## Package 'BPmodel'

September 23, 2021

$\mathbf{r}$
Type Package
Title Beta-Prime Regression Model
Version 1.1.2
Author Manoel Santos-Neto [aut, cre], Marcelo Bourguignon [aut], Mário de Castro [aut], Tales Ribeiro-Bezerra [aut]
Maintainer Manoel Santos-Neto <manoel.ferreira@professor.ufcg.edu.br></manoel.ferreira@professor.ufcg.edu.br>
<ul><li><b>Description</b> A new regression model for positive random variables with skewed and long tail.</li><li><b>License</b> GPL-3</li></ul>
<pre>URL https://github.com/santosneto/BPmodel</pre>
BugReports https://github.com/santosneto/BPmodel/issues
<b>Depends</b> R (>= 3.1.2)
<b>Imports</b> base, extraDistr, gamlss, gamlss.dist, ggplot2, Deriv, dplyr, graphics, pracma
Suggests testthat
Encoding UTF-8
RoxygenNote 7.1.2
NeedsCompilation no
Repository CRAN
<b>Date/Publication</b> 2021-09-23 09:00:04 UTC
R topics documented:
BP       2         diag.BP       3         envelope       4         test       5
Index 7

Reparameterized Beta Prime (BP) distribution for fitting a GAMLSS

2 BP

Description

of the BP distribution.

ΒP

# The function BP() defines the BP distribution, a two parameter distribution, for a gamlss.family object to be used in GAMLSS fitting using using the function gamlss(), with mean equal to the parameter mu and sigma equal the precision parameter. The functions dBP, pBP, qBP and rBP define the density, distribution function, quantile function and random generation for the BP parameterization

#### Usage

```
BP(mu.link = "log", sigma.link = "log")

dBP(x, mu = 1, sigma = 1, log = FALSE)

pBP(q, mu = 1, sigma = 1, lower.tail = TRUE, log.p = FALSE)

rBP(n = 1, mu = 1, sigma = 1)

qBP(p, mu = 1, sigma = 1, lower.tail = TRUE, log.p = FALSE)
```

#### **Arguments**

mu.link	object for which the extraction of model residuals is meaningful.
sigma.link	type of residual to be used.
x, q	vector of quantiles.
mu	vector of scale parameter values.
sigma	vector of shape parameter values.
log	logical; if TRUE, quantiles are given as log.
lower.tail	logical; if TRUE (default), probabilities are $P[X \le x]$ , otherwise, $P[X > x]$ .
log.p	log.p logical; if TRUE, probabilities p are given as log(p).
n	number of observations. If $length(n) > 1$ , the length is taken to be the number required.
р	vector of probabilities.

#### Value

returns a gamlss.family object which can be used to fit a BP distribution in the gamlss() function.

#### Note

For the function BP(), mu is the mean and sigma is the precision parameter of the BP distribution.

diag.BP

#### Author(s)

Manoel Santos-Neto <manoel.ferreira at professor.ufcg.edu.br>

#### References

Rigby, R.A., Stasinopoulos, D.M., Heller, G.Z., and De Bastiani, F. Distributions for modeling location, scale, and shape: Using GAMLSS in R, London: Chapman and Hall/CRC, 2019.

Stasinopoulos D.M., Rigby R.A., Heller G., Voudouris V., and De Bastiani F. Flexible Regression and Smoothing: Using GAMLSS in R, London: Chapman and Hall/CRC, 2017

Bourguignon, M., Santos-Neto, M. and Castro, M. A new regression model for positive random variables with skewed and long tail. *METRON*, v. 79, p. 33–55, 2021. doi: 10.1007/s40300021-00203y

#### **Examples**

```
y <- rBP(n = 100)
hist(y)
plot(function(x) dBP(x), 0.0001, 8)
gamlss::gamlss(y ~ 1, family = BP)</pre>
```

diag.BP

Diagnostic Analysis - Local Influence

#### **Description**

Diagnostics for the BP, GA, IG and WEI regression models

#### Usage

```
diag.BP(model, mu.link = "log", sigma.link = "log", scheme = "case.weight")

diag.GA(
    model,
    mu.link = "log",
    sigma.link = "identity",
    scheme = "case.weight"
)

diag.IG(
    model,
    mu.link = "log",
    sigma.link = "identity",
    scheme = "case.weight"
)

diag.WEI(
```

4 envelope

```
model,
mu.link = "log",
sigma.link = "identity",
scheme = "case.weight"
)
res_pearson(model)
```

#### Arguments

model Object of class gamlss holding the fitted model.

mu.link Defines the mu.link, with "identity" link as the default for the mu parameter.

sigma.link Defines the sigma.link, with "identity" link as the default for the sigma parame-

ter.

scheme Default is "case.weight". But, can be "response".

#### Value

Local influence measures.

#### Author(s)

Manoel Santos-Neto <manoel.ferreira at professor.ufcg.edu.br>

#### References

Bourguignon, M., Santos-Neto, M. and Castro, M. A new regression model for positive random variables with skewed and long tail. *METRON*, v. 79, p. 33–55, 2021. doi: 10.1007/s40300021-00203y

envelope

Envelopes

#### **Description**

Computes simulation envelopes.

#### Usage

```
envelope(
  model,
  k = 100,
  color = "grey50",
  xlabel = "Theorical Quantile",
  ylabel = "Empirical Quantile",
  font = "serif"
)
```

test 5

#### Arguments

model object of class gamlss.

k number of replications for envelope construction. Default is 19.

color a specification for the default envelope color.

xlabel a label for the x axis. ylabel a label for the y axis.

font the name of a font family for x and y axis.

#### Value

A simulated envelope.

#### Author(s)

Manoel Santos-Neto <manoel.ferreira at professor.ufcg.edu.br>

#### References

Atkinson, A.C. Plots, transformations and regression: an introduction to graphical methods of diagnostic regression analysis. Oxford: Oxford Science Publications, 1985.

Bourguignon, M., Santos-Neto, M. and Castro, M. A new regression model for positive random variables with skewed and long tail. *METRON*, v. 79, p. 33–55, 2021. doi: 10.1007/s40300021-00203y

Leiva, V., Santos-Neto, M., Cysneiros, F.J.A, Barros, M. Birnbaum-Saunders statistical modeling: a new approach. *Statistical Modelling*, v. 14, p. 21–48, 2014. doi: 10.1177/1471082X13494532

Santos-Neto, M., Cysneiros, F.J.A., Leiva, V., Barros, M. Reparameterized Birnbaum-Saunders regression models with varying precision. *Electronic Journal of Statistics*, v. 10, p. 2825–2855, 2016. doi: 10.1214/16EJS1187.

test Precision test

#### **Description**

Tests the null hypothesis of precision fixed in RBS models against the alternative of precision variable.

#### Usage

```
grad_test_bp(modelh0, modelh1)
grad_test_ga(modelh0, modelh1)
grad_test_ig(modelh0, modelh1)
```

6 test

```
grad_test_wei(modelh0, modelh1)

dRBS(x, mu = 1, sigma = 1, log = FALSE)

score_test_bp(modelh0, modelh1)

score_test_rbs(modelh0, modelh1)

score_test_ga(modelh0, modelh1)

score_test_ig(modelh0, modelh1)

score_test_wei(modelh0, modelh1)

wald_test(modelh1)
```

#### Arguments

modelh0 model under null hypothesis.

model h1 model under alternative hypothesis.

x vector of quantiles.

mu, sigma the (positive) location and precision parameter.

logical; The logarithm of the density is returned if the value is TRUE.

#### Value

A list with class "htest" containing the following components:

statistic the value of the test statistic.

parameter the degrees of freedom for the test statistic.

p.value the p-value for the test.

method a character string indicating what type of likelihood ratio test was performed.

data. name a character string giving the name(s) of the data

#### Author(s)

Manoel Santos-Neto <manoelferreira@uaest.ufcg.edu.br>

### **Index**

```
BP, 2
dBP(BP), 2
diag.BP, 3
diag.GA (diag.BP), 3
diag. IG (diag. BP), 3
diag.WEI (diag.BP), 3
dRBS (test), 5
envelope, 4
grad_test_bp (test), 5
grad_test_ga(test), 5
grad_test_ig (test), 5
grad_test_rbs (test), 5
grad_test_wei (test), 5
hBP (BP), 2
meanBP (BP), 2
pBP (BP), 2
plotBP (BP), 2
qBP (BP), 2
rBP(BP), 2
res_pearson(diag.BP), 3
score_test_bp (test), 5
score_test_ga (test), 5
score_test_ig (test), 5
score_test_rbs (test), 5
score_test_wei (test), 5
test, 5
wald_test (test), 5
wald_test_bp (test), 5
wald_test_ga(test), 5
wald_test_ig (test), 5
wald_test_wei (test), 5
```