

Package ‘varptest’

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Description Performs 20 omnibus tests for testing the composite hypothesis of variance homogeneity.

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adjusted.taha.test *Adjusted Taha Test*

Description

adjusted.taha.test performs Adjusted Taha variance homogeneity test.

Usage

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

Arguments

formula	a formula of the form <code>lhs ~ rhs</code> where <code>lhs</code> gives the sample values and <code>rhs</code> the corresponding groups.
data	a tibble or data frame containing the variables in <code>formula</code> .
alpha	the level of significance to assess variance homogeneity. Default is set to <code>alpha = 0.05</code> .
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 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 "Adjusted Taha Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	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

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

Examples

```
library(vartest)

adjusted.taha.test(Sepal.Length ~ Species, data = iris)
```

`ansari.test`

Ansari Bradley Test

Description

`ansari.test` performs Ansari Bradley variance homogeneity test.

Usage

```
ansari.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.

Value

A list 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 "Ansari Bradley 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

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)
```

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

- | | |
|----------------------|--|
| <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. |

Value

A list 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 "Bartlett's 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

- 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(varptest)

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

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

formula	a formula of the form <code>lhs ~ rhs</code> where <code>lhs</code> gives the sample values and <code>rhs</code> the corresponding groups.
data	a tibble or data frame containing the variables in <code>formula</code> .
alpha	the level of significance to assess variance homogeneity. Default is set to <code>alpha = 0.05</code> .
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 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 "Capon Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	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)
```

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

Description

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

Usage

```
cochrans.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.

Value

A list containing the following components:

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

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(varptest)

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

david.barton.test *David Barton Test*

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

- Gibbons, J.D., Chakraborti, S. (2010). Nonparametric Statistical Inference. *CRC Press*, **5th Ed.**, 316-320.
- 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.

Examples

```
library(vartest)

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

duran.test

Duran Test

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

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)
```

f.test

Fisher's Test

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

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)
```

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

<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.

Value

A list 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 "Fligner-Killeen 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

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

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[fligner_test](#)**Examples**

```
library(vartest)

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

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

<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.

Value

A list containing the following components:

<code>statistic</code>	the corresponding test statistic.
<code>parameter</code>	the parameter(s) of the approximate F distribution of the test statistic.
<code>p.value</code>	the p-value of the test.
<code>method</code>	the character string "G 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

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

hartley.test

Hartley's Maximum F-Ratio Test

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 <code>lhs ~ rhs</code> where <code>lhs</code> gives the sample values and <code>rhs</code> the corresponding groups.
data	a tibble or data frame containing the variables in <code>formula</code> .
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 containing the following components:

<code>statistic</code>	the corresponding test statistic.
<code>parameter</code>	the parameter(s) of the approximate F distribution of the test statistic.
<code>p.value</code>	the p-value of the test.
<code>method</code>	the character string "Hartley's Maximum F-Ratio 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

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(varptest)

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

klotz.test

Klotz Test

Description

`klotz.test` performs Klotz variance homogeneity test.

Usage

```
klotz.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.

Value

A list 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 "Klotz 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

- 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)
```

<code>levene.test</code>	<i>Levene's Test</i>
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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

<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>center</code>	a character string to define how to center. "mean": mean, "median": median, and "trim.mean": trimmed mean.
<code>deviation</code>	a character string to define how to specify the deviation. "absolute": absolute deviation and "squared": squared deviation.
<code>trim.rate</code>	the rate of observations trimmed from each tail of the distribution. Default is set to 0.25.
<code>alpha</code>	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
<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.

Value

A list containing the following components:

<code>statistic</code>	the corresponding test statistic.
<code>parameter</code>	the parameter(s) of the approximate F distribution of the test statistic.
<code>p.value</code>	the p-value of the test.
<code>method</code>	the character string "Levene's 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

- 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")
```

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

- 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)
```

`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

<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.

Value

A list containing the following components:

<code>statistic</code>	the corresponding test statistic.
<code>parameter</code>	the parameter(s) of the approximate F distribution of the test statistic.
<code>p.value</code>	the p-value of the test.
<code>method</code>	the character string "Modified Z Variance 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

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(varptest)

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

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

<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>center</code>	a character string to define how to center. "mean": mean, "median": median, and "trim.mean": trimmed mean.
<code>trim.rate</code>	the rate of observations trimmed from each tail of the distribution. Default is set to 0.25.
<code>alpha</code>	the level of significance to assess variance homogeneity. Default is set to alpha = 0.05.
<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.

Value

A list containing the following components:

<code>statistic</code>	the corresponding test statistic.
<code>parameter</code>	the parameter(s) of the approximate F distribution of the test statistic.
<code>p.value</code>	the p-value of the test.
<code>method</code>	the character string "O'Brien 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

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

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

siegel.tukey.test *Siegel Tukey Test*

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

<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.

Value

A list 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 "Siegel Tukey 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

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

taha.test

Taha Test

Description

taha.test performs Taha variance homogeneity test.

Usage

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

Arguments

formula	a formula of the form <code>lhs ~ rhs</code> where <code>lhs</code> gives the sample values and <code>rhs</code> the corresponding groups.
data	a tibble or data frame containing the variables in <code>formula</code> .
alpha	the level of significance to assess variance homogeneity. Default is set to <code>alpha = 0.05</code> .
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 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 "Taha Test".
data	a data frame containing the variables in which NA values (if exist) are removed.
formula	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

- 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.
- Taha, M.A.H. (1964). Rank Test for Scale Parameter for Asymmetrical One-Sided Distributions. *Annales de l'ISUP*, **13:3**, 169-180.

See Also

[taha_test](#)

Examples

```
library(vartest)

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

talwar.gentle.test *Talwar and Gentle Test*

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

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

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

<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.

Value

A list containing the following components:

<code>statistic</code>	the corresponding test statistic.
<code>parameter</code>	the parameter(s) of the approximate F distribution of the test statistic.
<code>p.value</code>	the p-value of the test.
<code>method</code>	the character string "Z Variance 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

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

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