Augusta University: STAT 7630

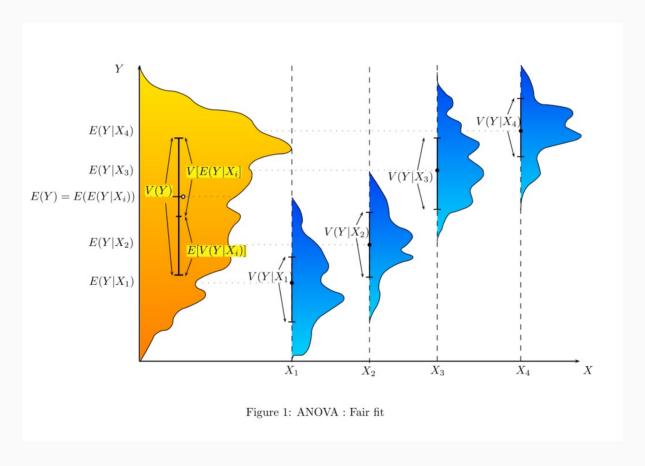
Applied Linear Models

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Lecture 2

- Review of Confidence Intervals
- One-way Analysis of Variance (ANOVA)



Let
$$X_1,\ldots,X_n\stackrel{iid}{\sim}\mathcal{N}(\mu_X,\sigma^2)$$
 and $Y_1,\ldots,Y_m\stackrel{iid}{\sim}\mathcal{N}(\mu_Y,\sigma^2)$.

We wish to test

$$H_0: \mu_X = \mu_Y$$

 $H_1: \mu_X \neq \mu_Y$

If σ^2 is **known**, we can use the two-sample z-test, which has test statistic

$$T(X) = rac{\overline{X} - \overline{Y}}{\sigma \sqrt{rac{1}{n} + rac{1}{m}}} \stackrel{H_0}{\sim} \mathcal{N}(0,1).$$

Let
$$X_1,\ldots,X_n\stackrel{iid}{\sim}\mathcal{N}(\mu_X,\sigma^2)$$
 and $Y_1,\ldots,Y_m\stackrel{iid}{\sim}\mathcal{N}(\mu_Y,\sigma^2)$.

We wish to test

$$H_0: \mu_X = \mu_Y \ H_1: \mu_X
eq \mu_Y$$

If σ^2 is **unknown**, we instead use the two-sample t-test, with pooled variance estimator s_p calculated as

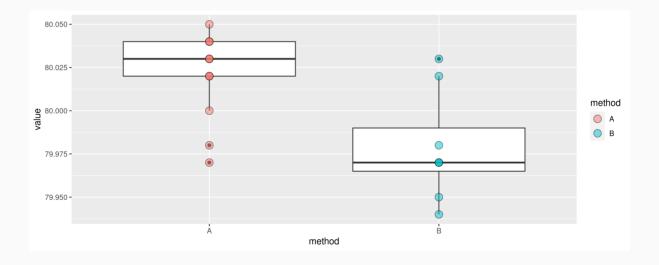
$$s_p^2 = rac{(n-1)s_X^2 + (m-1)s_Y^2}{m+n-2}.$$

The test statistic is

$$T(X) = rac{\overline{X} - \overline{Y}}{s_p \sqrt{rac{1}{n} + rac{1}{m}}} \stackrel{H_0}{\sim} t(m+n-2).$$

Example A (Rice, p423)

Two methods, A and B, were used in a determination of the latent heat of fusion of ice (Matrella 1963). The investigators wanted to find out by how much the methods differed. The data give the change in total heat from ice at $-0.72^{\circ}\mathrm{C}$ to water $0^{\circ}\mathrm{C}$ in calories per gram of mass:



```
A \leftarrow c(79.98, 80.04, 80.02, 80.04, 80.03, 80.03, 80.04, 79.97, 80.05, 80.03, 80.02, 80.00,
B \leftarrow c(80.02, 79.94, 79.98, 79.97, 79.97, 80.03, 79.95, 79.97)
n \leftarrow length(A)
m \leftarrow length(B)
dat \leftarrow data.frame(value = c(A, B), method = c(rep("A", n), rep("B", m)))
dat
      value method
##
      79.98
## 1
## 2
      80.04
                  Α
## 3
      80.02
                  Α
      80.04
## 4
                   Α
      80.03
## 5
                   Α
## 6
      80.03
                   Α
## 7 80.04
                   Α
      79.97
## 8
                   Α
      80.05
## 9
                   Α
## 10 80.03
                   Α
## 11 80.02
                   Α
## 12 80.00
                   Α
## 13 80.02
                   Α
                                                                                             6 / 23
## 14 80.02
                   В
```

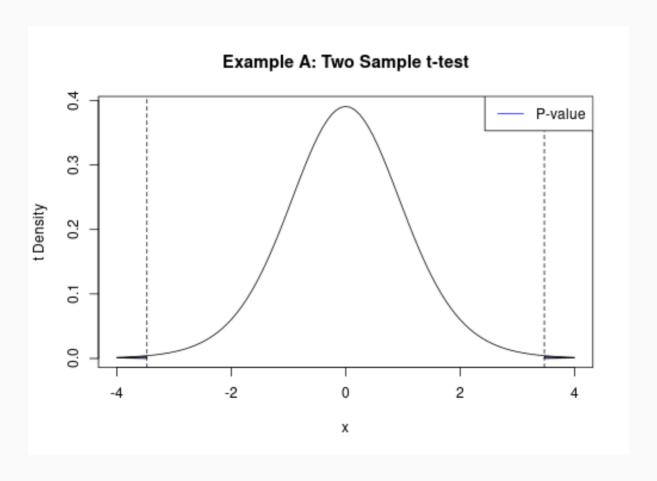
```
library(dplyr)
dat_summary ← dat %>% group_by(method) %>%
  summarize(mean = mean(value), sd = sd(value))
```

method	mean	sd
А	80.02077	0.0239658
В	79.97875	0.0313676

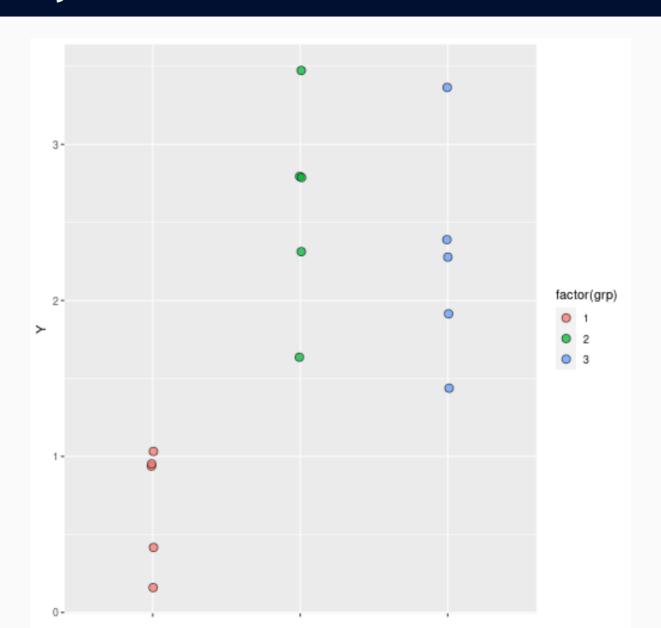
Example A

Using the built-in R function:

```
##
## Two Sample t-test
##
## data: A and B
## t = 3.4722, df = 19, p-value = 0.002551
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 0.01669058 0.06734788
## sample estimates:
## mean of x mean of y
## 80.02077 79.97875
```

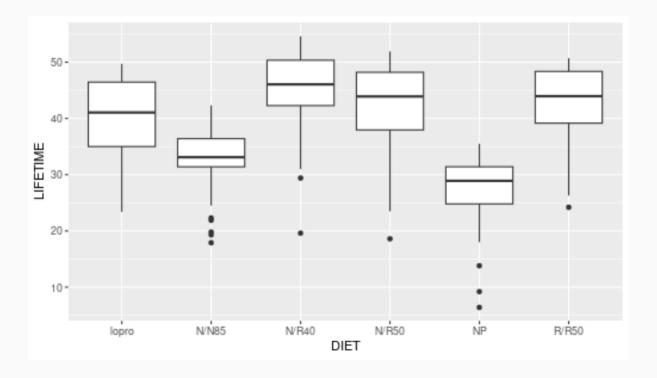


```
I \leftarrow 3
J \leftarrow 5
mu \leftarrow 1
alpha1 \leftarrow -0.5
alpha2 \leftarrow 2
alpha3 \leftarrow 1
sigma \leftarrow 0.75
set.seed(12345)
Y1 \leftarrow mu + alpha1 + rnorm(J, 0, sigma)
Y2 \leftarrow mu + alpha2 + rnorm(J, 0, sigma)
Y3 \leftarrow mu + alpha3 + rnorm(J, 0, sigma)
```



Diet and Longevity Study (12.5.33, Rice)

It is hypothesized that animals on restricted calorie diets have longer lifespans on average. A study by Weindruch et al. examined the effects of 6 different types of diets on lifespans in female mice.



Diet and Longevity Study (12.5.33, Rice)

We wish to test whether there is a significant difference in the average lifespans across the treatment groups.

 H_0 : All group mean lifespans are equal.

 H_1 : At least one group has a different mean lifespan.

Diet and Longevity Study (12.5.33, Rice)

DIET	mean	sd
N/N85	32.69123	5.125297
N/R40	45.11667	6.703406
N/R50	42.29718	7.768195
NP	27.40204	6.133701
R/R50	42.88571	6.683152
lopro	39.68571	6.991695

Diet and Longevity Study (12.5.33, Rice)

```
SSTOT ← sum((dat$LIFETIME - mean(dat$LIFETIME))^2)
SSTOT
## [1] 28031.36
SSW ← 0
for (subj in 1:n) {
  SSW ← SSW + (dat$LIFETIME[subj] - dat_summary$mean[which(dat_summary$DIET = dat$DIET[subj])
SSW
## [1] 15297.42
SSB ← SSTOT - SSW
SSB
## [1] 12733.94
```

Diet and Longevity Study (12.5.33, Rice)

```
MSB ← SSB / (I - 1)
MSB

## [1] 2546.788

MSE ← SSW / (n - I)
MSE

## [1] 44.59888
```

Diet and Longevity Study (12.5.33, Rice)

F statistic

```
test_stat ← MSB / MSE
test_stat
## [1] 57.10431
```

P-value

```
P_{val} \leftarrow pf(test_{stat}, df1 = I - 1, n - I, lower.tail = FALSE)
P_{val}
## [1] 4.111744e-43
```

Diet and Longevity Study (12.5.33, Rice)

We wish to test whether there is a significant difference in the average lifespans across the treatment groups.

```
fit ← lm(LIFETIME ~ DIET, data = dat)
anova(fit)

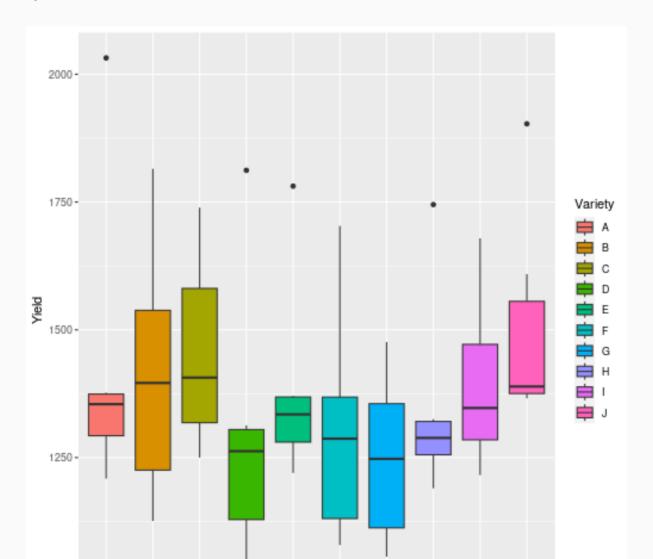
## Analysis of Variance Table
##
## Response: LIFETIME
## Df Sum Sq Mean Sq F value Pr(>F)
## DIET 5 12734 2546.8 57.104 < 2.2e-16 ***
## Residuals 343 15297 44.6
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1</pre>
```

Linseed Experiment

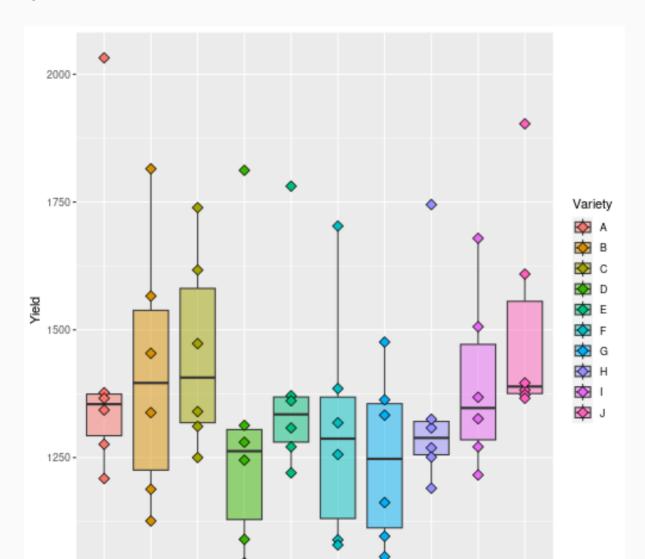
```
## Rows: 60 Columns: 3
## — Column specification -
## Delimiter: ","
## chr (1): Variety
## dbl (2): Plot. Yield
###
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show col types = FALSE` to quiet this message.
## # A tibble: 10 × 4
   Variety n mean
##
                             sd
     <chr>
             <int> <dbl> <dbl>
###
##
   1 A
                  6 1434. 300.
   2 B
                  6 1414. 255.
##
   3 C
                  6 1455 191.
###
   4 D
                  6 1297. 274.
###
   5 E
                  6 1385. 202.
##
###
   6 F
                  6 1305
                          230.
                  6 1248. 167.
   7 G
##
   8 H
                  6 1348
                          200.
###
   9 I
                  6 1394. 171.
##
## 10 J
                  6 1505.
                         216.
```

20 / 23

Linseed Experiment



Linseed Experiment



Linseed Experiment