

Package: LLIC (via r-universe)

November 29, 2024

Title Likelihood Criterion (LIC) Analysis for Laplace Regression Model

Version 3.0.0

Date 2024-11-23

Description Performs likelihood criterion analysis using the Laplace regression model to determine its optimal subset of variables. The methodology is based on Guo et al. (2023), LIC criterion for optimal subset selection in distributed interval estimation <doi:10.1080/02331888.2020.1823979>.

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Encoding UTF-8

RoxygenNote 7.3.2

Imports stats, VGAM, dplyr, LaplacesDemon, relliptical, ggplot2, rlang

NeedsCompilation no

Author Guangbao Guo [aut, cre], Yaxuan Wang [aut]

Maintainer Guangbao Guo <ggb11111111@163.com>

Depends R (>= 3.5.0)

Date/Publication 2024-11-28 12:00:02 UTC

Config/pak/sysreqs libxml2-dev

Repository <https://guangbaog.r-universe.dev>

RemoteUrl <https://github.com/cran/LLIC>

RemoteRef HEAD

RemoteSha 7d6409470aaecc10c8d3faf1b24655cac65543d7

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`data_pc`*Data Processing for LLIC Analysis*

Description

This function processes the data generated for the LLIC analysis, including filtering, mutation, and selection of specific columns.

Usage

```
data_pc(data)
```

Arguments

`data` A data frame containing the raw data generated for the LLIC analysis.

Value

A data frame with the following columns:

<code>X1</code>	The filtered values of the original 'X1' column, keeping only rows where 'X1' ≤ 2 .
<code>X2</code>	The original 'X2' column.
<code>X1_squared</code>	A new column containing the square of the 'X1' values.

Examples

```
set.seed(12)
library(dplyr)
library(VGAM)
raw_data <- data.frame(
  X1 = sample(1:3, 1200, replace = TRUE),
  X2 = sample(1:3, 1200, replace = TRUE),
  X3 = sample(1:3, 1200, replace = TRUE),
  X4 = sample(1:3, 1200, replace = TRUE),
  X5 = sample(1:3, 1200, replace = TRUE),
  Y = rlaplace(1200, 0, 1)
)
processed_data <- data_pc(raw_data)
```

 LLIC

LLIC for Lre Model

Description

This function carries out an Laplace LIC analysis utilizing the Lre model.

Usage

```
LLIC(X, y, alpha, K)
```

Arguments

X	Design matrix
y	Random response vector of observed values
alpha	Significance level
K	Number of subsets

Value

A list containing the following components:

MUopt	A vector of the means of the predictor variables in the optimal subset.
Bopt	A vector of the estimated regression coefficients from the final model fitted to the optimal subset.
MAEMUopt	The Mean Absolute Error (MAE) for the optimal subset.
MSEMUopt	The Mean Squared Error (MSE) for the optimal subset.
opt	Currently NULL, a placeholder for potential future use.
Yopt	A vector of the predicted values from the final model fitted to the optimal subset.

Examples

```
set.seed(12)
library(VGAM)
X <- matrix(data = sample(1:3, 1200 * 5, replace = TRUE), nrow = 1200, ncol = 5)
b <- sample(1:3, 5, replace = TRUE)
e <- rlaplace(1200, 0, 1)
Y <- X %*% b + e
alpha <- 0.05
K <- 10
result <- LLIC(X, Y, alpha, K)
MUopt <- result$MUopt
Bopt <- result$Bopt
MAEMUopt <- result$MAEMUopt
MSEMUopt <- result$MSEMUopt
opt <- result$opt
Yopt <- result$Yopt
```

`plot_LLIC`*advanced_plotting_LLIC for LLIC*

Description

This function visualizes the results of the LLIC analysis, including a comparison of actual and predicted values, and a bar chart of model coefficients.

Usage

```
plot_LLIC(X, Y, result)
```

Arguments

<code>X</code>	Design matrix used in the LLIC analysis.
<code>Y</code>	Random response vector of observed values used in the LLIC analysis.
<code>result</code>	A list containing the results of the Laplace LIC analysis from the LLIC function.

Value

A list containing two ‘ggplot’ objects:

`Actual_vs_Pred` A scatter plot comparing the actual vs predicted values.

`Coef` A bar chart displaying the model’s coefficients.

Examples

```
set.seed(12)
library(VGAM)
library(rlang)
library(dplyr)
library(ggplot2)
X <- matrix(data = sample(1:3, 1200 * 5, replace = TRUE), nrow = 1200, ncol = 5)
b <- sample(1:3, 5, replace = TRUE)
e <- rlaplace(1200, 0, 1)
Y <- X %*% b + e
alpha <- 0.05
K <- 10
result <- LLIC(X, Y, alpha, K)
plot_LLIC(X, Y, result)
plots <- plot_LLIC(X, Y, result)
print(plots$Actual_vs_Pred)
print(plots$Coef)
```

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