Package 'ppcc'

July 23, 2025

Type Package **Version** 1.2

Date 2020-02-01					
Title Probability Plot Correlation Coefficient Test					
Author Thorsten Pohlert					
Maintainer Thorsten Pohlert <thorsten.pohlert@gmx.de></thorsten.pohlert@gmx.de>					
Description Calculates the Probability Plot Correlation Coefficient (PPCC) between a continuous variable X and a specified distribution. The corresponding composite hypothesis test that was first introduced by Filliben (1975) <doi:10.1080 00401706.1975.10489279=""> can be performed to test whether the sample X is element of either the Normal, log-Normal, Exponential, Uniform, Cauchy, Logistic, Generalized Logistic, Gumbel (GEVI), Weibull, Generalized Extreme Value, Pearson III (Gamma 2), Mielke's Kappa, Rayleigh or Generalized Logistic Distribution. The PPCC test is performed with a fast Monte-Carlo simulation.</doi:10.1080>					
Depends $R(>=3.0.0)$					
Suggests VGAM (>= 1.0), nortest(>= 1.0)					
License GPL-3					
NeedsCompilation yes					
RoxygenNote 7.0.2					
Classification/MSC-2010 62G10					
Repository CRAN					
Date/Publication 2020-02-01 20:50:02 UTC					
Contents					
ppcc-package 2 ppccTest 2 ppPositions 6	2				
Index 8	}				

ppcc-package Goodness-of-Fit Tests using the Probability Plot Correlation Coeffi- cient	ppcc-package	Goodness-of-Fit Tests using the Probability Plot Correlation Coefficient
--	--------------	--

Description

The function ppccTest performs the Probability Plot Correlation Coefficient test for various continuous distribution functions.

ppccTest

Probability Plot Correlation Coefficient Test

Description

Performs the Probability Plot Correlation Coefficnt Test of Goodness-of-Fit

Usage

```
ppccTest(
    x,
    qfn = c("qnorm", "qlnorm", "qunif", "qexp", "qcauchy", "qlogis", "qgumbel",
        "qweibull", "qpearson3", "qgev", "qkappa2", "qrayleigh", "qglogis"),
    shape = NULL,
    ppos = NULL,
    mc = 10000,
    ...
)
```

Arguments

X	a numeric vector of data values; NA values will be silently ignored.
qfn	a character vector naming a valid quantile function
shape	numeric, the shape parameter for the relevant distribution, if applicable; defaults to $\ensuremath{\text{NULL}}$
ppos	character, the method for estimating plotting point positions, default's to NULL, see Details for corresponding defaults and ppPositions for available methods
mc	numeric, the number of Monte-Carlo replications, defaults to 10000
	further arguments, currently ignored

Details

Filliben (1975) suggested a probability plot correlation coefficient test to test a sample for normality. The ppcc is defined as the product moment correlation coefficient between the ordered data $x_{(i)}$ and the order statistic medians M_i ,

$$r = \frac{\sum_{i=1}^{n} (x_{(i)} - \bar{x}) (M_i - \bar{M})}{\sqrt{\sum_{i=1}^{n} (x_{(i)} - \bar{x})^2 \sum_{j=1}^{n} (M_j - \bar{M})^2}},$$

whereas the ordered statistic medians are related to the quantile function of the standard normal distribution, $M_i = \phi^{-1}(m_i)$. The values of m_i are estimated by plotting-point position procedures (see ppPositions).

In this function the test is performed by Monte-Carlo simulation:

- 1. Calculate quantile-quantile \hat{r} for the ordered sample data x and the specified qfn distribution (with shape, if applicable) and given ppos.
- 2. Draw n (pseudo) random deviates from the specified qfn distribution, where n is the sample size of x.
- 3. Calculate quantile-quantile r_i for the random deviates and the specified qfn distribution with given ppos.
- 4. Repeat step 2 and 3 for $i = \{1, 2, ..., mc\}$.
- 5. Calculate $S = \sum_{i=1}^{n} \operatorname{sgn}(\hat{r} r_i)$ with sgn the sign-function.
- 6. The estimated p-value is p = S/mc.

The probability plot correlation coefficient is invariant for location and scale. Therefore, the null hypothesis is a composite hypothesis, e.g. $H0: X \in N(\mu, \sigma), \ \mu \in R, \ \sigma \in R_{>0}$. Furthermore, distributions with one (additional) specified shape parameter can be tested.

The magnitude of \hat{r} depends on the selected method for plotting-point positions (see ppPositions) and the sample size. Several authors extended Filliben's method to assess the goodness-of-fit to other distributions, whereas theoretical quantiles were used as opposed to Filliben's medians.

The default plotting positions (see ppPositions) depend on the selected qfn.

Distributions with none or one single scale parameter that can be tested:

Argument	Function	Default pppos	Reference
qunif	Uniform	Weibull	Vogel and Kroll (1989)
qexp	Exponential	Gringorton	
qgumbel	Gumbel	Gringorton	Vogel (1986)
qrayleigh	Rayleigh	Gringorton	

Distributions with location and scale parameters that can be tested:

Argument	Function	Default pppos	Reference
qnorm	Normal	Blom	Looney and Gulledge (1985)
qlnorm	log-Normal	Blom	Vogel and Kroll (1989)
gcauchy	Cauchy	Gringorton	

qlogis Logistic Blom

If Blom's plotting position is used for qnorm, than the ppcc-test is related to the Shapiro-Francia normality test (Royston 1993), where $W' = r^2$. See sf.test and example(ppccTest).

Distributions with additional shape parameters that can be tested:

Argument	Function	Default pppos	Reference
qweibull	Weibull	Gringorton	
qpearson3	Pearson III	Blom	Vogel and McMartin (1991)
qgev	GEV	Cunane	Chowdhury et al. (1991)
qkappa2	two-param. Kappa Dist.	Gringorton	
aglogis	Generalized Logistic	Gringorton	

If qfn = qpearson3 and shape = 0 is selected, the qnorm distribution is used. If qfn = qgev and shape = 0, the qgumbel distribution is used. If qfn = qglogis and shape = 0 is selected, the qglogis distribution is used.

Value

a list with class 'htest'

Note

As the pvalue is estimated through a Monte-Carlo simulation, the results depend on the selected seed (see set.seed) and the total number of replicates (mc).

The default of mc = 10000 re-runs is sufficient for testing the composite hypothesis on levels of $\alpha = [0.1, 0.05]$. If a level of $\alpha = 0.01$ is desired, than larger sizes of re-runs (e.g. mc = 100000) might be required.

References

- J. U. Chowdhury, J. R. Stedinger, L.-H. Lu (1991), Goodness-of-Fit Tests for Regional Generalized Extreme Value Flood Distributions, *Water Resources Research* 27, 1765–1776.
- J. J. Filliben (1975), The Probability Plot Correlation Coefficient Test for Normality, *Technometrics* 17, 111–117.
- S. Kim, H. Shin, T. Kim, J.-H. Heo (2010), Derivation of the Probability Plot Correlation Coefficient Test Statistics for the Generalized Logistic Distribution. Intern. Workshop Adv. in Stat. Hydrol., May 23 25, 2010 Taormina.
- S. W. Looney, T. R. Gulledge (1985), Use of Correlation Coefficient with Normal Probability Plots, *The American Statistician* 39, 75–79.
- P. W. Mielke (1973), Another family of distributions for describing and analyzing precipitation data. *Journal of Applied Meteorology* 12, 275–280.
- P. Royston, P. (1993), A pocket-calculator algorithm for the Shapiro-Francia test for non-normality: an application to medicine. *Statistics in Medicine* 12, 181-184.

R. M. Vogel (1986), The Probability Plot Correlation Coefficient Test for the Normal, Lognormal, and Gumbel Distributional Hypotheses, *Water Resources Research* 22, 587–590.

- R. M. Vogel, C. N. Kroll (1989), Low-flow frequency analysis using probability-plot correlation coefficients, *Journal of Water Resources Planning and Management* 115, 338–357.
- R. M. Vogel, D. E. McMartin (1991), Probability Plot Goodness-of-Fit and Skewness Estimation Procedures for the Pearson Type 3 Distribution, *Water Resources Research* 27, 3149–3158.

See Also

qqplot, qqnorm, ppoints, ppPositions, Normal, Lognormal, Uniform, Exponential, Cauchy, Logistic, qgumbel, Weibull, qgev.

Examples

```
## Filliben (1975, p.116)
## Note: Filliben's result was 0.98538
## decimal accuracy in 1975 is assumed to be less than in 2017
x < -c(6, 1, -4, 8, -2, 5, 0)
set.seed(100)
ppccTest(x, "qnorm", ppos="Filliben")
## p between .75 and .9
## see Table 1 of Filliben (1975, p.113)
set.seed(100)
## Note: default plotting position for
## qnorm is ppos ="Blom"
ppccTest(x, "gnorm")
## p between .75 and .9
## see Table 2 of Looney and Gulledge (1985, p.78)
##
##
set.seed(300)
x <- rnorm(30)
qn <- ppccTest(x, "qnorm")</pre>
## p between .5 and .75
## see Table 2 for n = 30 of Looney and Gulledge (1985, p.78)
##
## Compare with Shapiro-Francia test
if(require(nortest)){
   sn <- sf.test(x)</pre>
   print(sn)
   W <- sn$statistic
   rr <- qn$statistic^2
   names(W) <- NULL</pre>
   names(rr) <- NULL</pre>
   print(all.equal(W, rr))
}
ppccTest(x, "qunif")
ppccTest(x, "qlnorm")
old <- par()
```

6 ppPositions

```
par(mfrow=c(1,3))
xlab <- "Theoretical Quantiles"</pre>
ylab <- "Empirical Quantiles"</pre>
qqplot(x = qnorm(ppPositions(30, "Blom")),
       y = x, xlab=xlab, ylab=ylab, main = "Normal q-q-plot")
qqplot(x = qunif(ppPositions(30, "Weibull")),
       y = x, xlab=xlab, ylab=ylab, main = "Uniform q-q-plot")
qqplot(x = qlnorm(ppPositions(30, "Blom")),
       y = x, xlab=xlab, ylab=ylab, main = "log-Normal q-q-plot")
par(old)
##
if (require(VGAM)){
set.seed(300)
x \leftarrow rgumbel(30)
gu <- ppccTest(x, "qgumbel")</pre>
print(gu)
1000 * (1 - gu$statistic)
}
##
## see Table 2 for n = 30 of Vogel (1986, p.589)
## for n = 30 and Si = 0.5, the critical value is 16.9
##
set.seed(200)
x \leftarrow runif(30)
un <- ppccTest(x, "qunif")
print(un)
1000 * (1 - un$statistic)
## see Table 1 for n = 30 of Vogel and Kroll (1989, p.343)
## for n = 30 and Si = 0.5, the critical value is 10.5
set.seed(200)
x \leftarrow rweibull(30, shape = 2.5)
ppccTest(x, "qweibull", shape=2.5)
ppccTest(x, "qweibull", shape=1.5)
##
if (require(VGAM)){
set.seed(200)
x <- rgev(30, shape = -0.2)
ev <- ppccTest(x, "qgev", shape=-0.2)</pre>
print(ev)
1000 * (1 - ev$statistic)
##
## see Table 3 for n = 30 and shape = -0.2
## of Chowdhury et al. (1991, p.1770)
## The tabulated critical value is 80.
}
```

ppPositions 7

Description

Calculates plotting point positions according to different authors

Usage

```
ppPositions(
   n,
   method = c("Gringorton", "Cunane", "Filliben", "Blom", "Weibull", "ppoints")
)
```

Arguments

n numeric, the sample size
method a character string naming a valid method (see Details)

Details

The following methods can by selected:

"Gringorton" the plotting point positions are calculated as

$$m_i = (i - 0.44) / (n + 0.12)$$

"Cunane" the plotting point positions are calculated as

$$m_i = (i - 0.4) / (n + 0.2)$$

"Blom" the plotting point positions are calculated as

$$m_i = (i - 0.3175) / (n + 0.25)$$

"Filliben" the order statistic medians are calculated as:

$$m_i = \begin{cases} 1 - 0.5^{1/n} & i = 1\\ (i - 0.3175) / (n + 0.365) & i = 2, \dots, n - 1\\ 0.5^{1/n} & i = n \end{cases}$$

"ppoints" R core's default plotting point positions are calculated (see ppoints).

Value

a vector of class numeric that contains the plotting positions

Index

```
* Composite Hypothesis Test
    ppccTest, 2
* Goodness-of-Fit
    ppccTest, 2
* Homogeinity test
    ppccTest, 2
* htest
    {\tt ppccTest}, {\color{red} 2}
*\ misc
    ppPositions, 6
* npar
    ppccTest, 2
Cauchy, 5
Exponential, 5
Logistic, 5
Lognormal, 5
Normal, 5
ppcc (ppcc-package), 2
ppcc-package, 2
ppccTest, 2, 2
ppoints, 5, 7
ppPositions, 2, 3, 5, 6
qgev, 5
qgumbel, 5
qqnorm, 5
qqplot, 5
set.seed, 4
sf.test,4
Uniform, 5
Weibull, 5
```