Package 'smoothic'

August 22, 2023

Type Package

Title Variable Selection Using a Smooth Information Criterion

Version 1.2.0

Depends R (>= 3.5.0)

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Description Implementation of the SIC epsilon-telescope method, either using single or distributional (multiparameter) regression. Includes classical regression with normally distributed errors and robust regression, where the errors are from the Laplace distribution. The ``smooth generalized normal distribution" is used, where the estimation of an additional shape parameter allows the user to move smoothly between both types of regression. See O'Neill and Burke (2022)
``Robust Distributional Regression with Automatic Variable Selection" for more details. <arXiv:2212.07317>. This package also contains the data analyses from O'Neill and Burke (2023). ``Variable selection using a smooth information criterion for distributional regression models". <doi:10.1007/s11222-023-10204-8>.

License GPL-3

URL https://github.com/meadhbh-oneill/smoothic,

https://meadhbh-oneill.ie/smoothic/

Encoding UTF-8

LazyData true

RoxygenNote 7.2.3

Imports data.table, dplyr, ggplot2, MASS, numDeriv, purrr, rlang, stringr, tibble, tidyr, toOrdinal

Suggests knitr, rmarkdown

VignetteBuilder knitr

NeedsCompilation no

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Repository CRAN

Date/Publication 2023-08-22 14:40:02 UTC

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bostonhouseprice Boston House Price Data (Original)

Description

Original data, which come from a study by Harrison Jr and Rubinfeld (1978), examining the association between median house prices in a particular community with various community characteristics. See bostonhouseprice2 for the corrected version, with additional variables.

Usage

bostonhouseprice

Format

A data frame with 506 rows and 9 variables:

crime crimes committed per capita

rooms average number of rooms per house

radial index of accessibility to radial highways

stratio average student-teacher ratio of schools in the community

lowstat percentage of the population that are "lower status"

Inox log(annual average nitrogen oxide concentration (pphm))

lproptax log(property tax per \$1000)

ldist log(weighted distances to five employment centres in the Boston region)

lprice log(median house price (\$))

Source

https://CRAN.R-project.org/package=wooldridge

bostonhouseprice2

References

Harrison Jr, D. and Rubinfeld, D. L. (1978). Hedonic housing prices and the demand for clean air. Journal of environmental economics and management, 5(1):81-102.

Wooldridge, J. M. (2015). Introductory econometrics: A modern approach. Cengage learning.

bostonhouseprice2 Boston House Price Data (Corrected Version)

Description

Corrected data, which come from a study by Harrison Jr and Rubinfeld (1978), examining the association between median house prices in a particular community with various community characteristics. See bostonhouseprice for the original version.

Usage

bostonhouseprice2

Format

A data frame with 506 rows and 13 variables:

crim per capita crime rate by town
zn proportion of residential land zoned for lots over 25,000 sq.ft
indus proportion of non-retail business acres per town
rm average number of rooms per dwelling
age proportion of owner-occupied units built prior to 1940
rad index of accessibility to radial highways
ptratio pupil-teacher ratio by town
lnox log(nitric oxides concentration (parts per 10 million))
ldis log(weighted distances to five Boston employment centres)
ltax log(full-value property-tax rate per USD 10,000)
llstat log(percentage of lower status of the population)
chast Charles River dummy variable (=1 if tract bounds river; 0 otherwise)
lcmedv log(corrected median value of owner-occupied homes in USD 1000's)

Source

https://CRAN.R-project.org/package=mlbench

References

Harrison Jr, D. and Rubinfeld, D. L. (1978). Hedonic housing prices and the demand for clean air. Journal of environmental economics and management, 5(1):81-102.

Leisch F, Dimitriadou E (2021). mlbench: Machine Learning Benchmark Problems. R package version 2.1-3.

citycrime

Description

Data relating to crime rates per one million residents in 50 U.S cities, taken from Thomas (1990).

Usage

citycrime

Format

A data frame with 50 rows and 7 variables:

violent reported violent crime rate per 100,000 residents

funding annual police funding per resident (\$)

hs percentage of people 25 years+ with 4 years of high school

not_hs percentage of 16 to 19 year-olds not in high school and not high school graduates

college percentage of 18 to 24 year-olds in college

college4 percentage of people 25 years+ with at least 4 years of college

crime_rate total overall reported crime rate per 1 million residents

Source

https://hastie.su.domains/StatLearnSparsity_files/DATA/crime.txt

References

Thomas, G.S., 1990. The Rating Guide to Life in America's Small Cities. Prometheus Books, 59 John Glenn Drive, Amherst, NY 14228-2197.

Hastie, T., Tibshirani, R. and Wainwright, M., 2015. Statistical learning with sparsity: the lasso and generalizations. CRC press.

diabetes

Description

Data relating to a study of disease progression one year after baseline.

Usage

diabetes

Format

A data frame with 442 rows and 11 variables:

- AGE age of the patient
- SEX sex of the patient
- BMI body mass index of the patient
- **BP** blood pressure of the patient
- S1 blood serum measurement 1
- S2 blood serum measurement 2
- S3 blood serum measurement 3
- S4 blood serum measurement 4
- **S5** blood serum measurement 5
- S6 blood serum measurement 6

Y quantitative measure of disease progression one year after baseline

Source

https://CRAN.R-project.org/package=lars

References

Efron, B., Hastie, T., Johnstone, I., Tibshirani, R., et al. (2004). Least angle regression. The Annals of Statistics.

pcancer

Description

Data, which come from a study by Stamey et al. (1989), examining the correlation between the level of prostate-specific antigen (PSA) and various clinical measures in men who were about to receive a radical prostatectomy.

Usage

pcancer

Format

A data frame with 97 rows and 9 variables:

lcavol log(cancer volume (cm^3))

lweight log(prostate weight (g))

age age of the patient

lbph log(amount of benign prostatic hyperplasia (cm^2))

svi presence of seminal vesicle invasion (1=yes, 0=no)

lcp log(capsular penetration (cm))

gleason Gleason score

pgg45 percentage of Gleason scores four of five

lpsa log(PSA (ng/mL))

Source

https://web.stanford.edu/~hastie/ElemStatLearn/datasets/prostate.data

References

Stamey, T. A., Kabalin, J. N., McNeal, J. E., Johnstone, I. M., Freiha, F., Redwine, E. A., and Yang, N. (1989). Prostate specific antigen in the diagnosis and treatment of adenocarcinoma of the prostate. ii. radical prostatectomy treated patients. The Journal of urology, 141(5):1076-1083.

plot_effects

Description

This function plots the model-based conditional density curves for different effect combinations. For example, take a particular covariate that is selected in the final model. The other selected covariates are fixed at their median values by default (see covariate_fix to fix at other values) and then the plotted red and blue densities correspond to the modification of the chosen covariate as "low" (25th quantile by default) and "high" (75th quantile by default).

Usage

```
plot_effects(
    obj,
    what = "all",
    show_average_indiv = TRUE,
    p = c(0.25, 0.75),
    covariate_fix,
    density_range
)
```

Arguments

obj	An object of class "smoothic" which is the result of a call to smoothic.
what	The covariate effects to be plotted, default is what = "all". The user may supply a vector of covariate names to be plotted (only covariates selected in the final model can be plotted).
show_average_in	div
	Should a "baseline" or "average" individual be shown, default is show_average_indiv = TRUE. If show_average_indiv = FALSE then this is not shown.
	The probabilities given to the quantile function. This corresponds to the plotted red and blue density curves where the chosen covariate is modified as "low" and "high". The default is $p = c(0.25, 0.75)$ to show the 25th and 75th quantiles.
	Optional values to fix the covariates at that are chosen in the final model. When not supplied, the covariates are fixed at their median values. See the example for more detail.
density_range	Optional range for which the density curves should be plotted.

Value

A plot of the conditional density curves.

Author(s)

Meadhbh O'Neill

Examples

```
# Sniffer Data -----
# MPR Model ----
results <- smoothic(</pre>
 formula = y \sim .,
 data = sniffer,
 family = "normal",
 model = "mpr"
)
plot_effects(results)
# Only plot gastemp and gaspres
# Do not show the average individual plot
# Plot the lower and upper density curves using 10th quantile (lower) and 90th quantile (upper)
# Fix violent to its violent to 820 and funding to 40
plot_effects(results,
             what = c("gastemp", "gaspres"),
             show_average_indiv = FALSE,
             p = c(0.1, 0.9),
             covariate_fix = c("gastemp" = 70,
                               "gaspres" = 4))
# The curves for the gastemp variable are computed by fixing gaspres = 4 (as is specified
# in the input). The remaining variables that are not specified in covariate_fix are fixed
```

```
# to their median values (i.e., tanktemp is fixed at its median). gastemp is then modified
# to be low (10th quantile) and high (90th quantile), as specified by p in the function.
```

plot_paths

Plot the ϵ *-telescope coefficient paths*

Description

This function plots the standardized coefficient values with respect to the ϵ -telescope for the location (and dispersion) components.

Usage

```
plot_paths(
    obj,
    log_scale_x = TRUE,
    log_scale_x_pretty = TRUE,
    facet_scales = "fixed"
)
```

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predict.smoothic

Arguments

obj	An object of class "smoothic" which is the result of a call to smoothic.
log_scale_x	Default is log_scale_x = TRUE, which uses a log scale on the x-axis. If log_scale_x = FALSE, then the raw values of the ϵ -telescope are plotted.
log_scale_x_pr	retty
	Default is log_scale_x_pretty = TRUE, where the x-axis labels are "pretty". epsilon_1 and epsilon_T must be a number to the power of 10 for this to apply.
facet_scales	Default is facet_scales = "fixed". This is supplied to facet_wrap.

Value

A plot of the standardized coefficient values through the ϵ -telescope.

Author(s)

Meadhbh O'Neill

Examples

```
# Sniffer Data -----
# MPR Model ----
results <- smoothic(
  formula = y ~ .,
  data = sniffer,
  family = "normal",
  model = "mpr"
)
plot_paths(results)</pre>
```

predict.smoothic *Predict smoothic*

Description

predict method class "smoothic"

Usage

```
## S3 method for class 'smoothic'
predict(object, newdata, ...)
```

Arguments

object	an object of class "smoothic" which is the result of a call to smoothic.
newdata	new data object
	further arguments passed to or from other methods.

smoothic

Value

a matrix containing the predicted values for the location mu and scale s

Author(s)

Meadhbh O'Neill

Examples

```
# Sniffer Data -----
# MPR Model ----
results <- smoothic(
  formula = y ~ .,
  data = sniffer,
  family = "normal",
  model = "mpr"
)
predict(results)</pre>
```

smoothic

Variable Selection Using a Smooth Information Criterion (SIC)

Description

Implements the SIC ϵ -telescope method, either using single or multiparameter regression. Returns estimated coefficients, estimated standard errors and the value of the penalized likelihood function. Note that the function will scale the predictors to have unit variance, however, the final estimates are converted back to their original scale.

Usage

```
smoothic(
  formula,
  data,
  family = "sgnd",
 model = "mpr",
  lambda = "log(n)",
  epsilon_1 = 10,
  epsilon_T = 1e-04,
  steps_T = 100,
  zero_tol = 1e-05,
 max_{it} = 10000,
 kappa,
  tau,
 max_it_vec,
  stepmax_nlm
)
```

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smoothic

Arguments

-	
formula	An object of class "formula": a two-sided object with response on the left hand side and the model variables on the right hand side.
data	A data frame containing the variables in the model; the data frame should be unstandardized.
family	The family of the model, default is family = "sgnd" for the "Smooth General- ized Normal Distribution" where the shape parameter kappa is also estimated. Classical regression with normally distributed errors is performed when family = "normal". If family = "laplace", this corresponds to a robust regression with errors from a Laplace-like distribution. If family = "laplace", then the default value of tau = 0.15, which is used to approximate the absolute value in the Laplace density function.
model	The type of regression to be implemented, either model = "mpr" for multiparam- eter regression (i.e., location and scale), or model = "spr" for single parameter regression (i.e., location only). Defaults to model="mpr".
lambda	Value of penalty tuning parameter. Suggested values are " $log(n)$ " and "2" for the BIC and AIC respectively. Defaults to lambda =" $log(n)$ " for the BIC case. This is evaluated as an R expression, so it may be a number of some function of n.
epsilon_1	Starting value for ϵ -telescope. Defaults to 10.
epsilon_T	Final value for ϵ -telescope. Defaults to 1e-04.
steps_T	Number of steps in ϵ -telescope. Defaults to 100, must be greater than or equal to 10.
zero_tol	Coefficients below this value are treated as being zero. Defaults to 1e-05.
max_it	Maximum number of iterations to be performed before the optimization is ter- minated. Defaults to 1e+04.
kappa	Optional user-supplied positive kappa value (> 0.2 to avoid computational is- sues) if family = "sgnd". If supplied, the shape parameter kappa will be fixed to this value in the optimization. If not supplied, kappa is estimated from the data.
tau	Optional user-supplied positive smoothing parameter value in the "Smooth Gen- eralized Normal Distribution" if family = "sgnd" or family = "laplace". If not supplied then tau = 0.15. If family = "normal" then tau = 0 is used. Smaller values of tau bring the approximation closer to the absolute value function, but this can cause the optimization to become unstable. Some issues with standard error calculation with smaller values of tau when using the Laplace distribution in the robust regression setting.
<pre>max_it_vec</pre>	Optional vector of length steps_T that contains the maximum number of iter- ations to be performed in each ϵ -telescope step. If not supplied, max_it is the maximum number of iterations performed for 10 steps and then the maximum number of iterations to be performed reduces to 10 for the remainder of the telescope.
stepmax_nlm	Optional maximum allowable scaled step length (positive scalar) to be passed to nlm. If not supplied, default values in nlm are used.

A list with estimates and estimated standard errors.

- coefficients vector of coefficients.
- see vector of estimated standard errors.
- model the matched type of model which is called.
- plike value of the penalized likelihood function.
- kappa value of the estimated/fixed shape parameter kappa if family = "sgnd".

Author(s)

Meadhbh O'Neill

References

O'Neill, M. and Burke, K. (2023) Variable selection using a smooth information criterion for distributional regression models. <doi:10.1007/s11222-023-10204-8>

O'Neill, M. and Burke, K. (2022) Robust Distributional Regression with Automatic Variable Selection. <arXiv:2212.07317>

Examples

```
# Sniffer Data -----
# MPR Model ----
results <- smoothic(
  formula = y ~ .,
  data = sniffer,
  family = "normal",
  model = "mpr"
)
summary(results)</pre>
```

sniffer

Sniffer Data

Description

Data examining the factors that impact the amount of hydrocarbon vapour released when gasoline is pumped into a tank.

Usage

sniffer

Format

A data frame with 125 rows and 5 variables:

tanktemp initial tank temperature (degrees F)
gastemp temperature of the dispensed gasoline (degrees F)
tankpres initial vapour pressure in the tank (psi)
gaspres vapour pressure of the dispensed gasoline (psi)
y hydrocarbons emitted (g)

Source

https://CRAN.R-project.org/package=alr4

References

Bedrick, E.J. (2000). Checking for lack of fit in linear models with parametric variance functions. Technometrics 42 (3), 226–236.

Weisberg, S. (2014). Applied Linear Regression, 4th edition. Hoboken NJ: Wiley.

summary.smoothic Summarising Smooth Information Criterion (SIC) Fits

Description

summary method class "smoothic"

Usage

```
## S3 method for class 'smoothic'
summary(object, ...)
```

Arguments

object	an object of class "smoothic" which is the result of a call to smoothic.
	further arguments passed to or from other methods.

Value

A list containing the following components:

- model the matched model from the smoothic object.
- coefmat a typical coefficient matrix whose columns are the estimated regression coefficients, estimated standard errors (SEE) and p-values.
- plike value of the penalized likelihood function.

Author(s)

Meadhbh O'Neill

Examples

```
# Sniffer Data -----
# MPR Model ----
results <- smoothic(
  formula = y ~ .,
  data = sniffer,
  family = "normal",
  model = "mpr"
)
summary(results)</pre>
```

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