

ZeaMays Data

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- ZeaMays Data 는 “HistData” package 에 있음.

```
install.packages("HistData", repos="http://cran.rstudio.com")
```

```
##  
## The downloaded binary packages are in  
## /var/folders/_h/tglth9bd4h98rjjb5vy9gn3m0000gn/T//RtmpVR9Jf9/downloaded_packages
```

```
library(HistData)  
str(ZeaMays)
```

```
## 'data.frame':    15 obs. of  5 variables:  
## $ pair : int  1 2 3 4 5 6 7 8 9 10 ...  
## $ pot : Factor w/ 4 levels "1","2","3","4": 1 1 1 2 2 2 3 3 3 3 ...  
## $ cross: num  23.5 12 21 22 19.1 ...  
## $ self : num  17.4 20.4 20 20 18.4 ...  
## $ diff : num  6.12 -8.38 1 2 0.75 ...
```

ZeaMays

```
##      pair pot  cross  self  diff  
## 1      1   1 23.500 17.375  6.125  
## 2      2   1 12.000 20.375 -8.375  
## 3      3   1 21.000 20.000  1.000  
## 4      4   2 22.000 20.000  2.000  
## 5      5   2 19.125 18.375  0.750  
## 6      6   2 21.500 18.625  2.875  
## 7      7   3 22.125 18.625  3.500  
## 8      8   3 20.375 15.250  5.125  
## 9      9   3 18.250 16.500  1.750  
## 10     10   3 21.625 18.000  3.625  
## 11     11   3 23.250 16.250  7.000  
## 12     12   4 21.000 18.000  3.000  
## 13     13   4 22.125 12.750  9.375  
## 14     14   4 23.000 15.500  7.500  
## 15     15   4 12.000 18.000 -6.000
```

```
attach(ZeaMays)
```

- Competition 현상 확인.

```

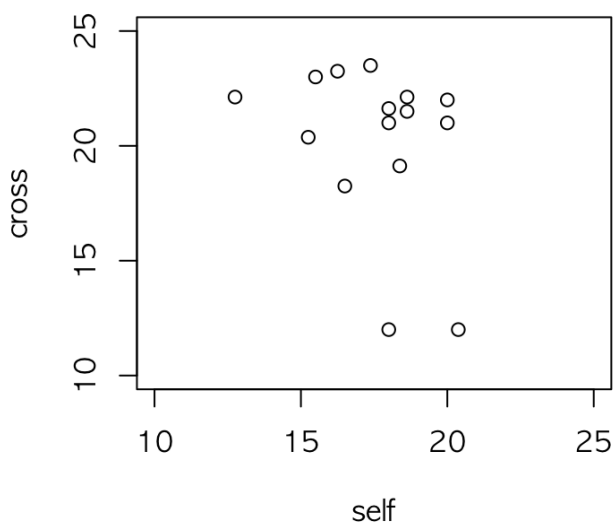
par(mfrow=c(1,2))
plot(x=self, y=cross, xlim=c(10, 25), ylim=c(10, 25))
title(main="Cross and Self Fertilisation")
plot(x=self, y=cross, pch=16, xlim=c(10, 25), ylim=c(10, 25))
title(main="Cross and Self Fertilisation")
abline(lsfit(x=self, y=cross)$coef, col="red")
cor(self, cross)

```

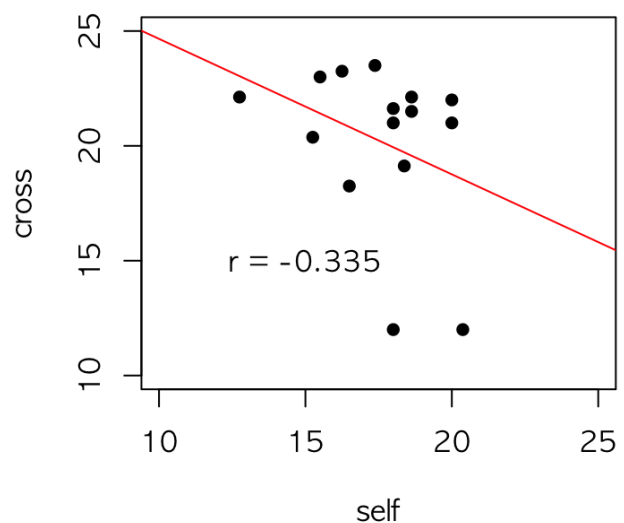
```
## [1] -0.3347553
```

```
text(x=15, y=15, labels=paste("r =", round(cor(cross, self), digits=3)))
```

Cross and Self Fertilisation



Cross and Self Fertilisation



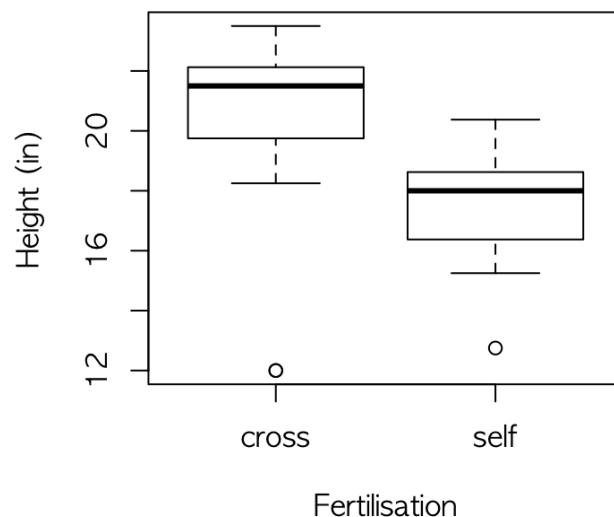
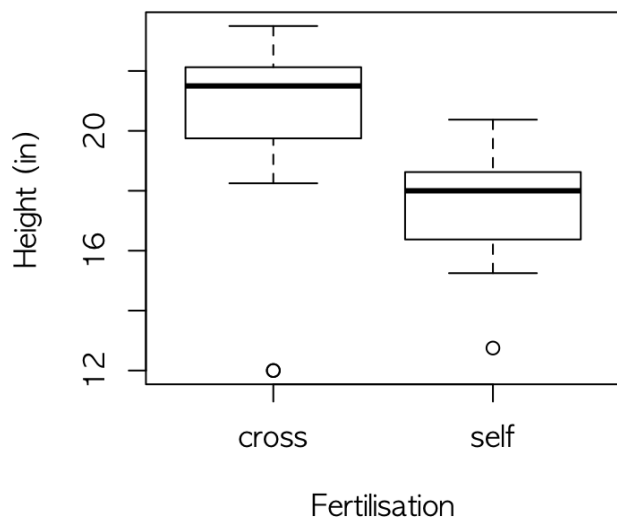
```
par(mfrow=c(1,1))
```

- boxplot 으로 비교하면 이 현상을 파악하기 어려움. 같은 결과를 갖는 두 가지 boxplot 작성 코드 비교.

```

par(mfrow=c(1,2))
boxplot(ZeaMays[, c("cross", "self")], ylab="Height (in)", xlab="Fertilisation")
boxplot(cross, self, names=c("cross", "self"), ylab="Height (in)", xlab="Fertilisation")

```



- 쌍으로 키우고 있으므로 성장의 차이는 paired one-sample t-test

```
t.test(x=cross, y=self, paired=T)
```

```
##
## Paired t-test
##
## data: cross and self
## t = 2.148, df = 14, p-value = 0.0497
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  0.003899165 5.229434169
## sample estimates:
## mean of the differences
##                2.616667
```

- 또는 둘 사이의 차이인 diff 대하여 one-sample t-test 를 수행하여도 동일한 결과. 이 때는 모든 매개변수는 디폴트 값으로 충족됨에 유의.

```
t.test(diff)
```

```
##
## One Sample t-test
##
## data: diff
## t = 2.148, df = 14, p-value = 0.0497
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
##  0.003899165 5.229434169
## sample estimates:
## mean of x
## 2.616667
```

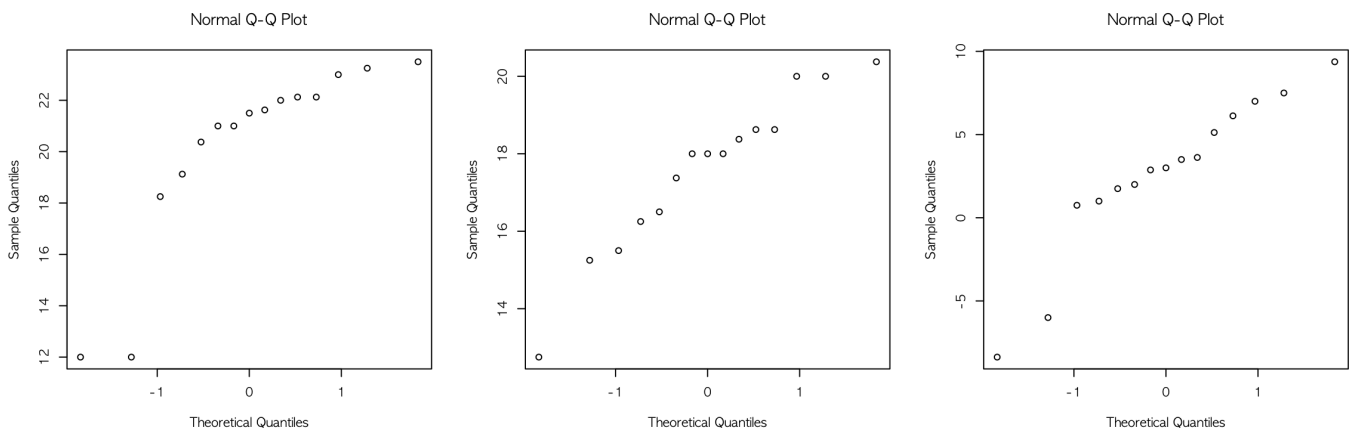
- 정규성에 대한 가정은 `ad.test`로 수행

```
library(nortest)
lapply(list(cross=cross, self=self, diff=diff), ad.test)
```

```
## $cross
##
##  Anderson-Darling normality test
##
## data:  X[[1L]]
## A = 1.4755, p-value = 0.0005057
##
##
## $self
##
##  Anderson-Darling normality test
##
## data:  X[[2L]]
## A = 0.3547, p-value = 0.4122
##
##
## $diff
##
##  Anderson-Darling normality test
##
## data:  X[[3L]]
## A = 0.6175, p-value = 0.08758
```

- `cross` 자료에 대한 정규성은 매우 의심되는 수준. `qqnorm`으로 파악.

```
par(mfrow=c(1,3))
lapply(list(cross=cross, self=self, diff=diff), qqnorm)
```



```
## $cross
## $cross$x
## [1] 1.8339146 -1.8339146 -0.3406948 0.3406948 -0.7279133 0.0000000
## [7] 0.5244005 -0.5244005 -0.9674216 0.1678940 1.2815516 -0.1678940
## [13] 0.7279133 0.9674216 -1.2815516
##
## $cross$y
## [1] 23.500 12.000 21.000 22.000 19.125 21.500 22.125 20.375 18.250 21.625
## [11] 23.250 21.000 22.125 23.000 12.000
##
##
## $self
## $self$x
## [1] -0.3406948 1.8339146 0.9674216 1.2815516 0.3406948 0.5244005
## [7] 0.7279133 -1.2815516 -0.5244005 -0.1678940 -0.7279133 0.0000000
## [13] -1.8339146 -0.9674216 0.1678940
##
## $self$y
## [1] 17.375 20.375 20.000 20.000 18.375 18.625 18.625 15.250 16.500 18.000
## [11] 16.250 18.000 12.750 15.500 18.000
##
##
## $diff
## $diff$x
## [1] 0.7279133 -1.8339146 -0.7279133 -0.3406948 -0.9674216 -0.1678940
## [7] 0.1678940 0.5244005 -0.5244005 0.3406948 0.9674216 0.0000000
## [13] 1.8339146 1.2815516 -1.2815516
##
## $diff$y
## [1] 6.125 -8.375 1.000 2.000 0.750 2.875 3.500 5.125 1.750 3.625
## [11] 7.000 3.000 9.375 7.500 -6.000
```

```
par(mfrow=c(1,1))
```

- t.test의 대안으로 wilcox.test 수행

```
wilcox.test(cross, self, paired=T)
```

```
##
## Wilcoxon signed rank test
##
## data: cross and self
## V = 96, p-value = 0.04126
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(diff)
```

```
##  
## Wilcoxon signed rank test  
##  
## data: diff  
## V = 96, p-value = 0.04126  
## alternative hypothesis: true location is not equal to 0
```

- cross와 self 가 성장률이 동일하다는 가설은 기각됨.