## Package 'RAZIAD'

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Title Regression Analysis of Zero-Inflated and Zero-Altered(Hurdle)

Data Version 0.0.1

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**Description** It performs regression analysis for zero-inflated and zero-altered data, enabling modeling of sparse response variable distributions and assessing their association with covariates. It computes Maximum Likelihood Estimates (MLE) and conducts model selection using AIC and BIC criteria. Additionally, it calculates Fisher Information, confidence intervals, standard errors, and z-scores for all model parameters.

**Encoding** UTF-8

RoxygenNote 7.3.1

Imports Matrix, extraDistr, stats

**Depends** R (>= 4.3.1.0)

LazyData true

Suggests testthat (>= 3.0.0)

Config/testthat/edition 3

**License** MIT + file LICENSE

NeedsCompilation no

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**Repository** CRAN

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DebTrivedi

#### Description

The DebTrivedi dataset contains data on health care utilization and insurance coverage.

#### Usage

data("DebTrivedi")

#### Format

A data frame with 4406 observations on the following 19 variables.

- ofp a numeric vector indicating physicians office visits
- ofnp a numeric vector
- opp a numeric vector
- opnp a numeric vector
- emer a numeric vector
- hosp a numeric vector indicating number of hospital stays
- health a factor with levels poor average excellent indicating self-perceived health status
- numchron a numeric vector indicating number of chronic conditions
- adldiff a factor with levels no yes
- region a factor with levels midwest noreast other west
- age a numeric vector
- black a factor with levels no yes
- gender a factor with levels female male
- married a factor with levels no yes
- school a numeric vector indicating number of years of education
- faminc a numeric vector
- employed a factor with levels no yes
- privins a factor with levels no yes indicating private insurance indicator
- medicaid a factor with levels no yes

#### fisher.reg

#### Details

Deb and Trivedi (1997) conducted an analysis on data comprising 4406 individuals aged 66 and above, enrolled in Medicare, a public insurance program. These data were initially sourced from the US National Medical Expenditure Survey (NMES) for the years 1987/88. They are accessible through the data archive of the Journal of Applied Econometrics at http://qed.econ.queensu.ca/jae/1997-v12.3/deb-trivedi/. Additionally, the dataset was compiled for an R package associated with Kleiber and Zeileis (2008) and is also retrievable as DebTrivedi.rda from the Journal of Statistical Software, alongside Zeileis (2006). The primary aim is to develop models for medical care demand, encompassing physician/non-physician office and hospital outpatient visits, based on available patient covariates.

#### Source

http://www.jstatsoft.org/v27/i08/paper

#### References

Zeileis, A. and Kleiber, C. and Jackma, S. (2008). "Regression Models for Count Data in R". JSS 27, 8, 1–25.

#### Examples

```
data(DebTrivedi)
plot(table(DebTrivedi$ofp), main = "physicians office visit (N=4406)", ylab="frequency")
#the response variable "physicians office visits" is highly right skewed
```

fisher.reg	Fisher Information computation for calculating the confidence inter-
	vals in Zero-inflated and Zero-altered regression models

#### Description

Fisher Information computation for calculating the confidence intervals in Zero-inflated and Zeroaltered regression models

#### Usage

```
fisher.reg(x, y, b0 = NULL, m, dist = "ZIP", link = "logit")
```

#### Arguments

x	a design matrix containing an intercept column (all ones) along with other avail- able covariates for the response variable
У	a zero-inflated or zero-altered(Hurdle) count response variable, represented as an integer vector
b0	the initial parameters for the model, calculated as the product of the number of parameters in the specified models and the number of covariates

m	M set in trigamma free approach only needed for ZIBNB, BNBH, ZINB, and NBH
dist	can be specified as follows: "ZIP" for "zero-inflated Poisson", "ZINB" for "zero- inflated negative binomial", "ZINB-r" for "zero-inflated negative binomial with fixed r", "ZIBNB" for "zero-inflated beta negative binomial", "ZIBB" for "zero- inflated beta binomial", "ZIBB-n" for "zero-inflated beta binomial with fixed n", "ZIBB-ab" for "zero-inflated beta binomial with fixed alpha and beta", "PH" for "zero-altered(hurdle) Poisson", "NBH" for "zero-altered(Hurdle) negative binomial", "NBH-r" for "zero-altered (Hurdle) negative binomial with fixed r", "BNBH" for " zero-altered (Hurdle) beta negative binomial", "BBH" for "zero- altered (Hurdle) beta binomial", "BBH-n" for "zero-altered(Hurdle) beta bino- mial with fixed n", and "BBH-ab" for "zero-altered(Hurdle) beta binomial with fixed alpha and beta".
link	can be set to one of four different options: "logit" for the logistic link function, "probit" for the probit link function, "loglog" for the log-log link function, and "cloglog" for the complementary log-log link function

#### Value

If dist = ZIP, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = ZINB, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = ZINB-r, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.

#### fisher.reg

- Estimated parameters.
- Estimation/length ratio.
- Standard error.
- Z-score.

If dist = ZIBNB, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = ZIBB, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = ZIBB-n, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = ZIBB-ab, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.

- Standard error.
- Z-score.

If dist = PH, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = NBH, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = NBH-r, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = BNBH, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

#### fisher.reg

If dist = BBH, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = BBH-n, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

If dist = BBH-ab, the following values are returned:

- FisherInformation: Fisher Information matrix for all the parameters and covariates in the model.
- ConfidenceIntervals: Contains the following information:
  - Lower and upper bounds of the confidence interval.
  - Estimated parameters.
  - Estimation/length ratio.
  - Standard error.
  - Z-score.

#### Examples

```
intercept<- rep(1,4406)
dt = DebTrivedi[, c(6:8, 13, 15, 18)]
dt = cbind(intercept, dt)
dt$gender.male <- ifelse(dt$gender == 'male', 1, 0)
dt$gender.female <- ifelse(dt$gender == 'female', 1, 0)
dt$health.poor <- ifelse(dt$health == 'poor', 1, 0)
dt$health.average <- ifelse(dt$health == 'average', 1, 0)
dt$health.excellent <- ifelse(dt$health == 'excellent', 1, 0)
dt$privins.yes <- ifelse(dt$privins == 'yes', 1, 0)
dt$privins.no <- ifelse(dt$privins == 'no', 1, 0)
y = DebTrivedi[,1]
x = data.matrix(dt[, c(1, 2, 4, 6, 8, 10, 12, 13)])
np = dim(x)[2]
```

reg.model

#### Regression Analysis for Zero-altered or Zero-inflated Data

#### Description

Regression Analysis for Zero-altered or Zero-inflated Data

#### Usage

reg.model(x, y, b0 = NULL, dist = "ZIP", link)

#### Arguments

x	a design matrix containing an intercept column (all ones) along with other available covariates available and selected for the response variable
У	a zero-inflated or zero-altered(Hurdle) count response variable, represented as an integer vector
b0	the initial parameters for the model, calculated as the product of the number of parameters in the specified models and the number of covariates. For simplicity one may put the MLE and intercept of the parameters and set the rest of covariates to zero or change them.
dist	can be specified as follows: "ZIP" for "zero-inflated Poisson", "ZINB" for "zero- inflated negative binomial", "ZINB-r" for "zero-inflated negative binomial with fixed r", "ZIBNB" for "zero-inflated beta negative binomial", "ZIBB" for "zero- inflated beta binomial", "ZIBB-n" for "zero-inflated beta binomial with fixed n", "ZIBB-ab" for "zero-inflated beta binomial with fixed alpha and beta", "PH" for "zero-altered(hurdle) Poisson", "NBH" for "zero-altered(Hurdle) negative binomial", "NBH-r" for "zero-altered (Hurdle) negative binomial with fixed r", "BNBH" for " zero-altered (Hurdle) beta negative binomial", "BBH" for "zero- altered (Hurdle) beta binomial", "BBH-n" for "zero-altered(Hurdle) beta bino- mial with fixed n", and "BBH-ab" for "zero-altered(Hurdle) beta binomial with fixed alpha and beta".
link	can be set to one of four different options: "logit" for the logistic link function, "probit" for the probit link function, "loglog" for the log-log link function, and "cloglog" for the complementary log-log link function

#### Value

A list containing AIC, BIC, the corresponding value of log likelihood, and the maximum likelihood estimate (MLE) of the unknown parameters in the model. If dist = ZIP, the following values are returned:

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- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log.likelihhod: The value of log likelihood with maximum likelihood estimate plugged-in.
- Estimated.Parameters: The maximum likelihood estimate of  $\Gamma$  and  $\beta$  for intercept and covariates included in the design matrix.

If dist = ZINB, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\Gamma)$ ,  $(\beta_1)$ , and  $(\beta_2)$  for intercept and covariates included in the design matrix.

If dist = ZINB-r, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\beta_1)$  for intercept only and  $(\Gamma)$ , and  $(\beta_2)$  for intercept and covariates included in the design matrix.

If dist = ZIBNB, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\Gamma)$ ,  $(\beta_1)$ ,  $(\beta_2)$ , and  $(\beta_3)$  for intercept and covariates included in the design matrix.

If dist = ZIBB, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.

- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\Gamma)$ ,  $(\beta_1)$ ,  $(\beta_2)$ , and  $(\beta_3)$  for intercept and covariates included in the design matrix.

If dist = ZIBB-n, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\beta_1)$  for intercept only and  $(\Gamma)$ ,  $(\beta_2)$ , and  $(\beta_3)$  for intercept and covariates included in the design matrix.

If dist = ZIBB-ab, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\beta_2)$  and  $(\beta_3)$  for intercept only and  $(\Gamma)$  and  $(\beta_1)$  for intercept and covariates included in the design matrix.

If dist = PH, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated.Parameters: The maximum likelihood estimate of Γ and β for intercept and covariates included in the design matrix.

If dist = NBH, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of (Γ), (β<sub>1</sub>), and (β<sub>2</sub>) for intercept and covariates included in the design matrix.

If dist = NBH-r, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\beta_1)$  for intercept only and  $(\Gamma)$ , and  $(\beta_2)$  for intercept and covariates included in the design matrix.

If dist = BNBH, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\Gamma)$ ,  $(\beta_1)$ ,  $(\beta_2)$ , and  $(\beta_3)$  for intercept and covariates included in the design matrix.

If dist = BBH, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\Gamma)$ ,  $(\beta_1)$ ,  $(\beta_2)$ , and  $(\beta_3)$  for intercept and covariates included in the design matrix.

If dist = BBH-n, the following values are returned:

- AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.
- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\beta_1)$  for intercept only and  $(\Gamma)$ ,  $(\beta_2)$ , and  $(\beta_3)$  for intercept and covariates included in the design matrix.

If dist = BBH-ab, the following values are returned:

• AIC: Akaike Information Criterion, a measure of the model's goodness of fit adjusted for the number of parameters.

- BIC: Bayesian Information Criterion, a criterion for model selection among a finite set of models.
- log-likelihood: The value of log-likelihood corresponding to the maximum likelihood estimate.
- Estimated Parameters: Maximum likelihood estimates of  $(\beta_2)$  and  $(\beta_3)$  for intercept only and  $(\Gamma)$  and  $(\beta_1)$ , for intercept and covariates included in the design matrix.

#### Examples

```
intercept<- rep(1,4406)
dt = DebTrivedi[, c(6:8, 13, 15, 18)]
dt = cbind(intercept, dt)
dt$gender.male <- ifelse(dt$gender == 'male', 1, 0)
dt$gender.female <- ifelse(dt$gender == 'female', 1, 0)
dt$health.poor <- ifelse(dt$health == 'poor', 1, 0)
dt$health.average <- ifelse(dt$health == 'average', 1, 0)
dt$health.excellent <- ifelse(dt$health == 'excellent', 1, 0)
dt$privins.yes <- ifelse(dt$privins == 'yes', 1, 0)
dt$privins.no <- ifelse(dt$privins == 'no', 1, 0)
y = DebTrivedi[,1]
x = data.matrix(dt[, c(1, 2, 4, 6, 8, 10, 12, 13)])
np = dim(x)[2]
b0 = c(rep(0.3, np), rep(0.1, np))
reg.model(x, y, b0=b0, dist="PH", link="logit")
```

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