

Package ‘vartest’

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Contents

ansari.test	2
bartlettstest	3
capon.test	5
cochrans.test	6
david.barton.test	7
duran.test	9
f.test	10
fk.test	11
g.test	12
hartley.test	14

klotz.test	15
levene.test	16
mood.test	18
mzv.test	19
obrien.test	21
print.summary.vht	22
select_vartest	23
siegel.tukey.test	24
summary.vht	25
talwar.gentle.test	26
zv.test	27
Index	29

ansari.test	<i>Ansari Bradley Test</i>
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Description

ansari.test performs Ansari Bradley variance homogeneity test.

Usage

```
ansari.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters (Conover et al., 1981).

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Ansari Bradley Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23**:4, 351-361.

See Also

[ansari_test](#)

Examples

```
library(vartest)

ansari.test(Sepal.Length ~ Species, data = iris)

out <- ansari.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

bartletts.test

Bartlett's Test

Description

bartletts.test performs Bartlett variance homogeneity test.

Usage

```
bartletts.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Bartlett's Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

- Lee, H.B., Katz, G.S., Restori, A.F. (2010). A Monte Carlo Study of Seven Homogeneity of Variance Tests. *Journal of Mathematics and Statistics*, **6:3**, 359-366.
- Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23:4**, 351-361.

See Also

[bartlett.test](#)

Examples

```
library(vartest)

bartletts.test(Sepal.Length ~ Species, data = iris)

out <- bartletts.test(Sepal.Length ~ Species, data = iris, verbose = FALSE)
summary(out)
```

`capon.test`*Capon Test*

Description

`capon.test` performs Capon variance homogeneity test.

Usage

```
capon.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

<code>formula</code>	a formula of the form <code>lhs ~ rhs</code> where <code>lhs</code> gives the sample values and <code>rhs</code> the corresponding groups.
<code>data</code>	a tibble or data frame containing the variables in <code>formula</code> .
<code>alpha</code>	the level of significance to assess variance homogeneity. Default is set to <code>alpha = 0.05</code> .
<code>na.rm</code>	a logical value indicating whether NA values should be stripped before the computation proceeds.
<code>verbose</code>	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters (Conover et al., 1981).

Value

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

<code>statistic</code>	the corresponding test statistic.
<code>parameter</code>	the parameter(s) of the approximate chi-squared distribution of the test statistic.
<code>p.value</code>	the p-value of the test.
<code>method</code>	the character string "Capon Test".
<code>data</code>	a data frame containing the variables in which NA values (if exist) are removed.
<code>formula</code>	a formula of the form <code>lhs ~ rhs</code> where <code>lhs</code> gives the sample values and <code>rhs</code> the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Capon, J. (1961). Asymptotic Efficiency of Certain Locally Most Powerful Rank Tests. *The Annals of Mathematical Statistics*, **32:1**, 88-100.

Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23:4**, 351-361.

Examples

```
library(vartest)

capon.test(Sepal.Length ~ Species, data = iris)

out <- capon.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

cochrans.test	<i>Cochran's C Test</i>
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Description

cochrans.test performs Cochran's C variance homogeneity test.

Usage

```
cochrans.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate F distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Cochran's C Test".

data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Wang, Y., Gil, P.R., Chen, Y.H., Kromrey, J.D., Kim, E.S., Pham, T., Nguyen, D., Romano, J.L. (2017). Comparing the Performance of Approaches for Testing the Homogeneity of Variance Assumption in One-Factor Anova Models. *Educational and Psychological Measurement*, **77:2**, 305-329.

Cochran, W.G. (1941). The Distribution of The Largest of a set of Estimated Variances as a Fraction of Their Total. *Annals of Eugenics*, **11:1**, 47-52.

See Also

[cochran.test](#)

Examples

```
library(vartest)

cochrans.test(Sepal.Length ~ Species, data = iris)

out <- cochrans.test(Sepal.Length ~ Species, data = iris, verbose = FALSE)
summary(out)
```

david.barton.test	<i>David Barton Test</i>
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Description

david.barton.test performs David Barton variance homogeneity test.

Usage

```
david.barton.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.

na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters (Conover et al., 1981).

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "David Barton Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

- Barton, D.E., David, F.N. (1958). A Test For Birth Order Effect. *Annals of Human Genetics*, **22:3**, 250-257.
- Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23:4**, 351-361.
- Gibbons, J.D., Chakraborti, S. (2010). Nonparametric Statistical Inference. *CRC Press*, **5th Ed.**, 316-320.

Examples

```
library(vartest)

david.barton.test(Sepal.Length ~ Species, data = iris)

out <- david.barton.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

duran.test	<i>Duran Test</i>
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Description

duran.test performs Duran variance homogeneity test.

Usage

```
duran.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters. This median-centered variant of the test is also referred to as the Conover-Iman test in literature (Conover and Iman, 1978).

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Duran Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Conover, W.J., Iman, R.L.(1978). Some Exact Tables For The Squared Ranks Test. *Communications in Statistics - Simulation and Computation*, **B7(5)**, 491-513.

Examples

```
library(vartest)

duran.test(Sepal.Length ~ Species, data = iris)

out <- duran.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

f.test	<i>Fisher's Test</i>
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Description

f.test performs Fisher's variance homogeneity test.

Usage

```
f.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate F distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Fisher's Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Gorbunova, A.A., Lemeshko, B.Y. (2012). Application of Parametric Homogeneity of Variances Tests under Violation of Classical Assumption. *2nd Stochastic Modeling Techniques and Data Analysis International Conference*, **5:8**, 253-260.

Examples

```
library(vartest)

f.test(Sepal.Length ~ Species, data = iris)

out <- f.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

fk.test

Fligner-Killeen Test

Description

fk.test performs Fligner-Killeen variance homogeneity test.

Usage

```
fk.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters (Conover et al., 1981).

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Fligner-Killeen Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23:4**, 351-361.

Fligner, M.A., Killeen, T.J.(1976). Distribution-Free Two-Sample Tests for Scale. *Journal of the American Statistical Association*, **71:353**, 210-213.

See Also

[fligner_test](#)

Examples

```
library(vartest)

fk.test(Sepal.Length ~ Species, data = iris)

out <- fk.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

g.test

G Test

Description

g.test performs G variance homogeneity test.

Usage

```
g.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate F distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "G Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

- UE't Lam, R. (2010). Scrutiny of Variance Results for Outliers: Cochran's Test Optimized. *Analytica Chimica Acta*, **659**(1-2), 68-84.
- Wang, Y., Gil, P.R., Chen, Y.H., Kromrey, J.D., Kim, E.S., Pham, T., Nguyen, D., Romano, J.L. (2017). Comparing the Performance of Approaches for Testing the Homogeneity of Variance Assumption in One-Factor Anova Models. *Educational and Psychological Measurement*, **77**:2, 305-329.

Examples

```
library(vartest)

g.test(Sepal.Length ~ Species, data = iris)

out <- g.test(Sepal.Length ~ Species, data = iris, verbose = FALSE)
summary(out)
```

hartley.test	<i>Hartley's Maximum F-Ratio Test</i>
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Description

hartley.test performs Hartley's Maximum F-Ratio variance homogeneity test.

Usage

```
hartley.test(formula, data, size = "mean", alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
size	a character string to define how to number of group observation. "mean": mean, "harmonic": harmonic mean, "maxn": maximum n, and "minvar": minimum variance.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the maximum F-ratio distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Hartley's Maximum F-Ratio Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Gorbunova, A.A., Lemeshko, B.Y. (2012). Application of Parametric Homogeneity of Variances Tests under Violation of Classical Assumption. *2nd Stochastic Modeling Techniques and Data Analysis International Conference*, **5:8**, 253-260.

Bhandary, M., Dai, H. (2008). An Alternative Test for the Equality of Variances for Several Populations When the Underlying Distributions are Normal. *Communications in Statistics-Simulation and Computation*, **38:1**, 109-117.

See Also

[hartleyTest](#)

Examples

```
library(vartest)

hartley.test(Sepal.Length ~ Species, data = iris, size = "mean")

out <- hartley.test(Sepal.Length ~ Species, data = iris, size = "mean")
summary(out)
```

klotz.test	<i>Klotz Test</i>
------------	-------------------

Description

klotz.test performs Klotz variance homogeneity test.

Usage

```
klotz.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters (Conover et al., 1981).

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Klotz Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23**:4, 351-361.

Klotz, J. (1962). Nonparametric Tests for Scale. *The Annals of Mathematical Statistics*, **33**:2, 498-512.

See Also

[klotz_test](#)

Examples

```
library(vartest)

klotz.test(Sepal.Length ~ Species, data = iris)

out <- klotz.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

levene.test

Levene's Test

Description

levene.test performs Levene variance homogeneity test.

Usage

```
levene.test(formula, data, center = "mean", deviation = "absolute",
  trim.rate = 0.25, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
center	a character string to define how to center. "mean": mean, "median": median, and "trim.mean": trimmed mean.
deviation	a character string to define how to specify the deviation. "absolute": absolute deviation and "squared": squared deviation.
trim.rate	the rate of observations trimmed from each tail of the distribution. Default is set to 0.25.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate F distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Levene's Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

- Wang, Y., Gil, P.R., Chen, Y.H., Kromrey, J.D., Kim, E.S., Pham, T., Nguyen, D., Romano, J.L. (2017). Comparing the Performance of Approaches for Testing the Homogeneity of Variance Assumption in One-Factor Anova Models. *Educational and Psychological Measurement*, **77:2**, 305-329.
- Brown, M.B., Forsythe, A.B. (1974). Robust Tests for the Equality of Variances. *Journal of the American Statistical Association*, **69:346**, 364-367.

See Also

[leveneTest](#)

Examples

```
library(vartest)

levene.test(Sepal.Length ~ Species, data = iris, center="median", deviation="absolute")
levene.test(Sepal.Length ~ Species, data = iris, center="median", deviation="squared")
levene.test(Sepal.Length ~ Species, data = iris, center="mean", deviation="absolute")
levene.test(Sepal.Length ~ Species, data = iris, center="mean", deviation="squared")
levene.test(Sepal.Length ~ Species, data = iris, center="trim.mean", deviation="absolute")
levene.test(Sepal.Length ~ Species, data = iris, center="trim.mean", deviation="squared")

out <- levene.test(Sepal.Length~Species, data=iris, center="median", deviation="absolute")
summary(out)
```

mood.test

Mood Test

Description

mood.test performs Mood variance homogeneity test.

Usage

```
mood.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters (Conover et al., 1981).

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Mood Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23**:4, 351-361.

Mood, A.M. (1954). On the Asymptotic Efficiency of Certain Nonparametric Two-Sample Tests. *The Annals of Mathematical Statistics*, **25**:3, 514-522.

See Also

[mood_test](#)

Examples

```
library(vartest)

mood.test(Sepal.Length ~ Species, data = iris)

out <- mood.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

mzv.test

Modified Z Variance Test

Description

mzv.test performs Modified Z Variance variance homogeneity test.

Usage

```
mzv.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate F distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Modified Z Variance Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

- Wang, Y., Gil, P.R., Chen, Y.H., Kromrey, J.D., Kim, E.S., Pham, T., Nguyen, D., Romano, J.L. (2017). Comparing the Performance of Approaches for Testing the Homogeneity of Variance Assumption in One-Factor Anova Models. *Educational and Psychological Measurement*, **77:2**, 305-329.
- Overall, J.E., Woodward, J.A. (1974). A Simple Test for Heterogeneity of Variance in Complex Factorial Designs. *Psychometrika*, **39:3**, 311-318.

Examples

```
library(vartest)

mzv.test(Sepal.Length ~ Species, data = iris)

out <- mvz.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

obrien.test*O'Brien Test*

Description

obrien.test performs O'Brien variance homogeneity test.

Usage

```
obrien.test(formula, data, center = "mean", trim.rate = 0.25, alpha = 0.05,  
             na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
center	a character string to define how to center. "mean": mean, "median": median, and "trim.mean": trimmed mean.
trim.rate	the rate of observations trimmed from each tail of the distribution. Default is set to 0.25.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate F distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "O'Brien Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

O'Brien, R.G. (1981). A Simple Test for Variance Effects in Experimental Designs. *Psychological Bulletin*, **89:3**, 570-574.

Gorbunova, A.A., Lemesko, B.Y. (2012). Application of Parametric Homogeneity of Variances Tests under Violation of Classical Assumption. *2nd Stochastic Modeling Techniques and Data Analysis International Conference*, **5:8**, 253-260.

Examples

```
library(vartest)

obrien.test(Sepal.Length ~ Species, data = iris, center = "mean")

out <- obrien.test(Sepal.Length ~ Species, data = iris, center = "mean")
summary(out)
```

print.summary.vht

Print Method for Summary of Variance Homogeneity Test Results

Description

Prints the formatted summary of a vht object to the console.

Usage

```
## S3 method for class 'summary.vht'
print(x, ...)
```

Arguments

x An object returned by summary.vht().

... Additional arguments.

Author(s)

Merve Kasikci and Osman Dag

See Also

[summary.vht](#)

Examples

```
out <- bartlettstest(Sepal.Length ~ Species, data = iris, verbose = FALSE)
summary(out)
```

select_vartest*Data-Driven Selection of the Optimal Variance Homogeneity Test*

Description

select_vartest performs an automated, simulation-based evaluation to identify the optimal variance homogeneity test for a dataset.

Usage

```
select_vartest(formula, data, nrep = 1000, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
nrep	an integer specifying the number of Monte Carlo replications to perform for the simulation. The default is 1000.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

The function calculates the empirical moments of the input data and uses the Pearson distribution system to generate random datasets. By evaluating 28 distinct methodological modifications of variance homogeneity tests on these generated datasets, the function recommends the method that yields the highest adjusted power (with the lowest Type I error).

Value

A data frame presenting empirical power, Type I error rate, and adjusted power for each variance homogeneity test.

Author(s)

Muhammed Ali Yilmaz and Osman Dag

Examples

```
library(vartest)

set.seed(1234)
select_vartest(Sepal.Length ~ Species, data = iris, nrep = 1000)
```

siegel.tukey.test	<i>Siegel Tukey Test</i>
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Description

siegel.tukey.test performs Siegel Tukey variance homogeneity test.

Usage

```
siegel.tukey.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters (Conover et al., 1981).

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Siegel Tukey Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

- Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23:4**, 351-361.
- Gibbons, J.D., Chakraborti, S. (2010). Nonparametric Statistical Inference. *CRC Press*, **5th Ed.**, 316-320.
- Siegel, S., Tukey, J.W. (1960). A Nonparametric Sum of Ranks Procedure for Relative Spread in Unpaired Samples. *Journal of the American Statistical Association*, **55:291**, 429-445.

See Also

[GSTTest](#)

Examples

```
library(vartest)

siegel.tukey.test(Sepal.Length ~ Species, data = iris)

out <- siegel.tukey.test(Sepal.Length ~ Species, data = iris)
summary(out)
```

summary.vht

Summary Method for Variance Homogeneity Test Results

Description

Provides a concise summary of the results from a variance homogeneity test in the package.

Usage

```
## S3 method for class 'vht'
summary(object, ...)
```

Arguments

object	An object of class vht, typically returned by variance homogeneity tests.
...	Additional arguments.

Details

This method is specifically designed for objects of class vht. It prints test method, dependent variable, grouping variable, test statistic, degrees of freedom, and p-value.

Value

Prints a summary to the console.

Author(s)

Merve Kasikci and Osman Dag

Examples

```
out <- bartlettstest(Sepal.Length ~ Species, data = iris, verbose = FALSE)
summary(out)
```

talwar.gentle.test	<i>Talwar and Gentle Test</i>
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Description

talwar.gentle.test performs Talwar and Gentle variance homogeneity test.

Usage

```
talwar.gentle.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Details

Prior to testing for variance homogeneity, the data within each group are centered by their corresponding group median to eliminate the confounding effects of location parameters (Conover et al., 1981).

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate chi-squared distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Talwar and Gentle Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Conover, W.J., Johnson, M.E., Johnson, M.M. (1981). A Comparative Study of Tests for Homogeneity of Variances, with Applications to the Outer Continental Shelf Bidding Data. *Technometrics*, **23:4**, 351-361.

Talwar, P.P., Gentle, J.E. (1976). A Robust Test for the Homogeneity of Scales. *Communications in Statistics - Theory and Methods*, **6:4**, 363-369.

Examples

```
library(vartest)

talwar.gentle.test(Sepal.Length ~ Species, data = iris)

out <- talwar.gentle.test(Sepal.Length ~ Species, data = iris, verbose = FALSE)
summary(out)
```

zv.test

Z Variance Test

Description

zv.test performs Z variance homogeneity test.

Usage

```
zv.test(formula, data, alpha = 0.05, na.rm = TRUE, verbose = TRUE)
```

Arguments

formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.
data	a tibble or data frame containing the variables in formula.
alpha	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
verbose	a logical for printing output to R console.

Value

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

statistic	the corresponding test statistic.
parameter	the parameter(s) of the approximate F distribution of the test statistic.
p.value	the p-value of the test.
method	the character string "Z Variance Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	a formula of the form lhs ~ rhs where lhs gives the sample values and rhs the corresponding groups.

Author(s)

Gozde Cosar and Osman Dag

References

Gorbunova, A.A., Lemeshko, B.Y. (2012). Application of Parametric Homogeneity of Variances Tests under Violation of Classical Assumption. *2nd Stochastic Modeling Techniques and Data Analysis International Conference*, **5:8**, 253-260.

Wang, Y., Gil, P.R., Chen, Y.H., Kromrey, J.D., Kim, E.S., Pham, T., Nguyen, D., Romano, J.L. (2017). Comparing the Performance of Approaches for Testing the Homogeneity of Variance Assumption in One-Factor Anova Models. *Educational and Psychological Measurement*, **77:2**, 305-329.

Overall, J.E., Woodward, J.A. (1974). A Simple Test for Heterogeneity of Variance in Complex Factorial Designs. *Psychometrika*, **39:3**, 311-318.

Examples

```
library(vartest)

zv.test(Sepal.Length ~ Species, data = iris)

out <- zv.test(Sepal.Length ~ Species, data = iris, verbose = FALSE)
summary(out)
```

Index

* functions

- ansari.test, [2](#)
- bartlettts.test, [3](#)
- capon.test, [5](#)
- cochrans.test, [6](#)
- david.barton.test, [7](#)
- duran.test, [9](#)
- f.test, [10](#)
- fk.test, [11](#)
- g.test, [12](#)
- hartley.test, [14](#)
- klotz.test, [15](#)
- levene.test, [16](#)
- mood.test, [18](#)
- mzv.test, [19](#)
- obrien.test, [21](#)
- print.summary.vht, [22](#)
- select_vartest, [23](#)
- siegel.tukey.test, [24](#)
- summary.vht, [25](#)
- talwar.gentle.test, [26](#)
- zv.test, [27](#)

- ansari.test, [2](#)
- ansari_test, [3](#)

- bartlett.test, [4](#)
- bartlettts.test, [3](#)

- capon.test, [5](#)
- cochran.test, [7](#)
- cochrans.test, [6](#)

- david.barton.test, [7](#)
- duran.test, [9](#)

- f.test, [10](#)
- fk.test, [11](#)
- fligner_test, [12](#)

- g.test, [12](#)

- GSTTest, [25](#)

- hartley.test, [14](#)
- hartleyTest, [15](#)

- klotz.test, [15](#)
- klotz_test, [16](#)

- levene.test, [16](#)
- leveneTest, [17](#)

- mood.test, [18](#)
- mood_test, [19](#)
- mzv.test, [19](#)

- obrien.test, [21](#)

- print.summary.vht, [22](#)

- select_vartest, [23](#)
- siegel.tukey.test, [24](#)
- summary.vht, [22](#), [25](#)

- talwar.gentle.test, [26](#)

- zv.test, [27](#)