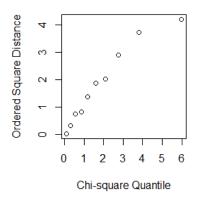
# **Multivariate Analysis**

Homework 2 M052040003 鍾冠毅

```
4.26.
       (a)
       > dii
           diag. D. sq.
            1.8753045
       2
            2.0203262
       3
            2.9009088
       4
            0.7352659
       5
            0.3105192
            0.0176162
       6
       7
            3.7329012
       8
            0.8165401
       9
            1.3753379
            4.2152799
       (b)
       > chisq.2.0.5 <- dii[dii <= qchisq(0.5, 2)]
> length(chisq.2.0.5)/length(dii)
       [1] 0.5
       (c)
       > order.d
         [1] 0.0176162 0.3105192 0.7352659 0.8165401 1.3753379 1.8753045 2.0203262
```

### **Chi-square Plot**

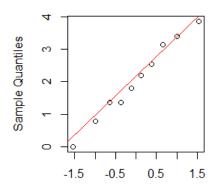
[8] 2.9009088 3.7329012 4.2152799



(d) 由(b)(c)結果可知,兩數據服從雙變數常態分配。

4.30. (a)  $\hat{\lambda}_1 = 0.3708906$ 

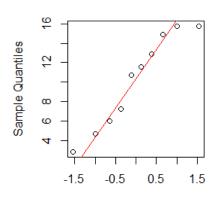
### **Normal Q-Q Plot**



Theoretical Quantiles

(b)  $\hat{\lambda}_2 = 0.9361967$ 

# Normal Q-Q Plot



Theoretical Quantiles

(c)  $(\hat{\lambda}_1, \hat{\lambda}_2) = (1.2732157, 0.0310405)$ 

若個別計算最佳的 $\hat{\lambda}$ ,則會使兩變數個別較趨近常態分佈。若同時計 $\hat{\lambda}_1$ 與 $\hat{\lambda}_2$ ,則會找到最佳的 $\hat{\lambda}$ ,使得兩變數較趨近於雙變量常態分佈。

### > df.norm

Indep Supp Benev Conform Leader df.ks.p 0.009751414 0.0002120256 0.03796667 0.005667049 5.349873e-05 df.lambda 0.523772578 1.3962618614 1.26187573 1.039266313 3.815471e-01 df.ks.p.n 0.125018888 0.0014273401 0.11409417 0.008039632 7.012327e-02

df.ks.p : 轉換前的 Kolmogorov-Smirnov 檢定之 p-value

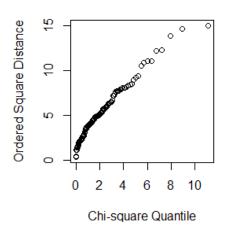
df.lamda :轉換所使用的最佳λ

df.ks.p :轉換後的 Kolmogorov-Smirnov 檢定之 p-value

(a) 由上表可知所有的 p-value 皆小於 0.05, 故拒絕常態之假設。

(b) 雖然在 Chi-square 圖中各點分布接近一斜直線,但是由(a)可知個變數之邊際分佈皆不符從常態分佈,故此資料不符從多變量常態分佈。

## **Chi-square Plot**



(c) 由上表可知,各變數經轉換後重新檢定之 p-value 皆有成長的趨勢,甚至大於 0.05,即服從常態分佈,如:Indep、Benev、Leader 等。

### Appendix

```
ylab = "Ordered Square Distance", main =
### 4.26 ###
                                                        "Chi-square Plot")
# (a) #
x1 <- c(1, 2, 3, 3, 4, 5, 6, 8, 9, 11)
                                                        ### 4.30. ###
x2 <- c(18.95, 19.00, 17.95, 15.54, 14.00,
                                                        install.packages("MASS")
     12.95, 8.94, 7.49, 6.00, 3.99)
                                                        library(MASS)
                                                        library(car)
df <- data.frame(cbind(x1, x2))
S \leftarrow cov(df)
                                                        # (a) #
x1.bar <- mean(x1)
                                                        lambda1 <- powerTransform(x1)</pre>
x2.bar <- mean(x2)
                                                        l1 <- lambda1$lambda</pre>
A \leftarrow t(cbind((x1 - x1.bar), (x2 - x2.bar)))
                                                        x1.n <- (x1^|1-1)/|1
D.sq <- t(A)%*%solve(S)%*%A
                                                        qqnorm(x1.n)
dii <- diag(D.sq)
                                                        qqline(x1.n, col="red")
                                                        # (b) #
# (b) #
                                                        lambda2 <- powerTransform(x2)</pre>
chisq.2.0.5 <- dii[dii <= qchisq(0.5, 2)]
                                                        I2 <- lambda2$lambda
length(chisq.2.0.5)/length(dii)
                                                        x2.n <- (x2^|2-1)/|2
                                                        qqnorm(x2.n)
# (c) #
                                                        qqline(x2.n, col="red")
order.d <- dii[order(dii)]
chi.10 <- sapply(1:10,
                                                        # (c) #
          function(j)
                                                        lambda12 <- powerTransform(cbind(x1,x2))</pre>
           qchisq((10-j+0.5)/10,2, lower.tail
= F))
                                                        l1.c <- lambda12$lambda[1]</pre>
plot(order.d ~ chi.10, xlab = "Chi-square
                                                        l2.c <- lambda12$lambda[2]</pre>
Quantile",
                                                        x1.c <- (x1^|1.c-1)/|1.c
```

```
x2.c <- (x1^12.c-1)/12.c
                                                         ylab = "Ordered Square Distance", main =
                                                      "Chi-square Plot")
### 4.39 ###
                                                      # (c) #
# (a) #
                                                      library(car)
#install.packages("nortest")
                                                      sapply(1:5, function(x)
library(nortest)
                                                      powerTransform(df[,x])$lambda)
df <- read.csv("4.39.csv", header = F, sep =
",")
names(df) <- c("Indep", "Supp", "Benev",
"Conform",
                                                      df.lambda <- sapply(1:5, function(x)
                                                      powerTransform(df[, x])$lambda)
        "Leader", "Gender", "Socio")
                                                      df.n <-
df.ks.p <- sapply(1:5, FUN = function(x)
lillie.test(df[,x])$p.value)
                                                      apply(df, MARGIN = 2,
df.sw.p <- sapply(1:5, FUN = function(x))
                                                          FUN = function(x) {
shapiro.test(df[,x])$p.value)
                                                           lam <- powerTransform(x)$lambda;</pre>
                                                           (x^lam-1)/lam})
# (b) #
df <- df[, 1:5]
S \leftarrow cov(df)
                                                      df.ks.p.n <- sapply(1:5, function(x)
A <- t(apply(df, MARGIN = 2, FUN =
                                                      lillie.test(df.n[, x])$p.value)
function(x) x-mean(x)))
D.sq <- t(A)%*%solve(S)%*%A
                                                      df.norm <- t(cbind(df.ks.p, df.lambda,
dii <- diag(D.sq)
                                                      df.ks.p.n))
order.d <- dii[order(dii)]
                                                      colnames(df.norm) <- names(df)
chi.130 <- sapply(1:130,
          FUN = function(j)
           qchisq((length(dii)-
j+0.5)/length(dii),2, lower.tail = F))
plot(order.d~chi.130, xlab = "Chi-square
Quantile",
```