# Package 'modsymmtest'

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Type Package

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**Title** Modified Symmetry Test

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<b>Description</b> Provides functionality of testing whether a dataset comes from a symmetric distribution when the center of symmetry is unknown, including modified wilcoxon test and modified sign test. In addition, sample size determination for both tests is provided.
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R topics documented:
modsymmtest-package
getV
get_quant_H0
mod.symm.test
model
n.symm.test
plasma.silicon
tree_model_sign
Index 9

2 getV

 $\begin{tabular}{ll} {\it modsymmtest-package} & {\it The modified Wilcoxon and Sign tests for symmetry about an unknown} \\ & {\it center} \\ \end{tabular}$ 

# Description

Provides functionality of testing whether a dataset comes from a symmetric distribution when the center of symmetry is unknown, including modified wilcoxon test and modified sign test. In addition, sample size determination is provided.

# Author(s)

Jiaojiao Zhou, Xinyu Gao, Albert Vexler

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#### References

Vexler, A., Gao, X., & Zhou, J. (2023). How to implement signed-rank wilcox. test () type procedures when a center of symmetry is unknown. Computational Statistics & Data Analysis, 107746.

Gastwirth, J. L. (1971). On the sign test for symmetry. Journal of the American Statistical Association, 66(336), 821-823.

# **Examples**

```
## Not run:
    data("plasma.silicon")
    post <- plasma.silicon$postoperative
    pre <- plasma.silicon$preoperative
    # post <- c(0.21,0.24,0.1,0.12,0.28,0.25,0.22,0.21,0.22,0.23,0.22,0.24,0.45,0.38,
    # 0.23,0.22,0.18,0.15,0.04,0.14,0.24,0.20,0.24,0.18,0.19,0.15,0.26,0.3,0.22,0.24)
# pre <- c(0.15,0.13,0.39,0.2,0.39,0.42,0.24,0.18,0.26,0.12,0.1,0.11,0.19, 0.15,0.27,
    # 0.28,0.11,0.11,0.18,0.18,0.24,0.48,0.27,0.22,0.18,0.19,0.32,0.31,0.19,0.21)
    mod.symm.test(post, pre)

## End(Not run)</pre>
```

getV

Getting asymptotic variance of modified wilcox test statistic

## **Description**

Getting asymptotic variance of modified wilcox test statistic

## Usage

```
getV(x_{,} y_{-} = NULL)
```

get\_quant\_H0 3

# **Arguments**

x\_ A data set

y\_ Another data set, default is NULL

#### Value

Asymptotic variance of modified wilcoxon test statistic

get\_quant\_H0

Title

# **Description**

Title

# Usage

```
get_quant_H0(x)
```

# **Arguments**

Χ

pilot sample

mod.symm.test

Wilcoxon and Sign tests for symmetry about an unknown center

# Description

R built-in function 'wilcox.test()' is designed to perform both the one- and two-sample Wilcoxon test for symmetry under the assumption that the center of symmetry is specified. The procedure 'mod.symm.test()' extends the capabilities of 'wilcox.test()' for situations where the center of symmetry is unknown. Such cases can be found in e.g., regression residuals evaluations, as well as in the book 'Nonparametric statistical methods using R' by Kloke and McKean, and in the scholarly work of Gastwirth.

# Usage

```
mod.symm.test(
    x,
    y = NULL,
    alternative = c("two.sided", "left.skewed", "right.skewed"),
    method = "wilcox"
)
```

4 mod.symm.test

## **Arguments**

х	numeric vector of data values. Non-finite (e.g., infinite or missing) values will be omitted.
У	an optional numeric vector of data values: as with x non-finite values will be omitted.
alternative	a character string specifying the alternative hypothesis, must be one of "two sided" (default), "right.skewed", or "left.skewed". You can specify just the initial letter. "right.skewed": test whether positively skewed, "left.skewed": test whether negatively skewed.
method	a character string specifying which symmetry test to be used, "wilcox" refers to Wilcoxon signed-rank test, and "sign" is sign test.

#### **Details**

When "wilcox", the default method, is used, the test statistic W has the form of the Wilcoxon test statistic with unknown center of symmetry replaced by its sample estimator. For details, see Vexler et al. (2023)

When method="sign", the test statistic S is the total number of the observations that smaller than sample mean. For more details, see Gastwitrh (1971).

#### Value

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

- statistic the value of the test statistic.
- var the asymptotic variance.
- alternative a character string describing the alternative hypothesis.
- p.value the p-value for the test.
- method the type of test applied.

## Author(s)

Jiaojiao Zhou, Xinyu Gao, Albert Vexler

## References

Vexler, A., Gao, X., & Zhou, J. (2023). How to implement signed-rank 'wilcox.test()' type procedures when a center of symmetry is unknown. Computational Statistics & Data Analysis, 107746.

Gastwirth, J. L. (1971). On the Sign Test for Symmetry. Journal of the American Statistical Association, 66(336), 821-823.

## **Examples**

```
## Not run:
    # A study measures the plasma silicon levels before and after silicone implants surgery
# in 30 women to evaluate the effect of the surgery. Informally speaking, we can be interested
# in that there is an unknown constant shift such that the plasma silicon level of post-surgery
# can be explained completely based on that of pre-surgery. This can be stated as the null hypothesis
# `H_0` The difference of plasma silicon level between post-surgery pre-surgery has a symmetric distribution
# around a shift that is unknown.
data("plasma.silicon")
```

model 5

```
post <- plasma.silicon$postoperative</pre>
pre <- plasma.silicon$preoperative</pre>
# post <- c(0.21,0.24,0.1,0.12,0.28,0.25,0.22,0.21,0.22,0.23,0.22,0.24,0.45,0.38,
        0.23, 0.22, 0.18, 0.15, 0.04, 0.14, 0.24, 0.2, 0.24, 0.18, 0.19, 0.15, 0.26, 0.3, 0.22, 0.24)
# pre <- c(0.15,0.13,0.39,0.2,0.39,0.42,0.24,0.18,0.26,0.12,0.1,0.11,0.19, 0.15,0.27,
            0.28, 0.11, 0.11, 0.18, 0.18, 0.24, 0.48, 0.27, 0.22, 0.18, 0.19, 0.32, 0.31, 0.19, 0.21)
mod.symm.test(x=post, y=pre, alternative ="two.sided", method = "wilcox")
Result:
Modified Wilcoxon signed-rank test
data: post and pre
W = 238, p-value = 0.767
alternative hypothesis: two.sided
Interpretation:
Test statistic `W` is the number of walsh average higher than sample mean, see more details
```

in paper authored by Vexler, etc.

p-value is 0.767, which implies there is no clue to reject the null hypothesis that the distribution of the difference of plasma silicon levels before and after silicone implants surgery is symmetric.

## End(Not run)

model

Tree model

# **Description**

Tree model object for wilcox method to find the optimal alp

#### Usage

model

#### **Format**

An rpart object that contains a tree model which fits alp to npilot, sk, kur, sd and distance\_p2\_half.

n.symm.test

Sample size determination for nonparametric tests of symmetry when the center is unknown

#### **Description**

Determine the sample size required for one-sample Wilcoxon signed-rank test or Sign test with an unknown symmetry center estimated by sample mean given a target power. The function uses learning sample in order to predict (calculate) sample size needed to reach a preassumed power based on the underlying data that is exemplified by the learning sample.

6 n.symm.test

#### **Usage**

```
n.symm.test(
    x,
    sig.level = 0.05,
    power = 0.8,
    method = "wilcox",
    alternative = c("two.sided", "right.skewed", "left.skewed")
)
```

# **Arguments**

x learning sample data, numeric vector of data values
sig.level significance level (Type I error probability), the default value is 0.05
power power of test (1 minus Type II error probability), the default value is 0.8
method a character string specifying which symmetry test to be used, "wilcox" refers to Wilcoxon signed-rank test, and "sign" sign test
alternative a character string specifying the alternative hypothesis, must be one of "two.sided" (default), "right.skewed", or "left.skewed". Can be abbreviated.

"two.sided": test whether skewed,
"right.skewed": test whether positively skewed,
"left.skewed": test whether negatively skewed

#### **Details**

A Normal approximation to the power requires specification of some unknown quantities in the non-parametric context. In this regard, empirical smoothed CDF and Bootstrap methods were leveraged to estimate these quantities using learning sample 'x'.

Remark: If the test provides a power, say, P, based on the learning data and P is higher than the target power, a warning message will be shown. However, a needed sample size N to reach the target power will be conducted.

## Value

A list of class "power.htest" containing the following components:

- N sample size estimated
- sig.level significance level (Type I error probability)
- power power of test (1 minus Type II error probability)
- method the test method applied
- alternative one- or two-sided test. Can be abbreviated

#### References

Chakraborti, S., Hong, B., & van de Wiel, M. A. (2006). A Note on Sample Size Determination for a Nonparametric Test of Location. Technometrics, 48(1), 88-94.

Vexler, A., Gao, X., & Zhou, J. (2023). How to implement signed-rank 'wilcox.test()' type procedures when a center of symmetry is unknown. Computational Statistics & Data Analysis, 107746.

Gastwirth, J. L. (1971). On the Sign Test for Symmetry. Journal of the American Statistical Association, 66(336), 821-823.

plasma.silicon 7

#### **Examples**

```
## Not run:
data("plasma.silicon")
post <- plasma.silicon$postoperative</pre>
 pre <- plasma.silicon$preoperative</pre>
diff <- post - pre
n.symm.test(diff, sig.level = 0.05, power = 0.5, method = "wilcox", alternative = "two.sided")
 Sample size calculation under wilcox procedure
          N = 83
  sig.level = 0.05
      power = 0.5
       type = wilcox
alternative = two.sided
 Interpretation:
Given the pilot sample `diff` and the significance level 0.05. The sample size of the data that is expected to
provide the target power 0.5 of the Wilcoxon test procedure is computed as 83.
## End(Not run)
```

# **Description**

plasma.silicon

A study was conducted to measure the plasma silicon level in blood within 30 women who have been taken a silicone implants surgery. Plasma silicon levels (microg per gram dry weight) were taken prior to surgical placement of the implants. A post-surgery washout period was allowed, and the plasma silicon levels were retaken.

Dataset: plasma.silicon

# Usage

```
load(plasma.silicon.rda)
```

#### **Format**

A data frame with 30 rows and 3 variables:

PatientNo Patient Number.

preoperative pre-operative measurement of plasma silicion level.

postoperative post-operative measurement of plasma silicion level.

## **Source**

Riffenburgh, R. H., & Gillen, D. L. (2020). Statistics in medicine. Academic press. Appendix 2 databases

#### References

Riffenburgh, R. H., & Gillen, D. L. (2020). Statistics in medicine. Academic press.

8 tree\_model\_sign

tree\_model\_sign

 $Tree\_model\_sign$ 

# Description

Tree model object to find the optimal alp for sign test method.

# Usage

```
tree_model_sign
```

# **Format**

An rpart object that contains a tree model which fits alp to npilot, sk, kur, sd.

# **Index**

```
* datasets
    plasma.silicon, 7
* package
    modsymmtest-package, 2
* rpart
    model, 5
    tree_model_sign, 8

get_quant_H0, 3
getV, 2

mod.symm.test, 3
model, 5
modsymmtest-package, 2

n.symm.test, 5

plasma.silicon, 7

tree_model_sign, 8
```