## STAT 8010 R Lab 10

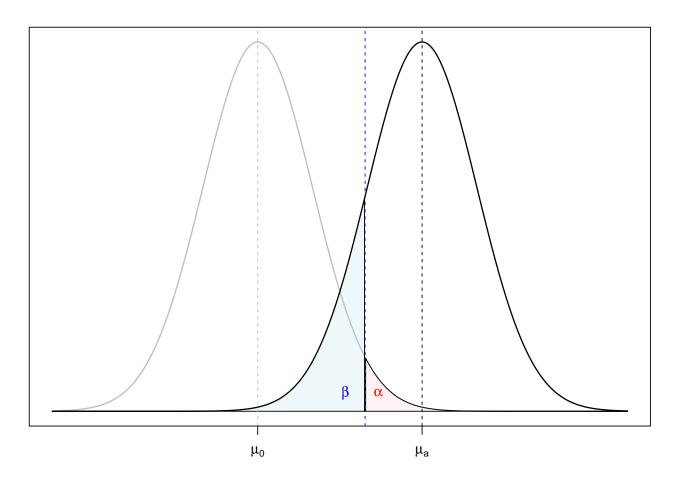
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### Type I and Type II Errors

```
library(scales)
curve(dnorm(x), from = -3.75, to = 6.75, n = 1001, lwd = 1.5, col = "gray",
      xlab = "",
      ylab = "", las = 1,
      xaxt = "n", yaxt = "n")
abline(v = 1.96, lty = 2, col = "blue")
x_{grid} \leftarrow seq(-3.75, 6.75, 0.01)
y_grid <- dnorm(x_grid)</pre>
polygon(c(x_grid[x_grid > 1.96], rev(x_grid[x_grid > 1.96])),
        c(y_grid[x_grid > 1.96], rep(0, length(y_grid[x_grid > 1.96]))), col = alpha("pink", 0.2))
curve(dnorm(x, mean = 3), from = -3.75, to = 6.75, n = 1001, lwd = 1.5, add = T,
      xlab = "",
      ylab = "", las = 1)
x_{grid} \leftarrow seq(-3.75, 6.75, 0.01)
y_grid <- dnorm(x_grid, mean = 3)</pre>
polygon(c(x_grid[x_grid < 1.96], rev(x_grid[x_grid < 1.96])),</pre>
        c(y_grid[x_grid < 1.96], rep(0, length(y_grid[x_grid < 1.96]))), col = alpha("lightblue", 0.2))</pre>
abline(v = 0, lty = 2, col = "gray")
abline(v = 3, lty = 2)
axis(1, at = 0, labels = expression(mu[0]))
axis(1, at = 3, labels = expression(mu[a]))
text(2.2, 0.02, expression(alpha), col = "red")
text(1.6, 0.02, expression(beta), col = "blue")
```



## Power analysis

```
library(asbio)
## Warning: package 'asbio' was built under R version 3.6.2
## Loading required package: tcltk
power.z.test(sigma = 10, n = 25, power = NULL, alpha = 0.05,
            effect = 4, test = c("one.tail"))
## $sigma
## [1] 10
##
## $n
## [1] 25
##
## $power
## [1] 0.63876
##
## $alpha
## [1] 0.05
## $effect
## [1] 4
##
## $test
```

```
## [1] "one.tail"
power.z.test(sigma = 10, n = NULL, power = 0.8, alpha = 0.05,
             effect = 4, test = c("one.tail"))
## $sigma
## [1] 10
##
## $n
## [1] 38.64098
##
## $power
## [1] 0.8
##
## $alpha
## [1] 0.05
##
## $effect
## [1] 4
##
## $test
## [1] "one.tail"
power.t.test(n = 25, delta = 4, sd = 10, sig.level = 0.05,
             power = NULL, type = "one.sample", alternative = "one.sided")
##
##
        One-sample t test power calculation
##
##
                 n = 25
##
             delta = 4
##
                sd = 10
##
         sig.level = 0.05
##
            power = 0.617259
       alternative = one.sided
power.t.test(n = NULL, delta = 4, sd = 10, sig.level = 0.05,
             power = 0.8, type = "one.sample", alternative = "one.sided")
##
##
        One-sample t test power calculation
##
##
                 n = 40.02908
##
             delta = 4
##
                sd = 10
##
         sig.level = 0.05
##
             power = 0.8
##
       alternative = one.sided
Inference for two population means
```

#### Tapeworm example

```
treatment <- c(18, 43, 28, 50, 16, 32, 13, 35, 38, 33, 6, 7)
control <- c(40, 54, 26, 63, 21, 37, 39, 23, 48, 58, 28, 39)
dat <- data.frame(cbind(treatment, control))
```

```
summary(dat)
                      control
##
     treatment
                          :21.00
##
  Min. : 6.00 Min.
  1st Qu.:15.25
                   1st Qu.:27.50
## Median :30.00
                   Median :39.00
## Mean :26.58
                   Mean :39.67
## 3rd Qu.:35.75
                   3rd Qu.:49.50
          :50.00
## Max.
                   Max.
                          :63.00
boxplot(dat, boxwex = 0.3, las = 1)
60 -
50
40
30
20
10 -
                      treatment
                                                              control
apply(dat, 2, mean)
## treatment
              control
## 26.58333 39.66667
apply(dat, 2, sd)
## treatment
              control
## 14.36193 13.85859
var.test(treatment, control)
##
## F test to compare two variances
##
## data: treatment and control
## F = 1.074, num df = 11, denom df = 11, p-value = 0.9079
## alternative hypothesis: true ratio of variances is not equal to 1
```

```
## 0.3091686 3.7306092
## sample estimates:
## ratio of variances
             1.073959
# Assuming $\sigma_{1}=\sigma_{2}$
t.test(treatment, control, var.equal = T)
##
##
   Two Sample t-test
##
## data: treatment and control
## t = -2.2709, df = 22, p-value = 0.03329
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -25.031761 -1.134906
## sample estimates:
## mean of x mean of y
## 26.58333 39.66667
# Assuming $\sigma_{1}\neq sigma_{2}$
t.test(treatment, control, var.equal = F)
##
  Welch Two Sample t-test
##
## data: treatment and control
## t = -2.2709, df = 21.972, p-value = 0.03331
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -25.032642 -1.134025
## sample estimates:
## mean of x mean of y
## 26.58333 39.66667
# Left-tailed test
t.test(treatment, control, alternative = "less")
##
## Welch Two Sample t-test
## data: treatment and control
## t = -2.2709, df = 21.972, p-value = 0.01665
## alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
        -Inf -3.189613
##
## sample estimates:
## mean of x mean of y
## 26.58333 39.66667
Two sample t test with only sample statistics
t.test.from.summary.data <- function(mean1, sd1, n1, mean2, sd2, n2, ...) {
   data1 \leftarrow scale(1:n1)*sd1 + mean1
   data2 <- scale(1:n2)*sd2 + mean2</pre>
```

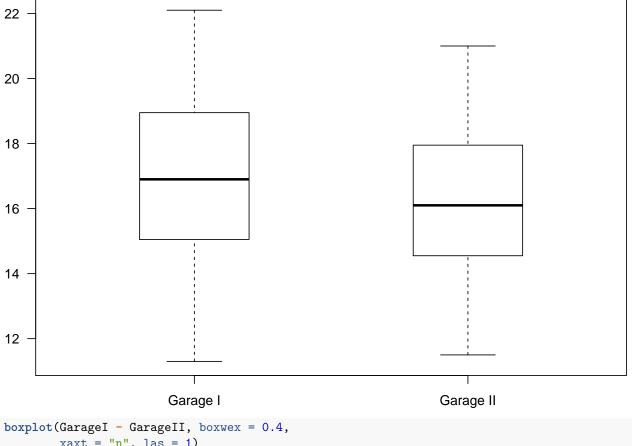
## 95 percent confidence interval:

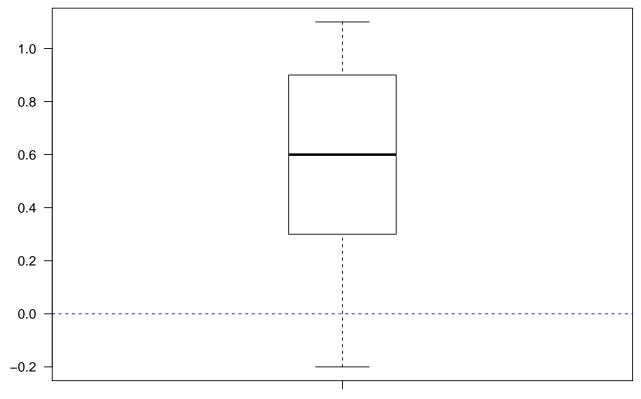
```
t.test(data1, data2, ...)
}
t.test.from.summary.data(12.5, 7.63, 10, 27.5, 15.3, 10)
##
## Welch Two Sample t-test
##
## data: data1 and data2
## t = -2.7744, df = 13.216, p-value = 0.01558
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -26.660768 -3.339232
## sample estimates:
## mean of x mean of y
##
        12.5
                   27.5
## Check
df = ((4.3^2)/37 + (2.2^2)/31)^2 / (((4.3^2)/37)^2 / 36 + ((2.2^2)/31)^2 / 30)
se \leftarrow sqrt(4.3<sup>2</sup> / 37 + 2.2<sup>2</sup> / 31)
tstat \leftarrow (19.45 - 18.2) / se
Pvalue \leftarrow 2 * (1 - pt(1.5435, df))
```

#### Paired T-Test

```
repair \leftarrow c(17.6, 17.3, 20.2, 19.1,
             19.5, 18.4, 11.3, 11.5,
            13.0, 12.7, 16.3, 15.8,
            15.3, 14.9, 16.2, 15.3,
             12.2, 12.0, 14.8, 14.2,
             21.3, 21.0, 22.1, 21.0,
             16.9, 16.1, 17.6, 16.7,
             18.4, 17.5)
GarageI <- repair[seq(1, 29, 2)]</pre>
GarageII <- repair[seq(2, 30, 2)]</pre>
dat <- cbind(GarageI, GarageII)</pre>
apply(dat, 2, mean)
## GarageI GarageII
## 16.84667 16.23333
library(scales)
plot(GarageI, GarageII,
     pch = 16, col = alpha("blue", 0.5), las = 1,
     xlim = c(11, 23),
     ylim = c(11, 23))
abline(0, 1, col = "gray", lwd = 1.5)
```

```
22 - 20 - 18 - 16 - 14 - 12 - 12 14 16 18 20 22 GarageI , GarageII ,
```





Garage I - Garage II

```
t.test(GarageI, GarageII, alternative = c("greater"), paired = T)
```