

# Package ‘symmetry’

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**Title** Testing for Symmetry of Data and Model Residuals

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**Description** Implementations of a large number of tests for symmetry and their bootstrap variants, which can be used for testing the symmetry of IID samples around a known or unknown mean. Functions are also there for testing the symmetry of model residuals around zero. Currently, the supported models are linear models and GARCH models (fitted with the 'fGarch' package). All tests are implemented using the 'Rcpp' package which ensures great performance of the code.

**Depends** R (>= 3.1.0)

**License** MIT + file LICENSE

**Encoding** UTF-8

**LazyData** true

**Imports** Rcpp, Rdpack

**RdMacros** Rdpack

**LinkingTo** Rcpp, RcppArmadillo

**RoxygenNote** 6.1.1

**SystemRequirements** C++11

**Suggests** knitr, rmarkdown, sn, fGarch, testthat

**Roxygen** list(markdown = TRUE)

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rmixnorm	<i>Mixture of 2 normal distributions</i>
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**Description**

Generates random numbers from a mixture of 2 normal distributions

**Usage**

```
rmixnorm(n, mean1 = 0, sd1 = 1, mean2 = 0, sd2 = 1, p = 0.5)
```

**Arguments**

n	number of observations
mean1	mean of the first normal
sd1	standard deviation of the first normal
mean2	mean of the second normal
sd2	standard deviation of the second normal
p	probability of the first normal

**Value**

Vector of random numbers from the specified mixture of normals.

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rs1	<i>Azzalini skew logistic distribution</i>
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**Description**

Generates random numbers from the skew logistic distribution

**Usage**

```
rs1(n = 1, xi = 0, omega = 1, alpha = 0, dp = NULL)
```

**Arguments**

n	sample size.
xi	vector of location parameters.
omega	vector of (positive) scale parameters.
alpha	vector of slant parameters.
dp	a vector of length 3 whose elements represent the parameters described above. If dp is specified, the individual parameters cannot be set.

**Value**

Vector of random numbers from Azzalini skew logistic distribution.

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symmetry	<i>symmetry: A package which implements tests for symmetry of IID data, linear models and GARCH models</i>
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**Description**

The package contains a large number of tests for symmetry (and their bootstrap variants), which can be used to test the symmetry of IID samples or of model residuals. Currently, the supported models are linear models and GARCH models (fitted with the fGarch package). The tests are implemented using Rcpp which ensures great performance.

**Details**

To see the available tests, see [TestStatistics](#)

For documentation on how to perform the tests, see [symmetry\\_test](#)

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symmetry_test	<i>Perform symmetry tests</i>
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**Description**

This is a generic function used to perform symmetry tests on numeric vectors or objects of class lm (linear models) and objects of class fGARCH (GARCH models fitted with the fGarch package).

**Usage**

```

symmetry_test(x, ...)

## Default S3 method:
symmetry_test(x, stat, mu = NULL, bootstrap = FALSE,
  B = 1000, boot_method = c("sign", "reflect"), trim = 0, k = 0,
  ...)

## S3 method for class 'lm'
symmetry_test(x, stat, B = 1000, boot_method = c("sign",
  "reflect"), k = 0, ...)

## S3 method for class 'fGARCH'
symmetry_test(x, stat, B = 1000, burn = 0,
  boot_method = c("sign", "reflect"), k = 0, approximate = FALSE,
  ...)

```

**Arguments**

<code>x</code>	an object of class <code>numeric</code> , <code>lm</code> or <code>fGARCH</code>
<code>...</code>	not used
<code>stat</code>	a character vector indicating the test statistic to be used (see <a href="#">Available Test Statistics</a> )
<code>mu</code>	the centre parameter around which to test symmetry
<code>bootstrap</code>	a logical indicating whether to use bootstrap
<code>B</code>	the number of bootstrap replications
<code>boot_method</code>	the method of bootstrap sample generation (see <a href="#">Details</a> )
<code>trim</code>	the trim value used for estimating the centre (as used in "mean")
<code>k</code>	the <code>k</code> parameter of the statistic, ignored if the test statistic doesn't depend on a parameter (see <a href="#">Test Statistics</a> )
<code>burn</code>	the number of elements to remove from the beginning of the time series for testing
<code>approximate</code>	a logical indicating whether to use the faster approximate bootstrap method (see <a href="#">Details</a> )

**Details**

The tests are performed using bootstrap procedures or using asymptotic results, where applicable. Currently, two methods of generating a bootstrap sample from the null distribution are available. The "sign" method generates the bootstrap sample by multiplying the existing sample by -1 or 1 at random (with equal probabilities), essentially randomizing the sign of the data, giving a symmetric distribution. The "reflect" method reflects the sample around zero and samples `length(x)` elements with replacement. In practice, it has been shown that the "sign" method is almost always better, thus is the default.



```

                                skew = 1.9),
                                cond.dist = "snorm")

x <- fGarch::garchSim(specskew19, n = 500)
g <- fGarch::garchFit(~garch(1,1), x, cond.dist = "QMLE",
                      include.mean = FALSE, trace = FALSE)
symmetry_test(g, "CH", B=400, burn = 100) # slower
symmetry_test(g, "CH", B=400, burn = 100, approximate = TRUE)

```

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TestStatistics

*Available test statistics for symmetry tests*


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### Description

The list of implemented test statistics and their functions

### Usage

B1(X)

BHI(X)

BHK(X)

CH(X)

CM(X)

K2U(X)

K2(X)

KS(X)

SGN(X)

WCX(X)

MGG(X)

MOI(X, k)

MOK(X, k)

M(X)

NAI(X, k)

NAK(X, k)

S1(X, k)

S2(X, k)

T1(X, k)

T2(X, k)

### Arguments

X	the numeric vector for which to calculate the test statistic
k	the 'k' parameter in the formula (if applicable)

### Details

Below is a list of the implemented test statistics in the package. Each statistic is listed by it's name, a code string (e.g. 'B1', 'CM', 'MOI') and the formula of the statistic which is evaluated. The code string is used as an argument to the [symmetry\\_test](#) function. Some statistics depend on a parameter 'k' which can be seen from the formulas and is also passed as an argument.

Each statistic is implemented as a function with the same name as the code string, so the name of the function is passed as the argument "stat" to the [symmetry\\_test](#) function

### Value

The value of the test statistic.

### Test statistics

The list of available statistics in the format "code(s) : name (reference)"

- M : The Mira test statistic (see (Mira 1999))
- CM : The Cabilio–Masaro test statistic (see (Cabilio and Masaro 1996))
- MGG : The Miao, Gel and Gastwirth test statistic (see (Miao et al. 2006))
- B1 : The  $\sqrt{b_1}$  test statistic (see (Milošević and Obradović 2019))
- KS : The Kolmogorov–Smirnov test statistic (see (Milošević and Obradović 2019))
- SGN : The Sign test statistic (see (Milošević and Obradović 2019))
- KS : The Wilcoxon test statistic (see (Milošević and Obradović 2019))
- CH : The characterization based test statistic (see (Feuerverger et al. 1977))
- BHI : The Litvinova test statistic (see (Litvinova 2001))
- BHK : The Baringhaus and Henze test statistic (see (Baringhaus and Henze 1992))
- MOI and MOK : The Milošević and Obradović test statistics (see (Milošević and Obradović 2016))
- NAI and NAK : The Nikitin and Ahsanullah test statistics (see (Nikitin and Ahsanullah 2015))

- K2 and K2U : The Božin, Milošević, Nikitin and Obradović Kolmogorov type statistics based on V- and U- statistics respectively (see (Božin et al. 2018))
- T1, T2, S1 and S2 : The Allison and Pretorius test statistics (see (Allison and Pretorius 2017))

## References

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