



113-2 生物統計學一

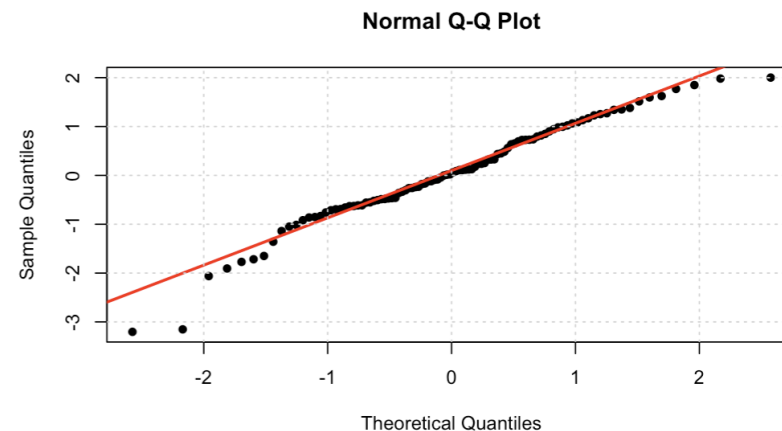
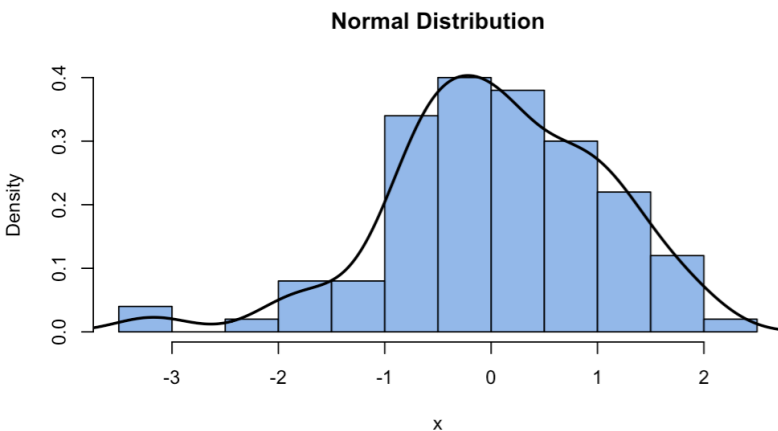
# 無母數分析

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# 常態檢定

1. 圖形檢驗法(graphical test): 密度曲線(density plot)、QQ plot (Quantile–Quantile Plot), ...
2. 統計假設檢定: Shapiro–Wilk test, Kolmogorov–Smirnov test, Anderson–Darling test, ...



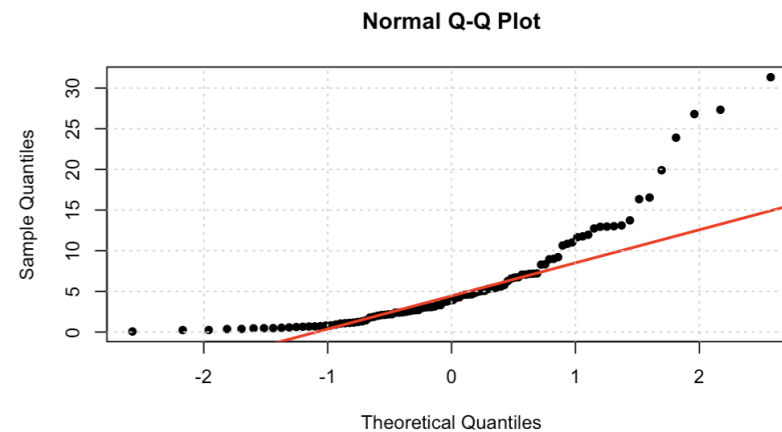
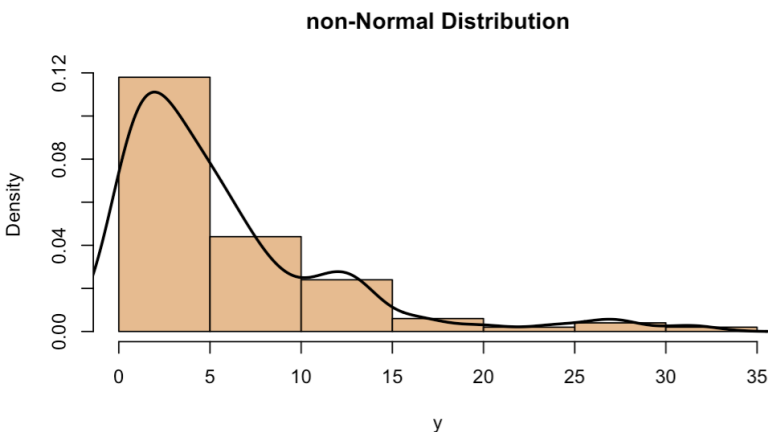
```
> shapiro.test(x)
```

Shapiro-Wilk normality test

```
data: x  
W = 0.97194, p-value = 0.03117
```

```
> ks.test(x, "pnorm", mean(x), sd(x))
```

Asymptotic one-sample Kolmogorov-Smirnov test



```
data: x  
D = 0.065759, p-value = 0.7802  
alternative hypothesis: two-sided
```

# Signed Test

- 適用情境：檢定單一樣本的母體中位數是否等於某定值  
(台大學生每週通勤時數的中位數是否為 5 小時?)

資料裡大於5小時的個數  
> `binom.test(29, 50, p = 0.5)`

Exact binomial test

```
data: 29 and 50
number of successes = 29, number of trials = 50, p-value = 0.3222
alternative hypothesis: true probability of success is not equal to 0.5
95 percent confidence interval:
 0.4320604 0.7181178
sample estimates:
probability of success
              0.58
```



$H_0$ 為真下，檢定統計量服從Bin(n, p)

其中  $p = P(X > M_0) = 0.5$

# Signed Test

- 適用情境：檢定**單一樣本**的母體中位數是否等於某定值  
(台大學生每週通勤時數的中位數是否為 5 小時?)
  - > `install.packages("BSDA")` (使用非R內建函數需要另外下載套件)
  - > `library("BSDA")` (使用前須先將下載套件匯入)
  - > `SIGN.test(studata1$Hours_Commute, md = 5)`

## One-sample Sign-Test

```
data: studata1$Hours_Commute
s = 29, p-value = 0.3222
alternative hypothesis: true median is not equal to 5
95 percent confidence interval:
 3 9
sample estimates:
median of x
 6
```

# Wilcoxon Signed Rank Test

- 適用情境：檢定配對樣本的母體中位數是否等於某定值  
(台大學生本學期 (credit\_current) 與上學期 (credit\_last) 的修課學分數是否相同？)

```
> wilcox.test(studata1$Credit_Last, studata1$Credit_Current, paired = TRUE)
```

```
Wilcoxon signed rank test with continuity correction
```

```
data: studata1$Credit_Last and studata1$Credit_Current
```

```
V = 522, p-value = 0.3788
```

```
alternative hypothesis: true location shift is not equal to 0
```

Warning messages:

```
1: In wilcox.test.default(studata1$Credit_Last, studata1$Credit_Current,  :  
   cannot compute exact p-value with ties
```

```
2: In wilcox.test.default(studata1$Credit_Last, studata1$Credit_Current,  :  
   cannot compute exact p-value with zeroes
```

- 有太多差異為0或太多資料排名相同(tie)的情況，因此以近似的分佈計算p-value

# Wilcoxon rank sum test / Mann-Whitney U test

- 適用情境：檢定兩獨立樣本的母體中位數是否相等  
(台大公衛系與非公衛系學生 (department), 每周通勤時數的中位數是否相同? )

```
> wilcox.test(studata1$Hours_Commute ~ studata1$Department, paired = F)
```

Wilcoxon rank sum test with continuity correction

```
data: studata1$Hours_Commute by studata1$Department
```

```
W = 137, p-value = 0.7148
```

```
alternative hypothesis: true location shift is not equal to 0
```

Warning message:

```
In wilcox.test.default(x = DATA[[1L]], y = DATA[[2L]], ...) :  
cannot compute exact p-value with ties
```

# Kruskal-Wallis Test

- 適用情境：檢定三組或以上獨立樣本的母體中位數是否相同  
(若以目前居住縣市 (county) 來分組，請問住在不同縣市的同學，每週通勤時數的中位數是否相同？)

```
> kruskal.test(studata1$Hours_Commute ~ studata1$County)
```

```
Kruskal-Wallis rank sum test
```

```
data: studata1$Hours_Commute by studata1$County  
Kruskal-Wallis chi-squared = 10.529, df = 3, p-value = 0.01457
```

- 事後檢定

```
> pairwise.wilcox.test(studata1$Hours_Commute, studata1$County, p.adjust.method = "bonf")
```

```
Pairwise comparisons using Wilcoxon rank sum test with continuity correction
```

```
data: studata1$Hours_Commute and studata1$County
```

	1	2	3
2	0.026	-	-
3	1.000	1.000	-
4	1.000	0.057	1.000