leftarrow mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
# ProxGrad - Single Group
proxgrad.out <- ProxGrad(x.train, y.train,</pre>
                           glm_type = "Logistic",
                           include_intercept = TRUE,
                           alpha_s = 3/4,
                           lambda_sparsity = 0.01,
                           tolerance = 1e-5, max_iter = 1e5)
```

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```
# Coefficients
coef(proxgrad.out)
```

cpg

Competing Proximal Gradients Library for Ensembles of Generalized Linear Models

Description

cpg computes the coefficients for ensembles of generalized linear models via competing proximal gradients.

Usage

```
cpg(
   x,
   y,
   glm_type = c("Linear", "Logistic")[1],
   G = 5,
   include_intercept = TRUE,
   alpha_s = 3/4,
   alpha_d = 1,
   lambda_sparsity,
   lambda_diversity,
   tolerance = 1e-08,
   max_iter = 1e+05
)
```

Arguments

max_iter

Design matrix. Х Response vector. У glm_type Description of the error distribution and link function to be used for the model. Must be one of "Linear" or "Logistic". Default is "Linear". Number of groups in the ensemble. include_intercept Argument to determine whether there is an intercept. Default is TRUE. Sparsity mixing parmeter. Default is 3/4. alpha_s alpha_d Diversity mixing parameter. Default is 1. lambda_sparsity Sparsity tuning parameter value. lambda_diversity Diversity tuning parameter value. tolerance Convergence criteria for the coefficients. Default is 1e-8.

Maximum number of iterations in the algorithm. Default is 1e5.

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Value

An object of class cpg

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

```
coef.CPGLIB, predict.CPGLIB
```

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 300
beta.active \leftarrow c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 150
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5</pre>
diag(Sigma) <- 1</pre>
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
# CPGLIB - Multiple Groups
cpg.out <- cpg(x.train, y.train,</pre>
                glm_type = "Logistic",
                G = 5, include_intercept = TRUE,
                alpha_s = 3/4, alpha_d = 1,
                lambda_sparsity = 0.01, lambda_diversity = 1,
                tolerance = 1e-5, max_iter = 1e5)
# Predictions
cpg.prob <- predict(cpg.out, newx = x.test, type = "prob",</pre>
                     groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
cpg.class <- predict(cpg.out, newx = x.test, type = "prob",</pre>
                      groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
plot(prob.test, cpg.prob, pch = 20)
```

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```
abline(h = 0.5,v = 0.5)
mean((prob.test-cpg.prob)^2)
mean(abs(y.test-cpg.class))
```

cv.cpg

Competing Proximal Gradients Library for Ensembles of Generalized Linear Models - Cross-Validation

Description

cv.cpg computes and cross-validates the coefficients for ensembles of generalized linear models via competing proximal gradients.

Usage

```
cv.cpg(
   x,
   y,
   glm_type = c("Linear", "Logistic")[1],
   G = 5,
   full_diversity = FALSE,
   include_intercept = TRUE,
   alpha_s = 3/4,
   alpha_d = 1,
   n_lambda_sparsity = 100,
   n_lambda_diversity = 100,
   tolerance = 1e-08,
   max_iter = 1e+05,
   n_folds = 10,
   n_threads = 1
)
```

Arguments

```
x Design matrix.

y Response vector.

glm_type Description of the error distribution and link function to be used for the model.

Must be one of "Linear" or "Logistic". Default is "Linear".

G Number of groups in the ensemble.

full_diversity Argument to determine if the overlap between the models should be zero. Default is FALSE.

include_intercept
```

Argument to determine whether there is an intercept. Default is TRUE.

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```
Sparsity mixing parmeter. Default is 3/4.
alpha_s
                  Diversity mixing parameter. Default is 1.
alpha_d
n_lambda_sparsity
                  Number of candidates for sparsity tuning parameter. Default is 100.
n_lambda_diversity
                  Number of candidates for diveristy tuning parameter. Default is 100.
tolerance
                  Convergence criteria for the coefficients. Default is 1e-8.
max_iter
                  Maximum number of iterations in the algorithm. Default is 1e5.
n_folds
                  Number of cross-validation folds. Default is 10.
n_threads
                  Number of threads. Default is a single thread.
```

Value

An object of class cv.cpg

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

```
coef.cv.CPGLIB, predict.cv.CPGLIB
```

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 300
beta.active <- c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 150
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5
diag(Sigma) <- 1
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test \leftarrow mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
```

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```
# CV CPGLIB - Multiple Groups
cpg.out <- cv.cpg(x.train, y.train,</pre>
                  glm_type = "Logistic",
                  G = 5, include_intercept = TRUE,
                  alpha_s = 3/4, alpha_d = 1,
                  n_lambda_sparsity = 100, n_lambda_diversity = 100,
                  tolerance = 1e-5, max_iter = 1e5)
# Predictions
cpg.prob <- predict(cpg.out, newx = x.test, type = "prob",</pre>
                     groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
cpg.class <- predict(cpg.out, newx = x.test, type = "class",</pre>
                      groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
plot(prob.test, cpg.prob, pch = 20)
abline(h = 0.5, v = 0.5)
mean((prob.test-cpg.prob)^2)
mean(abs(y.test-cpg.class))
```

cv.ProxGrad

Generalized Linear Models via Proximal Gradients - Cross-validation

Description

cv.ProxGrad computes and cross-validates the coefficients for generalized linear models using proximal gradients.

Usage

```
cv.ProxGrad(
    x,
    y,
    glm_type = c("Linear", "Logistic")[1],
    include_intercept = TRUE,
    alpha_s = 3/4,
    n_lambda_sparsity = 100,
    tolerance = 1e-08,
    max_iter = 1e+05,
    n_folds = 10,
    n_threads = 1
)
```

Arguments

x Design matrix.

y Response vector.

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glm_type Description of the error distribution and link function to be used for the model.

Must be one of "Linear" or "Logistic". Default is "Linear".

include_intercept

Argument to determine whether there is an intercept. Default is TRUE.

alpha_s Elastic net mixing parmeter. Default is 3/4.

n_lambda_sparsity

Sparsity tuning parameter value. Default is 100.

tolerance Convergence criteria for the coefficients. Default is 1e-8.

max_iter Maximum number of iterations in the algorithm. Default is 1e5.

n_folds Number of cross-validation folds. Default is 10.n_threads Number of threads. Default is a single thread.

Value

An object of class cv.ProxGrad

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

```
coef.cv.ProxGrad, predict.cv.ProxGrad
```

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 1000
beta.active <- c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 100
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5
diag(Sigma) <- 1
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test \leftarrow mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
```

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predict.CPGLIB

Predictions for CPGLIB Object

Description

predict.CPGLIB returns the predictions for a CPGLIB object.

Usage

```
## S3 method for class 'CPGLIB'
predict(
  object,
  newx,
  groups = NULL,
  ensemble_type = c("Model-Avg", "Coef-Avg", "Weighted-Prob", "Majority-Vote")[1],
  class_type = c("prob", "class")[1],
  ...
)
```

Arguments

object An object of class CPGLIB.

newx New data for predictions.

groups The groups in the ensemble for the predictions. Default is all of the groups in

the ensemble.

ensemble_type The type of ensembling function for the models. Options are "Model-Avg",

"Coef-Avg" or "Weighted-Prob" for classifications predictions. Default is "Model-

Avg".

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```
class_type The type of predictions for classification. Options are "prob" and "class". Default is "prob".... Additional arguments for compatibility.
```

Value

The predictions for the CPGLIB object.

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

cpg

Examples

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 300
beta.active \leftarrow c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 150
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5</pre>
diag(Sigma) <- 1</pre>
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
# CPGLIB - Multiple Groups
cpg.out <- cpg(x.train, y.train,</pre>
                glm_type = "Logistic",
                G = 5, include_intercept = TRUE,
                alpha_s = 3/4, alpha_d = 1,
                lambda_sparsity = 0.01, lambda_diversity = 1,
                tolerance = 1e-5, max_iter = 1e5)
```

Predictions

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predict.cv.CPGLIB

Predictions for cv.ProxGrad Object

Description

predict.cv.CPGLIB returns the predictions for a ProxGrad object.

Usage

```
## S3 method for class 'cv.CPGLIB'
predict(
  object,
  newx,
  groups = NULL,
  ensemble_type = c("Model-Avg", "Coef-Avg", "Weighted-Prob", "Majority-Vote")[1],
    class_type = c("prob", "class")[1],
    ...
)
```

Arguments

object An object of class cv.CPGLIB.

New data for predictions.

groups The groups in the ensemble for the predictions. Default is all of the groups in the ensemble.

ensemble_type The type of ensembling function for the models. Options are "Model-Avg", "Coef-Avg" or "Weighted-Prob" for classifications predictions. Default is "Model-Avg".

class_type The type of predictions for classification. Options are "prob" and "class". Default is "prob".

Additional arguments for compatibility.

Value

The predictions for the cv.CPGLIB object.

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Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

```
cv.cpg
```

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 300
beta.active < c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 150
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5
diag(Sigma) <- 1</pre>
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test \leftarrow mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
mean(y.test)
# CV CPGLIB - Multiple Groups
cpg.out <- cv.cpg(x.train, y.train,</pre>
                   glm_type = "Logistic",
                   G = 5, include_intercept = TRUE,
                   alpha_s = 3/4, alpha_d = 1,
                   n_lambda_sparsity = 100, n_lambda_diversity = 100,
                   tolerance = 1e-5, max_iter = 1e5)
# Predictions
cpg.prob <- predict(cpg.out, newx = x.test, type = "prob",</pre>
                     groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
cpg.class <- predict(cpg.out, newx = x.test, type = "class",</pre>
                      groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
plot(prob.test, cpg.prob, pch = 20)
abline(h = 0.5, v = 0.5)
mean((prob.test-cpg.prob)^2)
mean(abs(y.test-cpg.class))
```

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predict.cv.ProxGrad Pred

Predictions for cv.ProxGrad Object

Description

predict.cv.ProxGrad returns the predictions for a ProxGrad object.

Usage

```
## S3 method for class 'cv.ProxGrad'
predict(object, newx, type = c("prob", "class")[1], ...)
```

Arguments

object An object of class cv.ProxGrad.

newx New data for predictions.

type The type of predictions for binary response. Options are "prob" (default) and

"class".

... Additional arguments for compatibility.

Value

The predictions for the cv.ProxGrad object.

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

```
cv.ProxGrad
```

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 1000
beta.active <- c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 100</pre>
```

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```
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5</pre>
diag(Sigma) <- 1</pre>
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
# CV ProxGrad - Single Group
proxgrad.out <- cv.ProxGrad(x.train, y.train,</pre>
                              glm_type = "Logistic",
                              include_intercept = TRUE,
                              alpha_s = 3/4,
                              n_{\text{lambda\_sparsity}} = 100,
                              tolerance = 1e-5, max_iter = 1e5)
# Predictions
proxgrad.prob <- predict(proxgrad.out, newx = x.test, type = "prob")</pre>
proxgrad.class <- predict(proxgrad.out, newx = x.test, type = "class")</pre>
plot(prob.test, proxgrad.prob, pch = 20)
abline(h = 0.5, v = 0.5)
mean((prob.test-proxgrad.prob)^2)
mean(abs(y.test-proxgrad.class))
```

predict.ProxGrad

Predictions for ProxGrad Object

Description

predict.ProxGrad returns the predictions for a ProxGrad object.

Usage

```
## S3 method for class 'ProxGrad'
predict(object, newx, type = c("prob", "class")[1], ...)
```

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Arguments

object An object of class ProxGrad

newx New data for predictions.

type The type of predictions for binary response. Options are "prob" (default) and "class".

Additional arguments for compatibility.

Value

The predictions for the ProxGrad object.

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

ProxGrad

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 1000
beta.active <- c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 100
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5</pre>
diag(Sigma) <- 1
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test \leftarrow mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
# ProxGrad - Single Group
proxgrad.out <- ProxGrad(x.train, y.train,</pre>
                           glm_type = "Logistic",
                           include_intercept = TRUE,
```

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ProxGrad

Generalized Linear Models via Proximal Gradients

Description

ProxGrad computes the coefficients for generalized linear models using proximal gradients.

Usage

```
ProxGrad(
    x,
    y,
    glm_type = c("Linear", "Logistic")[1],
    include_intercept = TRUE,
    alpha_s = 3/4,
    lambda_sparsity,
    tolerance = 1e-08,
    max_iter = 1e+05
)
```

Arguments

max_iter

x Design matrix.

y Response vector.

glm_type Description of the error distribution and link function to be used for the model.

Must be one of "Linear" or "Logistic" . Default is "Linear".

include_intercept
Argument to determine whether there is an intercept. Default is TRUE.

alpha_s Elastic net mixing parmeter. Default is 3/4.

lambda_sparsity
Sparsity tuning parameter value.

tolerance Convergence criteria for the coefficients. Default is 1e-8.

Maximum number of iterations in the algorithm. Default is 1e5.

ProxGrad ProxGrad

Value

An object of class ProxGrad.

Author(s)

Anthony-Alexander Christidis, <anthony.christidis@stat.ubc.ca>

See Also

```
coef.ProxGrad, predict.ProxGrad
```

```
# Data simulation
set.seed(1)
n <- 50
N <- 2000
p <- 1000
beta.active \leftarrow c(abs(runif(p, 0, 1/2))*(-1)^rbinom(p, 1, 0.3))
# Parameters
p.active <- 100
beta <- c(beta.active[1:p.active], rep(0, p-p.active))</pre>
Sigma <- matrix(0, p, p)</pre>
Sigma[1:p.active, 1:p.active] <- 0.5</pre>
diag(Sigma) <- 1</pre>
# Train data
x.train <- mvnfast::rmvn(n, mu = rep(0, p), sigma = Sigma)
prob.train <- exp(x.train %*% beta)/</pre>
               (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)</pre>
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/</pre>
              (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)</pre>
# ProxGrad - Single Group
proxgrad.out <- ProxGrad(x.train, y.train,</pre>
                          glm_type = "Logistic",
                           include_intercept = TRUE,
                           alpha_s = 3/4,
                           lambda_sparsity = 0.01,
                           tolerance = 1e-5, max_iter = 1e5)
# Predictions
proxgrad.prob <- predict(proxgrad.out, newx = x.test, type = "prob")</pre>
proxgrad.class <- predict(proxgrad.out, newx = x.test, type = "class")</pre>
plot(prob.test, proxgrad.prob, pch = 20)
abline(h = 0.5, v = 0.5)
mean((prob.test-proxgrad.prob)^2)
```

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mean(abs(y.test-proxgrad.class))

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