

# Package ‘CPGLIB’

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**Type** Package

**Title** Competing Proximal Gradients Library

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**Description** Functions to generate ensembles of generalized linear models using competing proximal gradients. The optimal sparsity and diversity tuning parameters are selected via an alternating grid search.

**License** GPL (>= 2)

**Biarch** true

**Imports** Rcpp (>= 1.0.3)

**LinkingTo** Rcpp, RcppArmadillo

**RoxygenNote** 7.1.1

**Suggests** testthat, vctrs, mvnfast

**NeedsCompilation** yes

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coef.CPGLIB	<i>Coefficients for CPGLIB Object</i>
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## 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 ensemble. 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](#)

**Examples**

```

# 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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)

# CPGLIB - Multiple Groups
cpg.out <- cpg(x.train, y.train,
  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)

```

---

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, ...)

```

**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 ensemble. 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](#)

**Examples**

```
# 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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)
mean(y.test)

# CV CPGLIB - Multiple Groups
```

```
cpg.out <- cv.cpg(x.train, y.train,
  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)
cpg.coef <- coef(cpg.out)

# Coefficients for each group
cpg.coef <- coef(cpg.out, ensemble_average = FALSE)
```

---

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](#)

**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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)

# CV ProxGrad - Single Group
proxgrad.out <- cv.ProxGrad(x.train, y.train,
  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, ...)

```

**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](#)

**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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)

# ProxGrad - Single Group
proxgrad.out <- ProxGrad(x.train, y.train,
  glm_type = "Logistic",
  include_intercept = TRUE,
  alpha_s = 3/4,
  lambda_sparsity = 0.01,
  tolerance = 1e-5, max_iter = 1e5)
```

```
# 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

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.
include_intercept	Argument to determine whether there is an intercept. Default is TRUE.
alpha_s	Sparsity mixing parameter. Default is 3/4.
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.
max_iter	Maximum number of iterations in the algorithm. Default is 1e5.



**Value**

An object of class cpg

**Author(s)**

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

**See Also**

[coef.CPGLIB](#), [predict.CPGLIB](#)

**Examples**

```
# 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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)

# CPGLIB - Multiple Groups
cpg.out <- cpg(x.train, y.train,
  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",
  groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
cpg.class <- predict(cpg.out, newx = x.test, type = "prob",
  groups = 1:cpg.out$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.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.

alpha_s	Sparsity mixing parameter. Default is 3/4.
alpha_d	Diversity mixing parameter. Default is 1.
n_lambda_sparsity	Number of candidates for sparsity tuning parameter. Default is 100.
n_lambda_diversity	Number of candidates for diversity 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](#)

**Examples**

```
# 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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)
```

```

# CV CPGLIB - Multiple Groups
cpg.out <- cv.cpg(x.train, y.train,
  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",
  groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
cpg.class <- predict(cpg.out, newx = x.test, type = "class",
  groups = 1:cpg.out$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.

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 parameter. 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](#)

**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))
Sigma <- matrix(0, p, p)
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) /
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta) /
  (1+exp(x.test %*% beta))
```

```

y.test <- rbinom(N, 1, prob.test)

# ProxGrad - Single Groups
proxgrad.out <- cv.ProxGrad(x.train, y.train,
                           glm_type = "Logistic",
                           include_intercept = TRUE,
                           alpha_s = 3/4,
                           n_lambda_sparsity = 100,
                           tolerance = 1e-5, max_iter = 1e5)

# Predictions
proxgrad.prob <- predict(proxgrad.out, newx = x.test, type = "prob")
proxgrad.class <- predict(proxgrad.out, newx = x.test, type = "class")
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.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".

`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 <- 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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)

# CPGLIB - Multiple Groups
cpg.out <- cpg(x.train, y.train,
  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",
                  groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
cpg.class <- predict(cpg.out, newx = x.test, type = "prob",
                   groups = 1:cpg.out$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))

```

---

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.
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".
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.



**Author(s)**

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

**See Also**

[cv.cpg](#)

**Examples**

```
# 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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)
mean(y.test)

# CV CPGLIB - Multiple Groups
cpg.out <- cv.cpg(x.train, y.train,
  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",
  groups = 1:cpg.out$G, ensemble_type = "Model-Avg")
cpg.class <- predict(cpg.out, newx = x.test, type = "class",
  groups = 1:cpg.out$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))
```

---

predict.cv.ProxGrad    *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](#)

### 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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %**% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %**% beta)/
  (1+exp(x.test %**% beta))
y.test <- rbinom(N, 1, prob.test)

# CV ProxGrad - Single Group
proxgrad.out <- cv.ProxGrad(x.train, y.train,
  glm_type = "Logistic",
  include_intercept = TRUE,
  alpha_s = 3/4,
  n_lambda_sparsity = 100,
  tolerance = 1e-5, max_iter = 1e5)

# Predictions
proxgrad.prob <- predict(proxgrad.out, newx = x.test, type = "prob")
proxgrad.class <- predict(proxgrad.out, newx = x.test, type = "class")
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], ...)

```

**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](#)

**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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)

# ProxGrad - Single Group
proxgrad.out <- ProxGrad(x.train, y.train,
  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")
proxgrad.class <- predict(proxgrad.out, newx = x.test, type = "class")
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))

```

---

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

<code>x</code>	Design matrix.
<code>y</code>	Response vector.
<code>glm_type</code>	Description of the error distribution and link function to be used for the model. Must be one of "Linear" or "Logistic" . Default is "Linear".
<code>include_intercept</code>	Argument to determine whether there is an intercept. Default is TRUE.
<code>alpha_s</code>	Elastic net mixing parameter. Default is 3/4.
<code>lambda_sparsity</code>	Sparsity tuning parameter value.
<code>tolerance</code>	Convergence criteria for the coefficients. Default is 1e-8.
<code>max_iter</code>	Maximum number of iterations in the algorithm. Default is 1e5.

**Value**

An object of class ProxGrad.

**Author(s)**

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

**See Also**

[coef.ProxGrad](#), [predict.ProxGrad](#)

**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))
Sigma <- matrix(0, p, p)
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)/
  (1+exp(x.train %*% beta))
y.train <- rbinom(n, 1, prob.train)
# Test data
x.test <- mvnfast::rmvn(N, mu = rep(0, p), sigma = Sigma)
prob.test <- exp(x.test %*% beta)/
  (1+exp(x.test %*% beta))
y.test <- rbinom(N, 1, prob.test)

# ProxGrad - Single Group
proxgrad.out <- ProxGrad(x.train, y.train,
  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")
proxgrad.class <- predict(proxgrad.out, newx = x.test, type = "class")
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))
```

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