

Package ‘StepReg’

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Title Stepwise Regression Analysis

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Description Stepwise regression is a statistical technique used for model selection. This package streamlines stepwise regression analysis by supporting multiple regression types, incorporating popular selection strategies, and offering essential metrics. It enables users to apply multiple selection strategies and metrics in a single function call, visualize variable selection processes, and export results in various formats. However, StepReg should not be used for statistical inference unless the variable selection process is explicitly accounted for, as it can compromise the validity of the results. This limitation does not apply when StepReg is used for prediction purposes. We validated StepReg's accuracy using public datasets within the SAS software environment. Additionally, StepReg features an interactive Shiny application to enhance usability and accessibility.

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BugReports <https://github.com/JunhuiLi1017/StepReg/issues>

VignetteBuilder knitr

Suggests knitr, testthat, BiocStyle, kableExtra

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Repository CRAN

Author Junhui Li [cre] (ORCID: <<https://orcid.org/0000-0003-3973-1700>>),

Junhui Li [aut],

Kai Hu [aut],

Xiaohuan Lu [aut],

Kun Cheng [ctb],

Sushmita N Nayak [ctb],

Cesar Bautista Sotelo [ctb],

Michael A Lodato [ctb],

Wenxin Liu [aut],
Lihua Julie Zhu [aut]
Maintainer Junhui Li <junhui.li11@umassmed.edu>
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creditCard	<i>creditCard</i>
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Description

Cross-section data on the credit history for a sample of applicants for a type of credit card. This dataset is from [CreditCard](#)

Usage

```
data(creditCard)
```

Format

A data frame containing 1,319 observations on 12 variables.

Details

- card Factor. Was the application for a credit card accepted?
- reports Number of major derogatory reports.
- age Age in years plus twelfths of a year.
- income Yearly income (in USD 10,000).
- share Ratio of monthly credit card expenditure to yearly income.
- expenditure Average monthly credit card expenditure.
- owner Factor. Does the individual own their home?
- selfemp Factor. Is the individual self-employed?

- dependents Number of dependents.
- months Months living at current address.
- majorcards Number of major credit cards held.
- active Number of active credit accounts.

For more information, refer to [CreditCard](#)

References

Greene, W.H. (2003). *Econometric Analysis*, 5th edition. Upper Saddle River, NJ: Prentice Hall.

plot.StepReg	<i>Plots from a StepReg object</i>
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Description

plot.StepReg visualizes the variable selection procedure using a StepReg object

Usage

```
## S3 method for class 'StepReg'
plot(
  x,
  strategy = attr(x, "nonhidden"),
  process = c("overview", "details"),
  num_digits = 6,
  ...
)
```

Arguments

x	StepReg object
strategy	Select which strategy to be displayed, default is the first name of StepReg object.
process	Select which process of stepwise regression to be displayed from 'details' and 'overview', default is 'overview'.
num_digits	The number of digits to keep when rounding the results. Default is 6.
...	Not used

Value

A list of plots comprising the selection detail plot and selection summary plot for each strategy.

Examples

```
## Not run:
data(mtcars)
formula <- mpg ~ .
x <- stepwise(formula = formula,
              data = mtcars,
              type = "linear",
              strategy = c("forward","bidirection","subset"),
              metric = c("AIC","BIC","SL"))

plot(x)
plot(x, strategy = "forward", process = "overview")

## End(Not run)
```

print.StepReg	<i>Prints from a StepReg object</i>
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Description

print.StepReg prints to console the from an object of class StepReg

Usage

```
## S3 method for class 'StepReg'
print(x, ...)
```

Arguments

x	StepReg object
...	further parameters

Value

only print some dataframe

remission	<i>remission</i>
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Description

A dataset containing the remission and 6 risk factors thought to be related to leukemia remission.

Usage

```
data(remission)
```

Format

A data frame with 27 rows and 7 columns.

Details

- remiss Indicates whether cancer remission occurred. A value of 1 indicates occurrence, while 0 indicates non-occurrence.
- cell Cellularity of the marrow clot section
- smear Smear differential percentage of blasts
- infil Percentage of absolute marrow leukemia cell infiltrate
- li Percentage labeling index of the bone marrow leukemia cells
- blast The absolute number of blasts in the peripheral blood
- temp The highest temperature before the start of treatment

References

Lee, E. T. (1974). "A Computer Program for Linear Logistic Regression Analysis." Computer Programs in Biomedicine 4:80–92.

<https://online.stat.psu.edu/stat501/book/export/html/1011>

report	<i>report from a StepReg object</i>
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Description

report output all tables in StepReg object to a report with format of html, docx, pptx, rtf, and xlsx.

Usage

```
report(x, report_name, format = c("html", "docx", "rtf", "pptx"))
```

Arguments

x	StepReg object
report_name	report name
format	the format of report, choose one or more from 'html', 'docx', 'rtf', 'pptx'. default is 'html'

Examples

```
## Not run:
data(mtcars)
mtcars$yes <- mtcars$wt
formula <- mpg ~ . + 0
x <- stepwise(formula = formula,
              data = mtcars,
              type = "linear",
              strategy = c("bidirection", "subset"),
              metric = c("AIC", "BIC"))
report(x, report_name = "report", format = c("html", "docx"))

## End(Not run)
```

StepRegShinyApp

StepReg Shiny App

Description

StepRegShinyApp is a Shiny application designed for performing stepwise regression analysis. In Step 1, users can upload their dataset, configure settings such as header, separator, and quotes, and select variables for distribution plots. In Step 2, users can choose the regression type (linear, logit, cox, poisson, gamma, or negbin), select dependent and independent variables, specify stepwise strategy (forward, backward, bidirectional, or subset), and set various metrics for model selection. The app dynamically adjusts input options based on the chosen regression type. Additionally, users can specify significant levels for entry and stay in the stepwise process. Finally, they can run the analysis to obtain stepwise regression results and visualize them through summary outputs and plots.

Usage

```
StepRegShinyApp()
```

stepwise

Main wrapper function for stepwise regression

Description

Select optimal model using various stepwise regression strategies, e.g., Forward Selection, Backward Elimination, Bidirectional Elimination; meanwhile, it also supports Best Subset method. Four types of models are currently implemented: linear regression, logistic regression, Cox regression, Poisson, and Gamma regression. For selection criteria, a.k.a, stop rule, users can choose from AIC, AICc, BIC, HQ, Significant Level, and more.

Usage

```

stepwise(
  formula,
  data,
  type = c("linear", "logit", "cox", "poisson", "gamma", "negbin"),
  strategy = c("forward", "backward", "bidirection", "subset"),
  metric = c("AIC", "AICc", "BIC", "CP", "HQ", "adjRsqr", "SL", "SBC", "IC(3/2)", "IC(1)"),
  sle = 0.15,
  sls = 0.15,
  include = NULL,
  test_method_linear = c("Pillai", "Wilks", "Hotelling-Lawley", "Roy"),
  test_method_glm = c("Rao", "LRT"),
  test_method_cox = c("efron", "breslow", "exact"),
  tolerance = 1e-07,
  weight = NULL,
  best_n = 3,
  num_digits = 6
)

```

Arguments

formula	(formula) The formula used for model fitting by defining the scope of dependent and independent variables. The formula takes the form of a '~' (tilde) symbol, with the response variable(s) on the left-hand side, and the predictor variable(s) on the right-hand side. The 'lm()' function uses this formula to fit a regression model. A formula can be as simple as 'y ~ x'. For multiple predictors, they must be separated by the '+' (plus) symbol, e.g. 'y ~ x1 + x2'. To include an interaction term between variables, use the ':' (colon) symbol: 'y ~ x1 + x1:x2'. Use the '.' (dot) symbol to indicate that all other variables in the dataset should be included as predictors, e.g. 'y ~ .'. In the case of multiple response variables (multivariate), the formula can be specified as 'cbind(y1, y2) ~ x1 + x2'. By default, an intercept term is always included in the models, to exclude it, include '0' or '- 1' in your formula: 'y ~ 0 + x1', 'y ~ x1 + 0', and 'y ~ x1 - 1'.
data	(data.frame) A dataset consisting of predictor variable(s) and response variable(s).
type	(character) The stepwise regression type. Choose from 'linear', 'logit', 'poisson', 'cox', 'gamma' and 'negbin'. Default is 'linear'. More information, see StepReg_vignettes
strategy	(character) The model selection strategy. Choose from 'forward', 'backward', 'bidirectional' and 'subset'. Default is 'forward'. More information, see StepReg_vignettes
metric	(character) The model selection criterion (model fit score). Used for the evaluation of the predictive performance of an intermediate model. Choose from 'AIC', 'AICc', 'BIC', 'CP', 'HQ', 'adjRsqr', 'SL', 'SBC', 'IC(3/2)', 'IC(1)'. Default is 'AIC'. More information, see StepReg_vignettes
sle	(numeric) Significance Level to Enter. It is the statistical significance level that a predictor variable must meet to be included in the model. E.g. if 'sle = 0.05',

	a predictor with a P-value less than 0.05 will 'enter' the model. Default is 0.15.
<code>sls</code>	(numeric) Significance Level to Stay. Similar to 'sle', 'sls' is the statistical significance level that a predictor variable must meet to 'stay' in the model. E.g. if 'sls = 0.1', a predictor that was previously included in the model but whose P-value is now greater than 0.1 will be removed.
<code>include</code>	(NULL character) A character vector specifying predictor variables that will always stay in the model. A subset of the predictors in the dataset.
<code>test_method_linear</code>	(character) Test method for multivariate linear regression analysis, choose from 'Pillai', 'Wilks', 'Hotelling-Lawley', 'Roy'. Default is 'Pillai'. For univariate regression, 'F-test' will be used.
<code>test_method_glm</code>	(character) Test method for logit, Poisson, Gamma, and negative binomial regression analysis, choose from 'Rao', 'LRT'. Default is 'Rao'. Only "Rao" is available for strategy = 'subset'.
<code>test_method_cox</code>	(character) Test method for cox regression analysis, choose from 'efron', 'breslow', 'exact'. Default is 'efron'.
<code>tolerance</code>	(numeric) A statistical measure used to assess multicollinearity in a multiple regression model. It is calculated as the proportion of the variance in a predictor variable that is not accounted for by the other predictor variables in the model. Default is 1e-07.
<code>weight</code>	(numeric) A numeric vector specifying the coefficients assigned to the predictor variables. The magnitude of the weight reflects the degree to which each predictor variable contributes to the prediction of the response variable. The range of weight should be from 0 to 1. Values greater than 1 will be coerced to 1, and values less than 0 will be coerced to 0. Default is NULL, which means that all weight are set equal.
<code>best_n</code>	(numeric(integer)) The number of models to be retained in the process output. Default is 3, indicating that only the top 3 best models with the same number of variables are displayed. If all models are displayed, set it to Inf.
<code>num_digits</code>	(numeric(integer)) The number of digits to keep when rounding the results. Default is 6.

Value

A list of multiple models, each with its associated strategy and metric, will be returned.

Author(s)

Junhui Li, Kai Hu, Xiaohuan Lu

References

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- Schwarz, G. (1978). Estimating the dimension of a model. *Annals of Statistics*, 6(2), pages. 15-18.

Examples

```
## perform multivariate linear stepwise regression with 'bidirection'
## strategy and 'AIC' stop rule, excluding intercept.
data(mtcars)
mtcars$yes <- mtcars$wt
formula <- cbind(mpg,drat) ~ . + 0
stepwise(formula = formula,
          data = mtcars,
          type = "linear",
          strategy = "bidirection",
          metric = "AIC")
## perform linear stepwise regression with 'bidirection' strategy and
## "AIC", "SBC", "SL", "AICc", "BIC", and "HQ" stop rule.
formula <- mpg ~ . + 1
stepwise(formula = formula,
```

```

data = mtcars,
type = "linear",
strategy = c("forward", "bidirection"),
metric = c("AIC", "SBC", "SL", "AICc", "BIC", "HQ"))

## perform logit stepwise regression with 'forward' strategy and significance
## level as stop rule.
data(remission)
formula <- remiss ~ .
stepwise(formula = formula,
         data = remission,
         type = "logit",
         strategy = "forward",
         metric = "SL",
         sle=0.05,
         sls=0.05)

```

tobacco

tobacco

Description

data on chemical components of 25 tobacco leaf

Usage

```
data(tobacco)
```

Format

A data frame containing 25 observations on 9 variables.

Details

- cigarette Rate of cigarette burn in inches per 1000 seconds.
- sugar Percent sugar in the leaf.
- nicotine Percent nicotine.
- nitrogen Percentage of nitrogen.
- chlorine Percentage of chlorine.
- potassium Percentage of potassium.
- phosphorus Percentage of phosphorus
- calcium Factor. Percentage of calcium.
- magnesium Percentage of magnesium.

References

Anderson, R. L. and Bancroft, T. A. (1952), Statistical Theory in Research, McGraw-Hill Book Company, Inc., New York, NY.

vote	<i>Vote for all models</i>
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Description

Votes for all models across all combinations of strategies and metrics

Usage

```
vote(x, ...)
```

Arguments

x	each dataframe from outputlist
...	further parameters

Value

A dataframe with column names "model" and combinations of strategy and metric. The first column represents the model formula, and a checkmark indicates that the corresponding model was supported by the given strategy and metric combination. Please note that for the subset strategy, the "vote" will report the single best model across all numbers of variables under Information Criteria (IC). However, this rule should not be applied to Significance Level (SL) because the F/Rao value is only comparable for models with the same number of variables.

Examples

```
data(mtcars)
formula <- mpg ~ .
x <- stepwise(formula = formula,
              data = mtcars,
              type = "linear",
              strategy = c("forward", "backward", "subset"),
              metric = c("AIC", "BIC"))
vote(x)
```

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