

Package ‘yager’

October 25, 2020

Title Yet Another General Regression Neural Network

Version 0.1.1

Author WenSui Liu

Maintainer WenSui Liu <liuwensui@gmail.com>

Description Another implementation of general regression neural network in R based on Specht (1991) <DOI:10.1109/72.97934>. It is applicable to the functional approximation or the classification.

URL <https://github.com/statcompute/yager>

Depends R (>= 3.6.0)

Imports datasets, stats, randtoolbox, lhs, MLmetrics, graphics,
parallel

License GPL (>= 2)

Encoding UTF-8

LazyData true

RoxygenNote 7.1.1

NeedsCompilation no

Repository CRAN

Date/Publication 2020-10-25 18:30:02 UTC

R topics documented:

folds	2
gen_latin	2
gen_sobol	3
gen_unifm	4
grnn.fit	4
grnn.imp	5
grnn.margin	6
grnn.optmiz_auc	6
grnn.parpred	7
grnn.partial	8

grnn.pfi	9
grnn.predict	10
grnn.predone	10
grnn.search_auc	11
grnn.search_rsq	12
grnn.x_imp	13
grnn.x_pfi	14

Index	15
--------------	-----------

folds	<i>Generate a list of index for the n-fold cross-validation</i>
-------	---

Description

The function `folds` generates a list of index for the n-fold cross-validation

Usage

```
folds(idx, n, seed = 1)
```

Arguments

idx	A vector of index list
n	The number of n folds
seed	The seed value to generate random n-fold index

Value

A list of n-fold index

Examples

```
folds(seq(10), n = 3, seed = 2020)
```

gen_latin	<i>Generate random numbers of latin hypercube sampling</i>
-----------	--

Description

The function `gen_latin` generates a vector of random numbers by latin hypercube sampling

Usage

```
gen_latin(min = 0, max = 1, n, seed = 1)
```

Arguments

min	The minimum value of random numbers
max	The maximum value of random numbers
n	The number of random numbers to generate
seed	The seed value of random number generation

Value

A vector of random numbers bounded by the min and max

Examples

```
gen_latin(0, 1, 10, 2020)
```

gen_sobol	<i>Generate sobol sequence</i>
-----------	--------------------------------

Description

The function gen_sobol generates a vector of scrambled sobol sequence

Usage

```
gen_sobol(min = 0, max = 1, n, seed = 1)
```

Arguments

min	The minimum value of random numbers
max	The maximum value of random numbers
n	The number of random numbers to generate
seed	The seed value of random number generation

Value

A vector of sobol sequence bounded by the min and max

Examples

```
gen_sobol(0, 1, 10, 2020)
```

gen_unifm	<i>Generate Uniform random numbers</i>
-----------	--

Description

The function `gen_unifm` generates a vector of uniform random numbers

Usage

```
gen_unifm(min = 0, max = 1, n, seed = 1)
```

Arguments

<code>min</code>	The minimum value of random numbers
<code>max</code>	The maximum value of random numbers
<code>n</code>	The number of random numbers to generate
<code>seed</code>	The seed value of random number generation

Value

A vector of uniform random numbers bounded by the min and max

Examples

```
gen_unifm(0, 1, 10, 2020)
```

grnn.fit	<i>Create a general regression neural network</i>
----------	---

Description

The function `grnn.fit` creates a general regression neural network (GRNN)

Usage

```
grnn.fit(x, y, sigma = 1, w = rep(1, length(y)))
```

Arguments

<code>x</code>	The matrix of predictors
<code>y</code>	The vector of response variable
<code>sigma</code>	The scalar of smoothing parameter
<code>w</code>	The vector of weights with default = 1 for each record

Value

A general regression neural network object

References

Donald Specht. (1991). A General Regression Neural Network.

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
```

grnn.imp

Derive the importance rank of all predictors used in the GRNN

Description

The function `grnn.imp` derives the importance rank of all predictors used in the GRNN. It essentially is a wrapper around the function `grnn.x_imp`.

Usage

```
grnn.imp(net, class = FALSE)
```

Arguments

<code>net</code>	The GRNN object generated by <code>grnn.fit()</code>
<code>class</code>	TRUE or FALSE, whether it is for the classification or not

Value

A dataframe with important values of all predictors in the GRNN

See Also

[grnn.x_imp](#)

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:3])
gnet <- grnn.fit(x = X, y = Y)
## Not run:
grnn.imp(net = gnet, class = TRUE)

## End(Not run)
```

grnn.margin	<i>Derive the marginal effect of a predictor used in a GRNN</i>
-------------	---

Description

The function `grnn.margin` derives the marginal effect of a predictor used in a GRNN by assuming mean values for the rest predictors

Usage

```
grnn.margin(net, i, plot = TRUE)
```

Arguments

<code>net</code>	The GRNN object generated by <code>grnn.fit()</code>
<code>i</code>	The <i>i</i> th predictor in the GRNN
<code>plot</code>	TRUE or FALSE to plot the marginal effect

Value

A plot of the marginal effect or a dataframe of the marginal effect

See Also

[grnn.partial](#)

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
grnn.margin(gnet, 1, plot = FALSE)
```

grnn.optimiz_auc	<i>Optimize the optimal value of GRNN smoothing parameter based on AUC</i>
------------------	--

Description

The function `grnn.optimiz_auc` optimize the optimal value of GRNN smoothing parameter by cross-validation. It is applicable to the classification.

Usage

```
grnn.optimiz_auc(net, lower = 0, upper, nfolds = 4, seed = 1, method = 1)
```

Arguments

net	A GRNN object generated by grnn.fit()
lower	A scalar for the lower bound of the smoothing parameter
upper	A scalar for the upper bound of the smoothing parameter
nfolds	A scalar for the number of n-fold, 4 by default
seed	The seed value for the n-fold cross-validation, 1 by default
method	A scalar referring to the optimization method, 1 for Golden section searc and 2 for Brent's method

Value

The best outcome

See Also

[grnn.search_auc](#)

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
## Not run:
grnn.optmiz_auc(net = gnet, lower = 3, upper = 7, nfolds = 2)

## End(Not run)
```

grnn.parpred

Calculate predicted values of GRNN by using parallelism

Description

The function grnn.parpred calculates a vector of GRNN predicted values based on an input matrix

Usage

```
grnn.parpred(net, x)
```

Arguments

net	The GRNN object generated by grnn.fit()
x	The matrix of input predictors

Value

A vector of predicted values

See Also[grnn.predict](#)**Examples**

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
grnn.parpred(gnet, X[seq(5), ])
```

`grnn.partial`*Derive the partial effect of a predictor used in a GRNN*

Description

The function `grnn.partial` derives the partial effect of a predictor used in a GRNN by average-out values of the rest predictors.

Usage

```
grnn.partial(net, i, plot = TRUE)
```

Arguments

<code>net</code>	The GRNN object generated by <code>grnn.fit()</code>
<code>i</code>	The <i>i</i> th predictor in the GRNN
<code>plot</code>	TRUE or FALSE to plot the partial effect

Value

A plot of the partial effect or a dataframe of the partial effect

See Also[grnn.margin](#)**Examples**

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
## Not run:
grnn.partial(gnet, 1, plot = FALSE)

## End(Not run)
```

`grnn.pfi`*Derive the PFI rank of all predictors used in the GRNN*

Description

The function `grnn.pfi` derives the PFI rank of all predictors used in the GRNN. It essentially is a wrapper around the function `grnn.x_pfi`.

Usage

```
grnn.pfi(net, class = FALSE, ntry = 1000, seed = 1)
```

Arguments

<code>net</code>	The GRNN object generated by <code>grnn.fit()</code>
<code>class</code>	TRUE or FALSE, whether it is for the classification or not
<code>ntry</code>	The number of random permutations to try, 1e3 times by default
<code>seed</code>	The seed value for the random permutation

Value

A dataframe with PFI values of all predictors in the GRNN

See Also

[grnn.x_pfi](#)

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:3])
gnet <- grnn.fit(x = X, y = Y)
## Not run:
grnn.pfi(net = gnet, class = TRUE)

## End(Not run)
```

grnn.predict	<i>Calculate predicted values of GRNN</i>
--------------	---

Description

The function `grnn.predict` calculates a vector of GRNN predicted values based on an input matrix

Usage

```
grnn.predict(net, x)
```

Arguments

<code>net</code>	The GRNN object generated by <code>grnn.fit()</code>
<code>x</code>	The matrix of input predictors

Value

A vector of predicted values

See Also

[grnn.predone](#)

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
grnn.predict(gnet, X[seq(5), ])
```

grnn.predone	<i>Calculate a predicted value of GRNN</i>
--------------	--

Description

The function `grnn.predone` calculates a predicted value of GRNN based on an input vector

The function `grnn.predone` calculates a predicted value of GRNN based on an input vector

Usage

```
grnn.predone(net, x, type = 1)
```

```
grnn.predone(net, x, type = 1)
```

Arguments

net	The GRNN object generated by grnn.fit()
x	The vector of input predictors
type	A scalar, 1 for euclidean distance and 2 for manhattan distance

Value

A scalar of the predicted value

A scalar of the predicted value

References

Donald Specht. (1991). A General Regression Neural Network.

Donald Specht. (1991). A General Regression Neural Network.

See Also

[grnn.fit](#)

[grnn.fit](#)

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
for (i in seq(5)) print(grnn.predone(gnet, X[i, ]))
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
for (i in seq(5)) print(grnn.predone(gnet, X[i, ]))
```

grnn.search_auc	<i>Search for the optimal value of GRNN smoothing parameter based on AUC</i>
-----------------	--

Description

The function `grnn.search_auc` searches for the optimal value of GRNN smoothing parameter by cross-validation. It is applicable to the classification.

Usage

```
grnn.search_auc(net, sigmas, nfold = 4, seed = 1)
```

Arguments

net	A GRNN object generated by grnn.fit()
sigmas	A numeric vector to search for the best smoothing parameter
nfolds	A scalar for the number of n-fold, 4 by default
seed	The seed value for the n-fold cross-validation, 1 by default

Value

The list of all searching outcomes and the best outcome

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
grnn.search_auc(net = gnet, sigmas = c(3, 5, 7), nfolds = 2)
```

grnn.search_rsq	<i>Search for the optimal value of GRNN smoothing parameter based on r-square</i>
-----------------	---

Description

The function grnn.search_rsq searches for the optimal value of GRNN smoothing parameter by cross-validation. It is applicable to the functional approximation

Usage

```
grnn.search_rsq(net, sigmas, nfolds = 4, seed = 1)
```

Arguments

net	A GRNN object generated by grnn.fit()
sigmas	A numeric vector to search for the best smoothing parameter
nfolds	A scalar for the number of n-fold, 4 by default
seed	The seed value for the n-fold cross-validation, 1 by default

Value

The list of all searching outcomes and the best outcome

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
grnn.search_rsqr(net = gnet, sigmas = seq(3), nfolds = 2)
```

grnn.x_imp

Derive the importance of a predictor used in the GRNN

Description

The function `grnn.x_imp` derives the importance of a predictor used in the GRNN by using the loss of predictability after eliminating the impact of the predictor in interest.

Usage

```
grnn.x_imp(net, i, class = FALSE)
```

Arguments

<code>net</code>	The GRNN object generated by <code>grnn.fit()</code>
<code>i</code>	The <i>i</i> th predictor in the GRNN
<code>class</code>	TRUE or FALSE, whether it is for the classification or not

Value

A vector with the variable name and two values of importance measurements, namely "imp1" and "imp2". The "imp1" measures the loss of predictability after replacing all values of the predictor with its mean. The "imp2" measures the loss of predictability after dropping the predictor from the GRNN.

See Also

[grnn.x_pfi](#)

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
grnn.x_imp(net = gnet, 1)
```

grnn.x_pfi	<i>Derive the permutation feature importance of a predictor used in the GRNN</i>
------------	--

Description

The function `grnn.x_pfi` derives the permutation feature importance (PFI) of a predictor used in the GRNN

Usage

```
grnn.x_pfi(net, i, class = FALSE, ntry = 1000, seed = 1)
```

Arguments

<code>net</code>	The GRNN object generated by <code>grnn.fit()</code>
<code>i</code>	The <i>i</i> th predictor in the GRNN
<code>class</code>	TRUE or FALSE, whether it is for the classification or not
<code>ntry</code>	The number of random permutations to try, 1e3 times by default
<code>seed</code>	The seed value for the random permutation

Value

A vector with the variable name and the PFI value.

See Also

[grnn.x_imp](#)

Examples

```
data(iris, package = "datasets")
Y <- ifelse(iris[, 5] == "setosa", 1, 0)
X <- scale(iris[, 1:4])
gnet <- grnn.fit(x = X, y = Y)
grnn.x_pfi(net = gnet, 1)
```

Index

`folds`, 2

`gen_latin`, 2

`gen_sobol`, 3

`gen_unifm`, 4

`grnn.fit`, 4, 11

`grnn.imp`, 5

`grnn.margin`, 6, 8

`grnn.optimiz_auc`, 6

`grnn.parpred`, 7

`grnn.partial`, 6, 8

`grnn.pfi`, 9

`grnn.predict`, 8, 10

`grnn.predone`, 10, 10

`grnn.search_auc`, 7, 11

`grnn.search_rsq`, 12

`grnn.x_imp`, 5, 13, 14

`grnn.x_pfi`, 9, 13, 14