# MVN package: Multivariate Normality Tests

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#### Abstract

Assessing the assumption of multivariate normality is required by many parametric multivariate statistical methods, such as discriminant analysis, principal component analysis, MANOVA, etc. Here, we present an R package to asses multivariate normality. The MVN package contains three most widely used multivariate normality tests, including Mardia's, Henze-Zirkler's and Royston's multivariate normality tests.

### Contents

1	Preparation of input data	2
2	Multivariate Normality Tests         2.1 Mardia's Multivariate Normality Test	
	2.3 Royston's Multivariate Normality Test	
3	Multivariate Normality Plots	6
	3.1 Q-Q Plot	
	3.2 Perspective and Contour Plots	7
4	Session info	10

### 1 Preparation of input data

MVN package expects a numeric matrix or a data frame that contains minimum two variables. In this vignette, we will work with the *Iris* data set. This data set is a multivariate data set introduced by Ronald A. Fisher (1936) as an application of discriminant analysis [1]. It is also called Anderson's Iris data set because Edgar Anderson collected the data to measure the morphologic variation of Iris flowers of three related species [2]. The data set consists of 50 samples from each of three species of Iris including *setosa*, *virginica* and *versicolor*. For each sample, four variables were measured including the length and the width of the *sepals* and *petals*, in centimeters. We will check the multivariate normality of the *Iris* data set by using three multivariate normality tests, including Mardia's, Royston's and Henze-Zirkler's multivariate normality tests.

First, we can call our data set using data function and display it using head function as follows:

```
data(iris)
head(iris)
     Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##
## 1
                5.1
                                            1.4
                             3.5
                                                          0.2
                                                               setosa
## 2
                4.9
                             3.0
                                            1.4
                                                          0.2
                                                               setosa
                4.7
                             3.2
                                                          0.2
## 3
                                            1.3
                                                               setosa
                4.6
                             3.1
                                            1.5
                                                          0.2
## 4
                                                               setosa
                5.0
                             3.6
## 5
                                            1.4
                                                          0.2
                                                               setosa
## 6
                5.4
                             3.9
                                            1.7
                                                          0.4
                                                               setosa
```

The Iris data is in data.frame format which consists of 5 variables (Sepal.Length, Sepal.Width, Petal.Length, Petal.Width, and Species) and 150 samples.

```
class(iris)
## [1] "data.frame"
dim(iris)
## [1] 150 5
```

For simplicity, we will work with a subset of the *Iris* data with first 50 samples without class label.

```
Iris = iris[1:50, 1:4]
head(Iris)
##
     Sepal.Length Sepal.Width Petal.Length Petal.Width
                5.1
                                            1.4
## 1
                             3.5
                                                          0.2
## 2
                4.9
                             3.0
                                            1.4
                                                          0.2
                4.7
                             3.2
                                            1.3
                                                          0.2
## 3
                                                          0.2
## 4
                4.6
                             3.1
                                            1.5
## 5
                5.0
                             3.6
                                            1.4
                                                          0.2
## 6
                5.4
                             3.9
                                            1.7
                                                          0.4
```

### 2 Multivariate Normality Tests

We will introduce three multivariate normality tests below, including Mardia's, Henze-Zirkler's and Royston's Multivariate Normality Tests.

Before using our multivariate normality tests, we need to load our MVN package as follows:

```
library(MVN)
```

#### 2.1 Mardia's Multivariate Normality Test

Mardia's test is based on multivariate extensions of *skewness* and *kurtosis* measures [3]. Now, we will check the multivariate normality of the *Iris* data using mardiaTest function in the MVN package. This function calculates the Mardia's multivariate skewness and kurtosis coefficients as well as their corresponding statistical tests. For large sample size the multivariate skewness is asymptotically distributed as a chi-square random variable; here it is corrected for small sample size. Likewise, the multivariate kurtosis is distributed as a unit-normal [4–6].

```
result <- mardiaTest(Iris, cov = TRUE, qqplot = FALSE)
result
##
      Mardia's Multivariate Normality Test
##
      data: Iris
##
##
                    : 3.08
      g1p
      chi.skew
               : 25.66
##
##
      p.value.skew : 0.1772
##
##
      g2p
                     : 26.54
##
      z.kurtosis
                    : 1.295
##
      p.value.kurt
                     : 0.1953
##
##
      chi.small.skew : 27.86
##
      p.value.small : 0.1128
##
##
      Result
                     : Data is multivariate normal.
```

#### Here:

```
g1p: Mardia's estimation of multivariate skewness,
chi.skew: Chi-square value of the skewness statistic,
p.value.skew: Significance value of skewness statistic,
g2p: Mardia's estimation of multivariate kurtosis,
z.kurtosis: z value of the kurtosis statistic,
p.value.kurt: Significance value of kurtosis statistic,
chi.small.skew: Chi-square value of the small sample skewness statistic,
p.value.small: Significance value of small sample skewness statistic.
```

As seen above results, both skewness (p = 0.1772) and kurtosis (p = 0.1953) values indicate multivariate normality.

mardiaTest function has an S4 class called mardia. We can use getSlots function in order to get the slots in this S4 class.

```
getSlots("mardia")
##
                                      p.value.skew chi.small.skew
                                                                    p.value.small
               g1p
                          chi.skew
                                         "numeric"
                                                         "numeric"
        "numeric"
                         "numeric"
                                                                          "numeric"
##
##
               g2p
                       z.kurtosis
                                      p.value.kurt
                                                              dname
                                                                         dataframe
##
        "numeric"
                         "numeric"
                                         "numeric"
                                                       "character"
                                                                      "data.frame"
```

To access the informations which are stored in these slots, we can use @ operator as follow:

```
result@p.value.skew

## [1] 0.1772

result@p.value.kurt

## [1] 0.1953
```

#### 2.2 Henze-Zirkler's Multivariate Normality Test

The Henze-Zirkler test is based on a non-negative functional distance that measures the distance between two distribution functions. If the data is multivariate normal, the test statistic is approximately log-normally distributed. It proceeds to calculate the mean, variance and smoothness parameter. Then, mean and variance are log-normalized and the p-value is estimated. We can use hzTest function in the MVN package to calculate the Henze-Zirkler's Multivariate Normality Test [7–11].

Here, HZ is the value of Henze-Zirkler statistic at significance level 0.05 and p-value is a p-value for the Henze-Zirkler's Multivariate Normality Test.

Since the p-value, which optain from the hzTest, lower than 0.05, one can conclude that this multivariate data set deviates from multivariate normality.

hzTest function has an S4 class called hz. We can use getSlots function in order to get the slots in this S4 class.

```
getSlots("hz")

## HZ p.value dname dataframe
## "numeric" "character" "data.frame"
```

To access the informations which are stored in these slots, we can use @ operator as follow:

```
result@HZ

## [1] 0.9488

result@p.value

## [1] 0.04995
```

#### 2.3 Royston's Multivariate Normality Test

Royston's H test uses Shapiro-Wilk's W statistic for multivariate normality. However, if kurtosis of the data greater than 3 then Shapiro-Francia test is used for leptokurtic samples else Shapiro-Wilk test is used for platykurtic samples [10,12–18].

Here, H is the value of Royston's H statistic at significance level 0.05 and p-value is an approximate p-value for the test with respect to equivalent degrees of freedom (edf).

According to the Royston's Multivariate Normality Test, the *Iris* data set does not appear to follow a multivariate normal distribution (p < 0.001).

roystonTest function has an S4 class called royston. We can use getSlots function in order to get the slots in this S4 class.

```
getSlots("hz")

## HZ p.value dname dataframe
## "numeric" "character" "data.frame"
```

To access the informations which are stored in these slots, we can use @ operator as follow:

```
result@H

## [1] 31.52

result@p.value

## [1] 2.188e-06
```

### 3 Multivariate Normality Plots

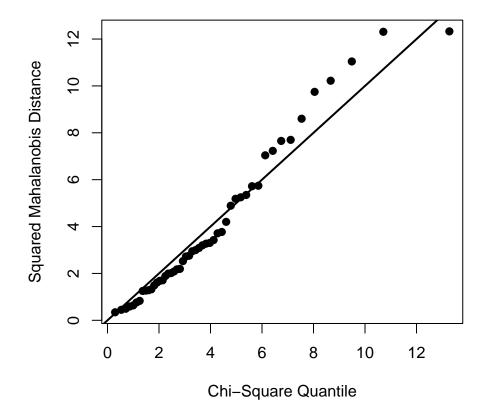
Our MVN package has ability to create three multivariate plots. We can use qqplot = TRUE option in the mardiaTest, hzTest and roystonTest functions to create a chi-suqare Q-Q plot. Furthermore, we can use mvnPlot function in our MVN package to create perspective and contour plots for binary data sets.

### 3.1 Q-Q Plot

We can create a chi-square Q-Q plot for our *Iris* data set to see whether there is any deviation from multivariate normality.

result <- roystonTest(Iris, qqplot = TRUE)</pre>

## Chi-Square Q-Q Plot

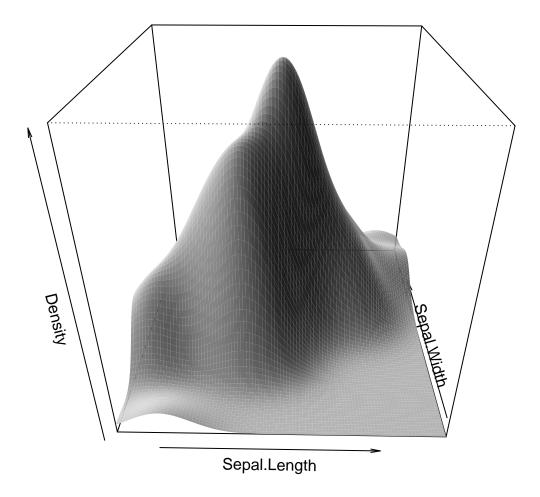


If the data set follows approximately a multivariate normal distribution, the resulting plot should be roughly straight line. As you can see from the chi-square Q-Q plot above, there are some deviations from the straight line and this indicates possible departures from a multivariate normal distribution.

#### 3.2 Perspective and Contour Plots

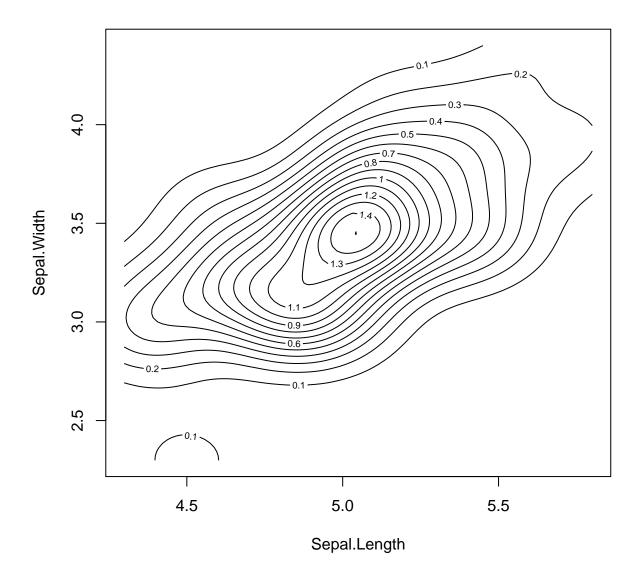
We can use the mvnPlot function in the MVN package to create a perspective plot for a binary data set. In order to get a perspective plot, we should continue with two variables, i.e., bivariate normal distribution. As an example, we subset first 50 rows and *sepal* measures of *Iris* data. Sepal measures of first 50 samples are bivariate normal. We can see that from the perspective plot. Perspective plot produces 3-dimensional bell-shaped graph when data is bivariate normal.

```
Iris = iris[1:50, 1:2]
result = hzTest(Iris)
mvnPlot(result, type = "persp", default = TRUE)
```



Another alternative is to use 2-dimensional contour graphs. We can use the mvnPlot function in the MVN package to create a contour plot for a binary data set. Contour graphs are very useful since it gives information about normality and correlation at the same time.

```
mvnPlot(result, type = "contour", default = TRUE)
```



From contour graph above, we can say that there is a positive correlation among sepal measures of flowers since contour lines lie around main diagonal.

### 4 Session info

```
sessionInfo()
## R version 3.1.0 (2014-04-10)
## Platform: x86_64-apple-darwin13.1.0 (64-bit)
## locale:
## [1] C/tr_TR.UTF-8/tr_TR.UTF-8/C/tr_TR.UTF-8/tr_TR.UTF-8
## attached base packages:
## [1] stats
                graphics grDevices utils
                                              datasets methods
                                                                  base
## other attached packages:
## [1] MVN_3.4
                    MASS 7.3-31
                                 moments_0.13 nortest_1.0-2 knitr_1.5
## loaded via a namespace (and not attached):
## [1] codetools_0.2-8 digest_0.6.4
                                      evaluate_0.5.3 formatR_0.10
## [5] highr_0.3
                   stringr_0.6.2 tools_3.1.0
```

### References

- [1] R. A. Fisher (1936). "The use of multiple measurements in taxonomic problems". Annals of Eugenics 7 (2): 179-188. doi:10.1111/j.1469-1809.1936.tb02137.x
- [2] Edgar Anderson (1936). "The species problem in Iris". Annals of the Missouri Botanical Garden 23 (3): 457-509. JSTOR 2394164.
- [3] Mardia, K. V. (1970). "Measures of multivariate skewness and kurtosis with applications". Biometrika 57 (3): 519-530. doi:10.1093/biomet/57.3.519.
- [4] Mardia, K. V. (1974), Applications of some measures of multivariate skewness and kurtosis for testing normality and robustness studies. Sankhy A, 36:115-128.
- [5] Trujillo-Ortiz, A. and R. Hernandez-Walls. (2003). Mskekur: Mardia's multivariate skewness and kurtosis coefficients and its hypotheses testing. A MATLAB file. URL http://www.mathworks.com/matlabcentral/fileexchange/loadFile.do?objectId=3519
- [6] Stevens, J. (1992), Applied Multivariate Statistics for Social Sciences. 2nd. ed. New-Jersey:Lawrance Erlbaum Associates Publishers. pp. 247-248.
- [7] Trujillo-Ortiz, A., R. Hernandez-Walls, K. Barba-Rojo and L. Cupul-Magana. (2007). HZmvntest:Henze-Zirkler's Multivariate Normality Test. A MATLAB file. URL http://www.mathworks.com/matlabcentral/fileexchange/loadFile.do?objectId=17931
- [8] Henze, N. and Zirkler, B. (1990), A Class of Invariant Consistent Tests for Multivariate Normality. Commun. Statist.-Theor. Meth., 19(10): 35953618.
- [9] Henze, N. and Wagner, Th. (1997), A New Approach to the BHEP tests for multivariate normality. Journal of Multivariate Analysis, 62:1-23.

- [10] Johnson, R.A. and Wichern, D. W. (1992). Applied Multivariate Statistical Analysis. 3rd. ed. New-Jersey:Prentice Hall.
- [11] Mecklin, C. J. and Mundfrom, D. J. (2003), On Using Asymptotic Critical Values in Testing for Multivariate Normality.
- [12] Mecklin, C.J. and Mundfrom, D.J. (2005). A Monte Carlo comparison of the Type I and Type II error rates of tests of multivariate normality. Journal of Statistical Computation and Simulation, 75:93-107.
- [13] Royston, J.P. (1982). An Extension of Shapiro and Wilks W Test for Normality to Large Samples. Applied Statistics, 31(2):115124.
- [14] Royston, J.P. (1983). Some Techniques for Assessing Multivariate Normality Based on the Shapiro-Wilk W. Applied Statistics, 32(2).
- [15] Royston, J.P. (1992). Approximating the Shapiro-Wilk W-Test for non-normality. Statistics and Computing, 2:117-119.121133.
- [16] Royston, J.P. (1995). Remark AS R94: A remark on Algorithm AS 181: The W test for normality. Applied Statistics, 44:547-551.
- [17] Shapiro, S. and Wilk, M. (1965). An analysis of variance test for normality. Biometrika, 52:591611.
- [18] Trujillo-Ortiz, A., R. Hernandez-Walls, K. Barba-Rojo and L. Cupul-Magana. (2007). Roystest:Royston's Multivariate Normality Test. A MATLAB file. URL http://www.mathworks.com/matlabcentral/fileexchange/17811