

Package: mvShapiroTest (via r-universe)

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Type Package

Title Generalized Shapiro-Wilk test for multivariate normality

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Description This package implements the generalization of the Shapiro-Wilk test for multivariate normality proposed by Villasenor-Alva and Gonzalez-Estrada (2009).

License GPL (>= 3)

Depends stats, datasets

LazyLoad yes

NeedsCompilation no

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Repository <https://elizagestrada.r-universe.dev>

RemoteUrl <https://github.com/cran/mvShapiroTest>

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`mvShapiro.Test`*Generalized Shapiro-Wilk test for multivariate normality*

Description

Given a d-dimensional random sample of size n, this function computes the test statistic and p-value of the Shapiro-Wilk test for multivariate normality proposed by Villasenor-Alva and Gonzalez-Estrada (2009).

Usage`mvShapiro.Test(X)`**Arguments**

X Numeric data matrix with d columns (vector dimension) and n rows (sample size).

Details

n must be larger than d.

When d=1, `mvShapiro.Test(X)` produces the same results as `shapiro.test(X)`.

Value

A list with class "htest" containing the following components.

statistic	the value of the generalized Shapiro-Wilk statistic for testing multivariate normality.
p.value	the p-value of the test.
method	the character string "Generalized Shapiro-Wilk test for multivariate normality".
data.name	a character string giving the name of the data set.

Author(s)

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References

Villasenor-Alva, J.A. and Gonzalez-Estrada, E. (2009). A generalization of Shapiro-Wilk's test for multivariate normality. *Communications in Statistics: Theory and Methods*, **38** 11,1870-1883.

See Also

[shapiro.test](#)

Examples

```
X <- matrix(rnorm(40),ncol=2) # Generating a two dimensional random sample of size 20
mvShapiro.Test(X)           # Testing multivariate normality on X

#-----
# iris.virginica contains a set of measurements corresponding to
# Iris virginica of the famous iris dataset.

iris.virginica <- as.matrix(iris[iris$Species == "virginica",1:4],ncol=4)
mvShapiro.Test(iris.virginica) # Testing multivariate normality on iris.virginica
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