

STAT 8010 R Lab 11

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AONVA Toy Examples

```
set.seed(1)
base1 <- rnorm(n = 36, sd = 2)
base2 <- rnorm(n = 36, sd = 6)
dat1 <- base1 + c(rep(5, 12), rep(10, 12), rep(15, 12))
dat2 <- base2 + c(rep(5, 12), rep(10, 12), rep(15, 12))
dat3 <- base1 + rep(5:7, each = 12)
level <- as.factor(rep(1:3, each = 12))
dat1 <- data.frame(x = dat1, Group = level)
dat2 <- data.frame(x = dat2, Group = level)
dat3 <- data.frame(x = dat3, Group = level)
library(dplyr)
g1summary <- dat1 %>%
  select(x, Group) %>%
  group_by(Group) %>%
  summarise(mean = mean(x), sd1 = sd(x))

g2summary <- dat2 %>%
  select(x, Group) %>%
  group_by(Group) %>%
  summarise(mean = mean(x), sd1 = sd(x))

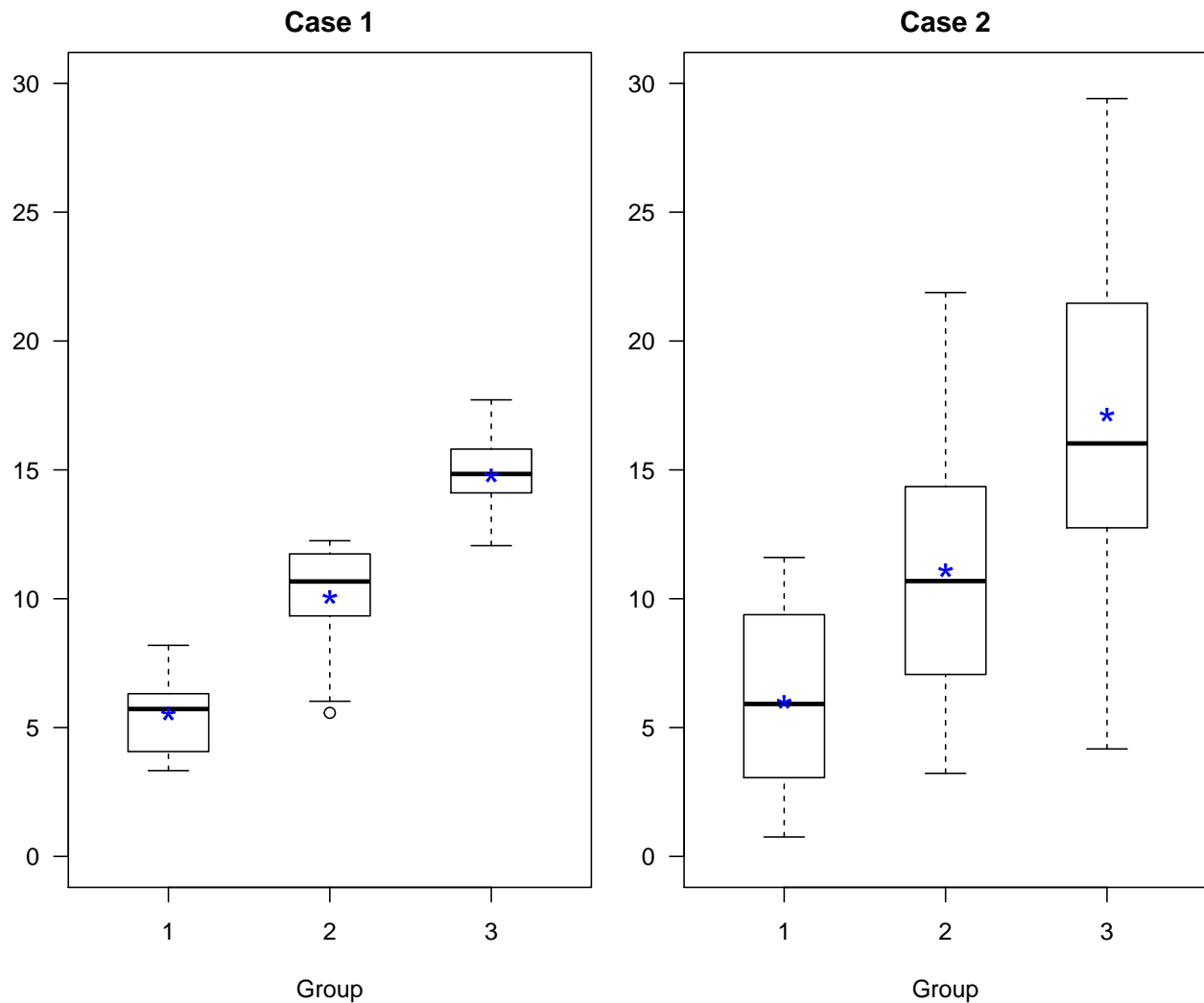
g3summary <- dat3 %>%
  select(x, Group) %>%
  group_by(Group) %>%
  summarise(mean = mean(x), sd1 = sd(x))

par(mfrow = c(1, 2), mar = c(4.1, 2.1, 2.1, 1.1))
boxplot(x ~ Group, data = dat1, las = 1, boxwex = 0.5,
        ylab = "", ylim = c(0, 30), main = "Case 1")
for (i in 1:3) points(i, g1summary$mean[i], pch = "*",
                      col = "blue", cex = 2)
boxplot(x ~ Group, data = dat2, las = 1, boxwex = 0.5,
```

```

      ylab = "", ylim = c(0, 30), main = "Case 2")
for (i in 1:3) points(i, g2summary$mean[i], pch = "*",
                      col = "blue", cex = 2)

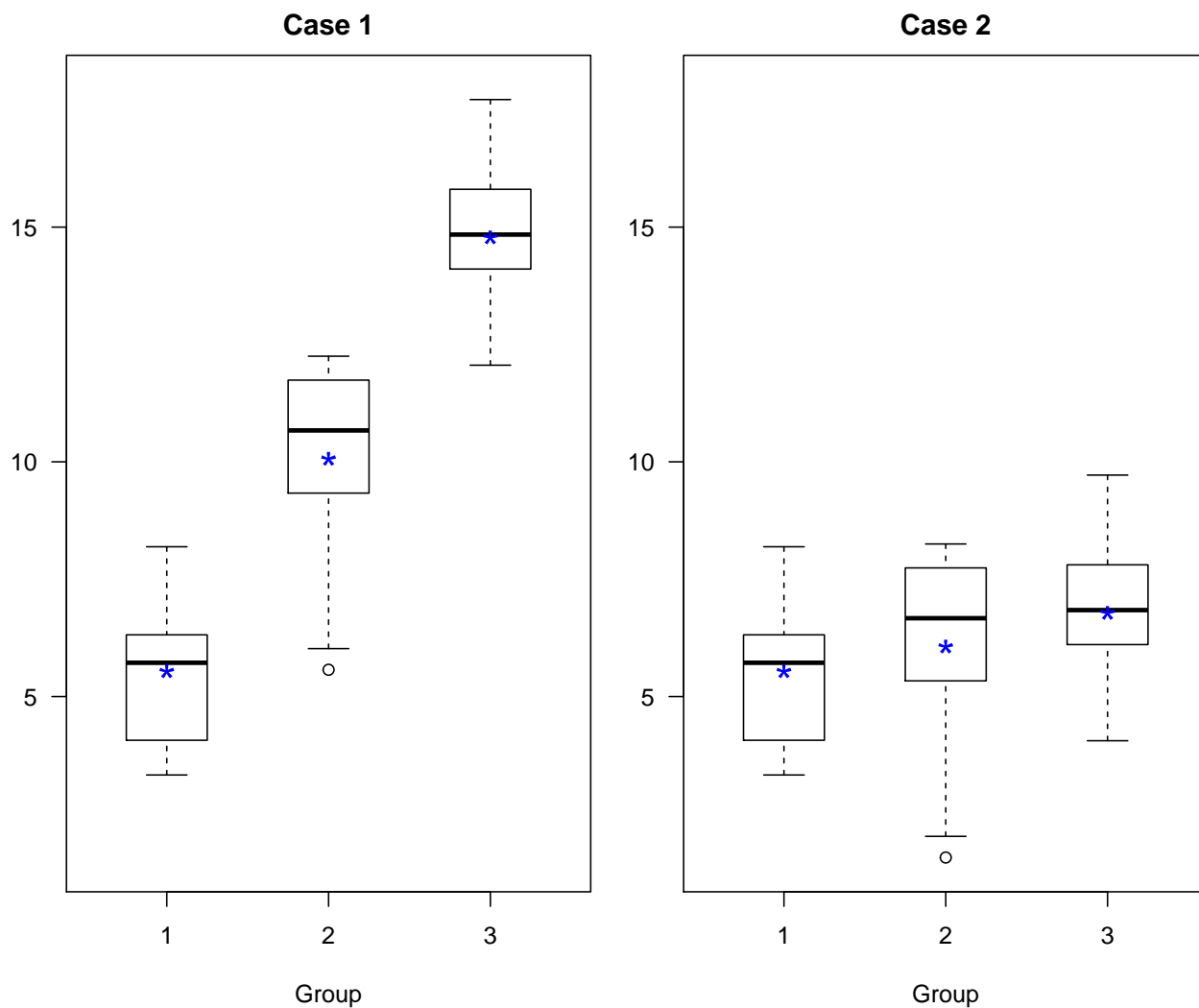
```



```

par(mfrow = c(1, 2), mar = c(4.1, 2.1, 2.1, 1.1))
boxplot(x ~ Group, data = dat1, las = 1, boxwex = 0.5,
        ylab = "", ylim = c(1.5, 18), main = "Case 1")
for (i in 1:3) points(i, g1summary$mean[i], pch = "*",
                      col = "blue", cex = 2)
boxplot(x ~ Group, data = dat3, las = 1, boxwex = 0.5,
        ylab = "", ylim = c(1.5, 18), main = "Case 2")
for (i in 1:3) points(i, g3summary$mean[i], pch = "*",
                      col = "blue", cex = 2)

```



```
model1 <- lm(x ~ Group, data = dat1)
model2 <- lm(x ~ Group, data = dat2)
anova(model1)
```

```
## Analysis of Variance Table
##
## Response: x
##          Df Sum Sq Mean Sq F value    Pr(>F)
## Group      2  512.54  256.271   75.443 4.902e-13 ***
## Residuals 33  112.10    3.397
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

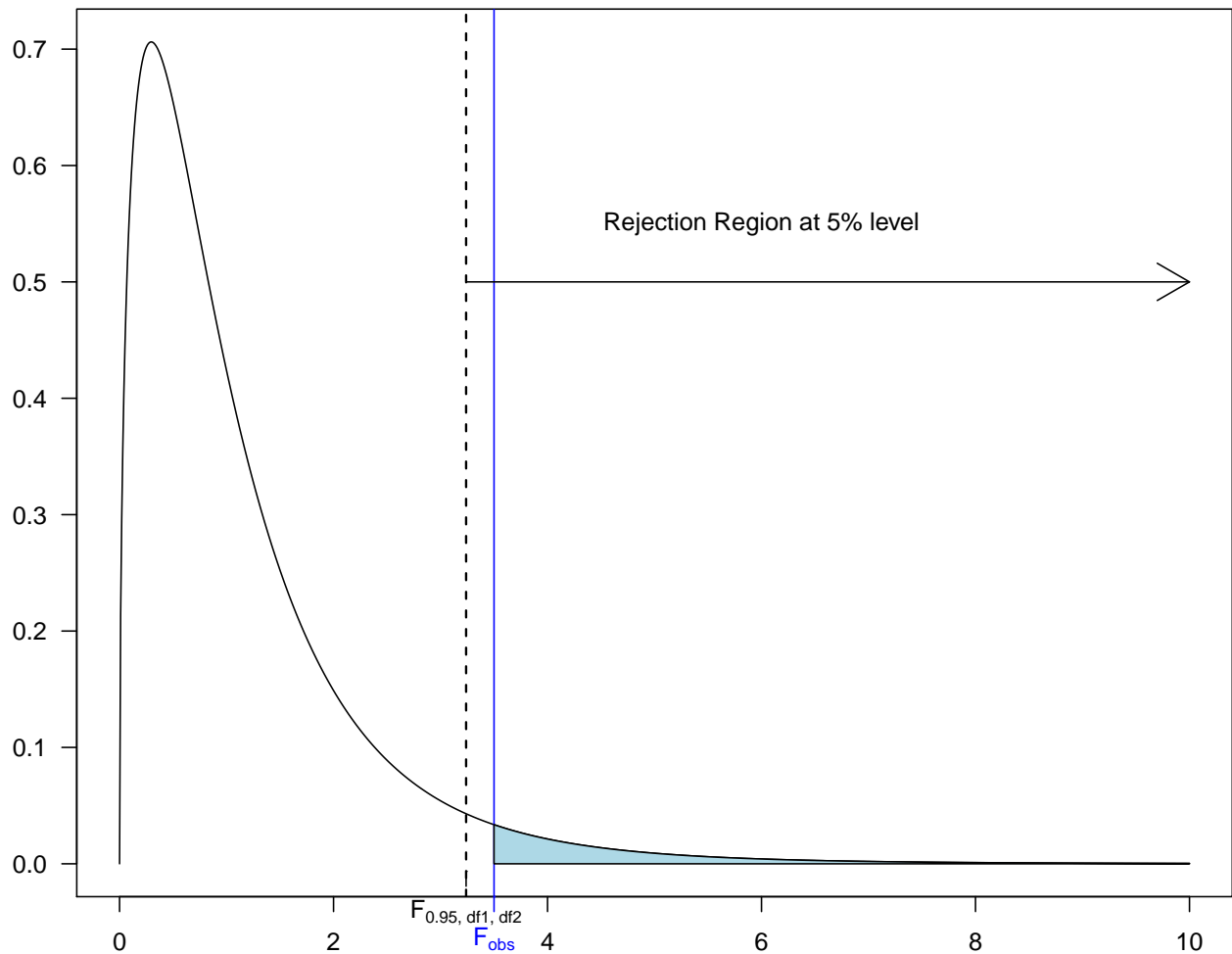
```
anova(model2)
```

```
## Analysis of Variance Table
##
## Response: x
##          Df Sum Sq Mean Sq F value    Pr(>F)
## Group      2   747.35   373.67  11.323 0.0001802 ***
## Residuals 33 1089.03    33.00
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

F Distribution

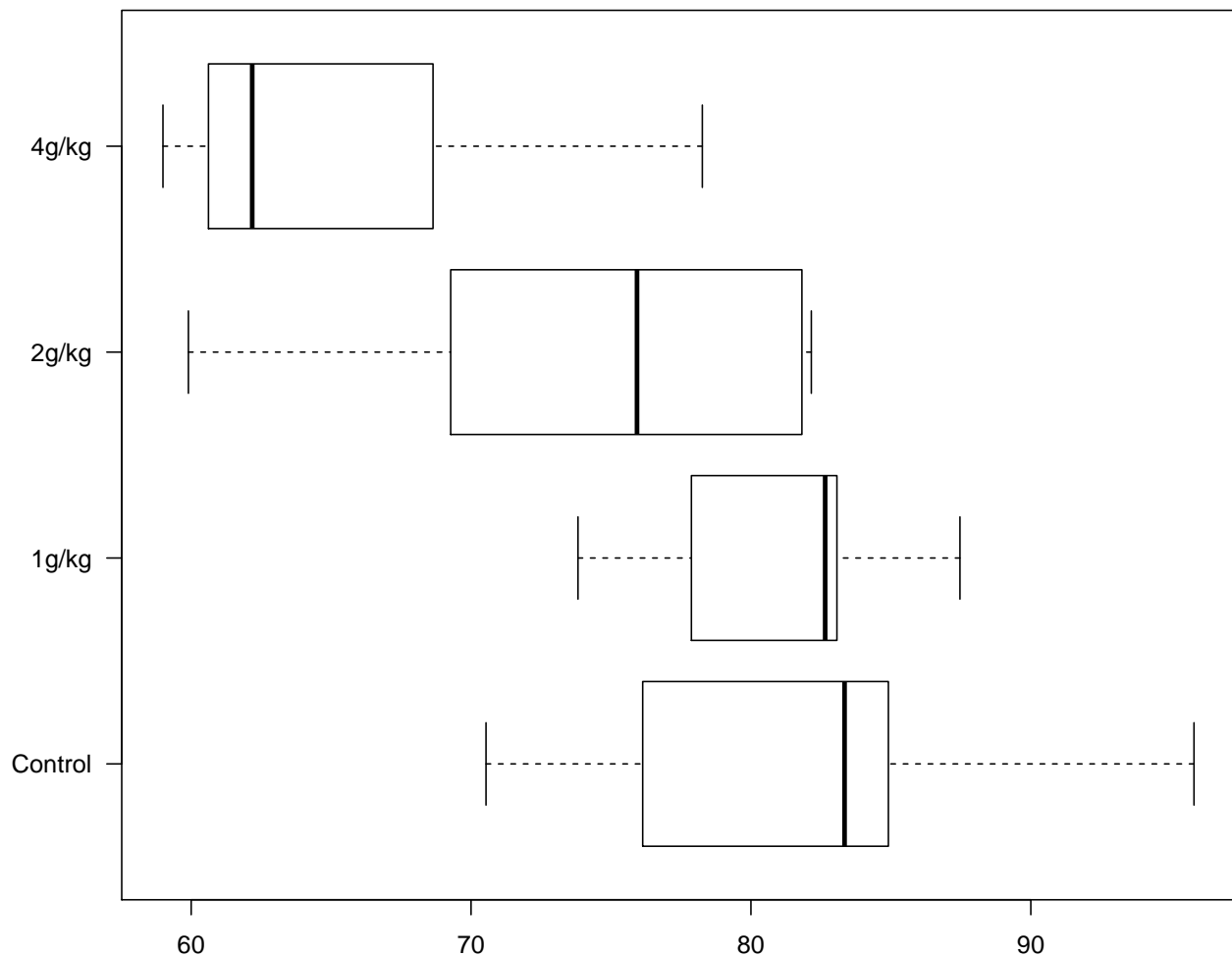
```
par(mar = c(4.1, 2.6, 1.1, 1.1))
curve(df(x, 3, 16), from = 0, to = 10, n = 1001, las = 1,
      xlab = "", ylab = "")
abline(v = 3.5, col = "blue")
abline(v = qf(0.95, 3, 16), lty = 2, lwd = 1.5)
xg <- seq(3.5, 10, 0.01)
yg <- df(xg, 3, 16)
polygon(c(xg[xg >= 3.5], rev(c(xg[xg >= 3.5]))), c(yg[xg >= 3.5], rep(0, length(yg[xg >= 3.5]))),
        col = "lightblue")
axis(1, at = 3.5, labels = expression(F["obs"]), col = "blue", col.axis = "blue")
axis(1, at = qf(0.95, 3, 16), line = -0.85, labels = expression(F[paste(0.95, ", df1", ", df2")]))
arrows(qf(0.95, 3, 16), 0.5, 10)
text(6, 0.55, "Rejection Region at 5% level")
```



Effects of Ethanol on Sleep Time Example

Data setup

```
set.seed(124)
g1 <- rnorm(5, 83, 9); g2 <- rnorm(5, 76, 9.5); g3 <- rnorm(5, 73, 9.2); g4 <- rnorm(5, 70, 9)
dat <- cbind(Response = c(g1, g2, g3, g4), Treatment = as.factor(rep(1:4, each = 5)))
dat <- data.frame(dat)
dat$Treatment <- as.factor(dat$Treatment)
par(mar = c(4.1, 4.1, 1.1, 1.1))
boxplot(Response ~ Treatment, data = dat, horizontal = T, yaxt = "n", ylab = "", xlab = "")
axis(2, at = 1:4, labels = c("Control", "1g/kg", "2g/kg", "4g/kg"), las = 1)
```



Data Summary

```
summary <- dat %>%
select(Response, Treatment) %>%
group_by(Treatment) %>%
summarise(mean = mean(Response),
           sdi = sd(Response))
lm <- lm(Response ~ Treatment, dat)
anova(lm)
```

```
## Analysis of Variance Table
##
## Response: Response
##           Df Sum Sq Mean Sq F value    Pr(>F)
## Treatment  3  861.13  287.044   4.2542 0.02173 *
## Residuals 16 1079.56   67.472
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Pairwise t-test

```
t.test(dat$Response[1:5], dat$Response[6:10], var.equal = T)
```

```
##
## Two Sample t-test
##
## data:  dat$Response[1:5] and dat$Response[6:10]
## t = 0.24012, df = 8, p-value = 0.8163
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -10.09426  12.44081
## sample estimates:
## mean of x mean of y
##  82.15052  80.97724
```

```
t.test(dat$Response[1:5], dat$Response[6:10], var.equal = F)
```

```
##
## Welch Two Sample t-test
##
## data:  dat$Response[1:5] and dat$Response[6:10]
## t = 0.24012, df = 6.2015, p-value = 0.818
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -10.68922  13.03577
## sample estimates:
## mean of x mean of y
##  82.15052  80.97724
```

Facebook Example

```
dat <- read.csv("FacebookFriends.csv")
head(dat); str(dat)
```

```
##   Friends Participant Score
## 1      102           1   3.8
## 2      102           2   3.6
## 3      102           3   3.2
## 4      102           4   2.4
## 5      102           5   4.8
## 6      102           6   3.0

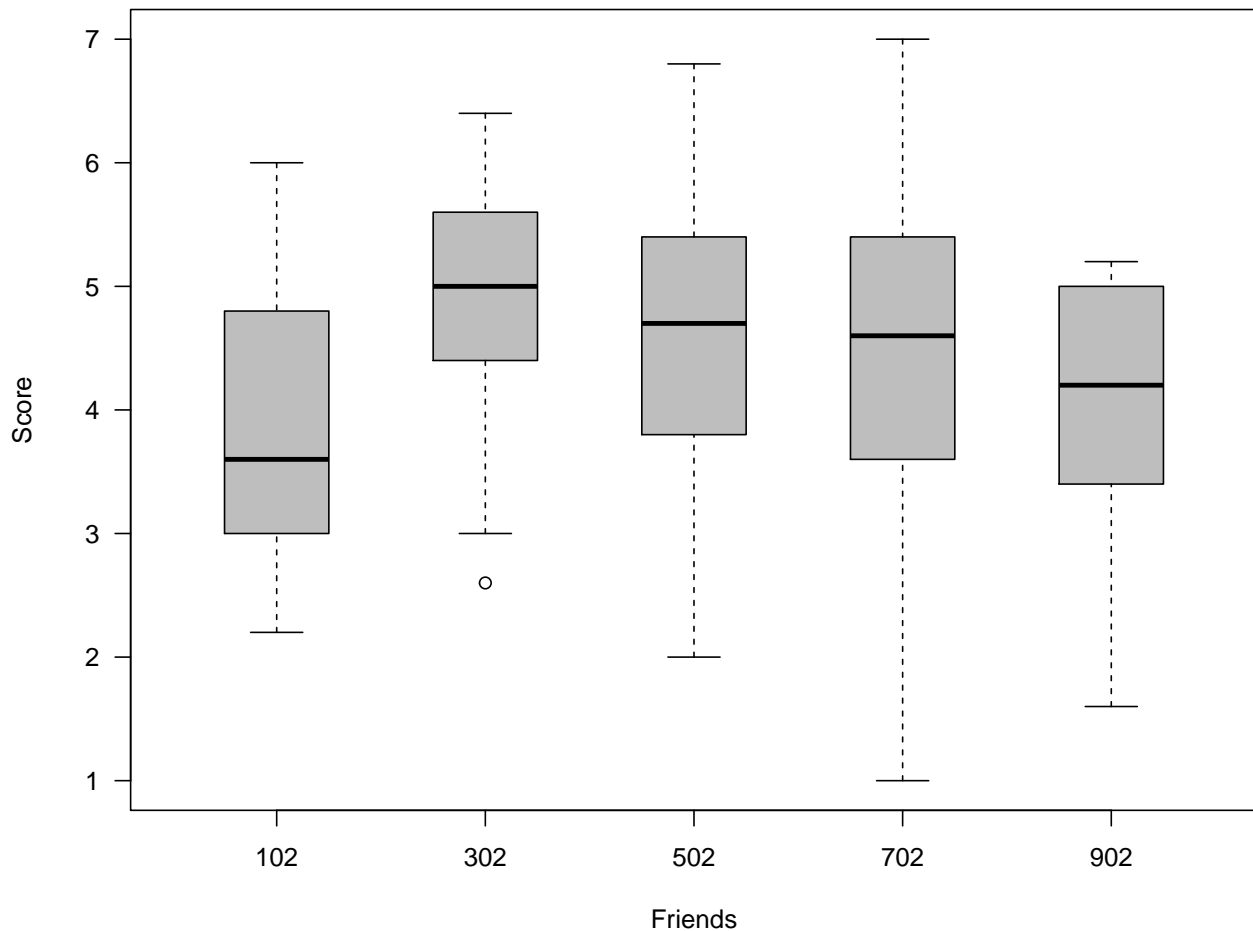
## 'data.frame':   134 obs. of  3 variables:
##  $ Friends      : int  102 102 102 102 102 102 102 102 102 102 ...
##  $ Participant: int   1 2 3 4 5 6 7 8 9 10 ...
```

```
## $ Score      : num  3.8 3.6 3.2 2.4 4.8 3 4.2 3.6 3.2 3 ...
```

```
dat$Friends <- as.factor(dat$Friends)
str(dat)
```

```
## 'data.frame': 134 obs. of 3 variables:
## $ Friends : Factor w/ 5 levels "102","302","502",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ Participant: int  1 2 3 4 5 6 7 8 9 10 ...
## $ Score : num  3.8 3.6 3.2 2.4 4.8 3 4.2 3.6 3.2 3 ...
```

```
boxplot(Score ~ Friends, data = dat, las = 1, col = "gray", boxwex = 0.5)
```



```
library(dplyr)
summary <- dat %>%
  select(Score, Friends) %>%
  group_by(Friends) %>%
  summarise(mean = mean(Score),
            sd1 = sd(Score))
summary
```

```
## # A tibble: 5 x 3
##   Friends mean  sd1
##   <fct>   <dbl> <dbl>
## 1 102     3.82 0.999
## 2 302     4.88 0.851
## 3 502     4.56 1.07
```

```
## 4 702      4.41 1.43
## 5 902      3.99 1.02
```

```
lm <- lm(Score ~ Friends, dat)
anova(lm)
```

```
## Analysis of Variance Table
```

```
##
```

```
## Response: Score
```

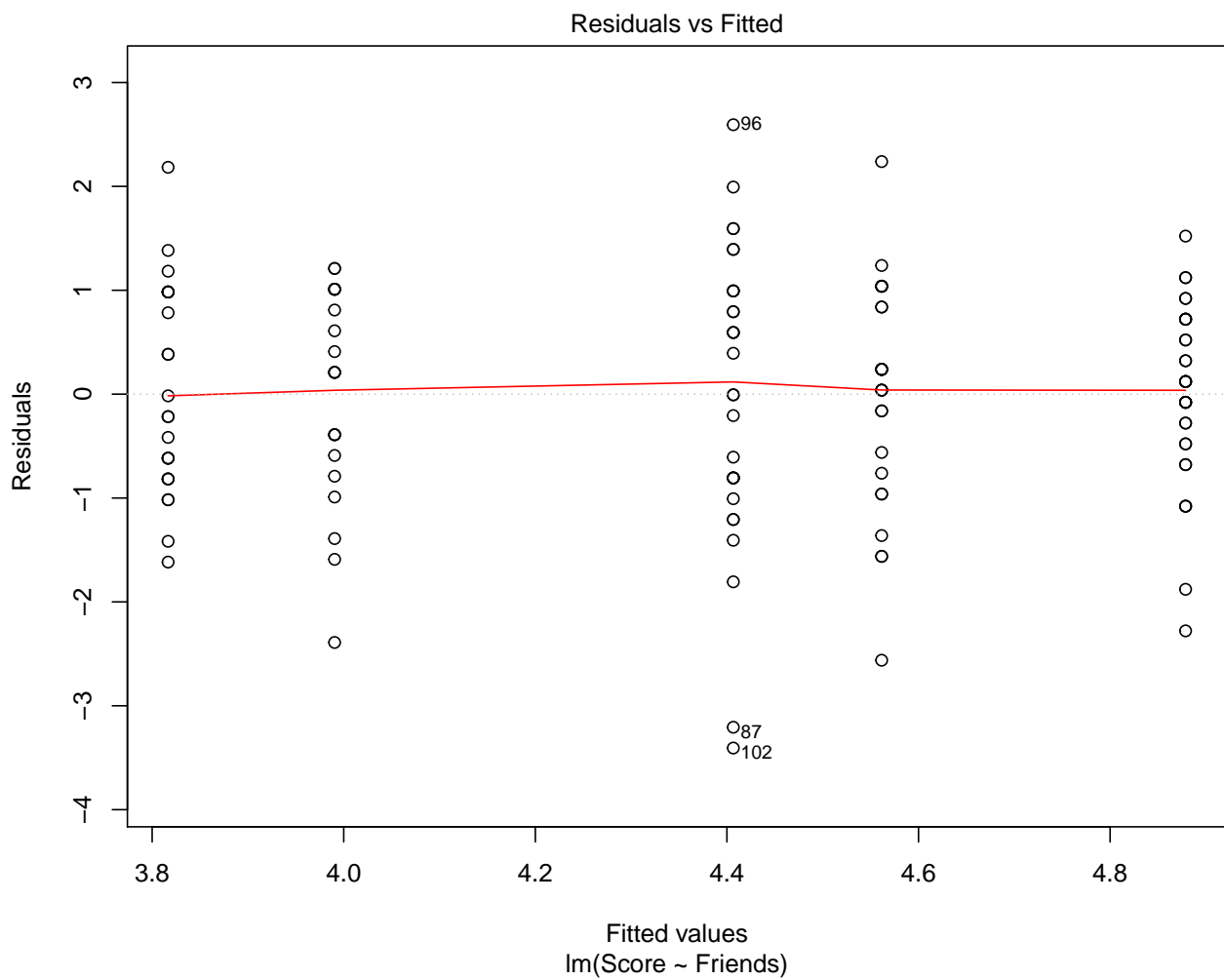
```
##           Df Sum Sq Mean Sq F value    Pr(>F)
## Friends      4   19.89   4.9726    4.142 0.00344 **
```

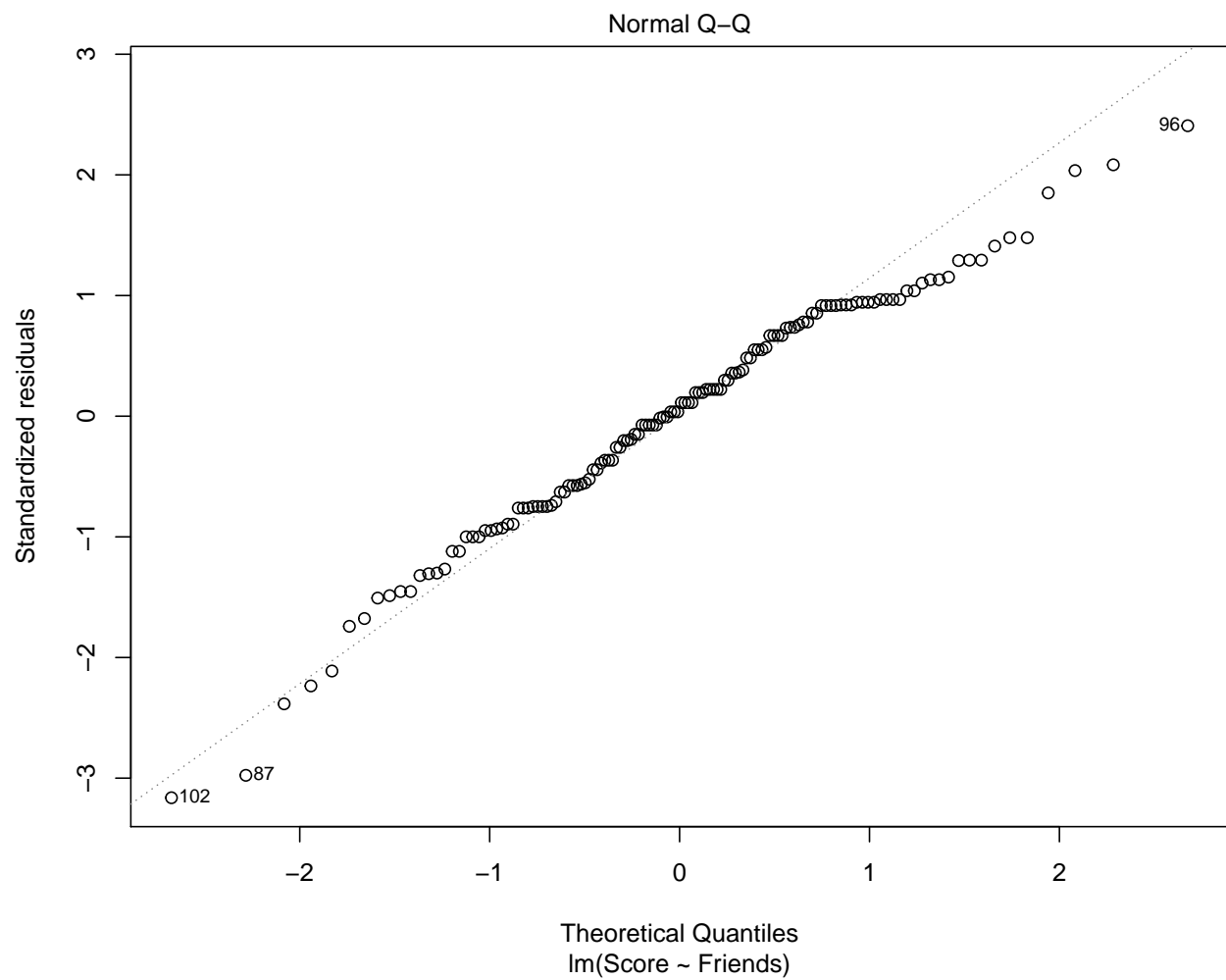
```
## Residuals 129  154.87   1.2005
```

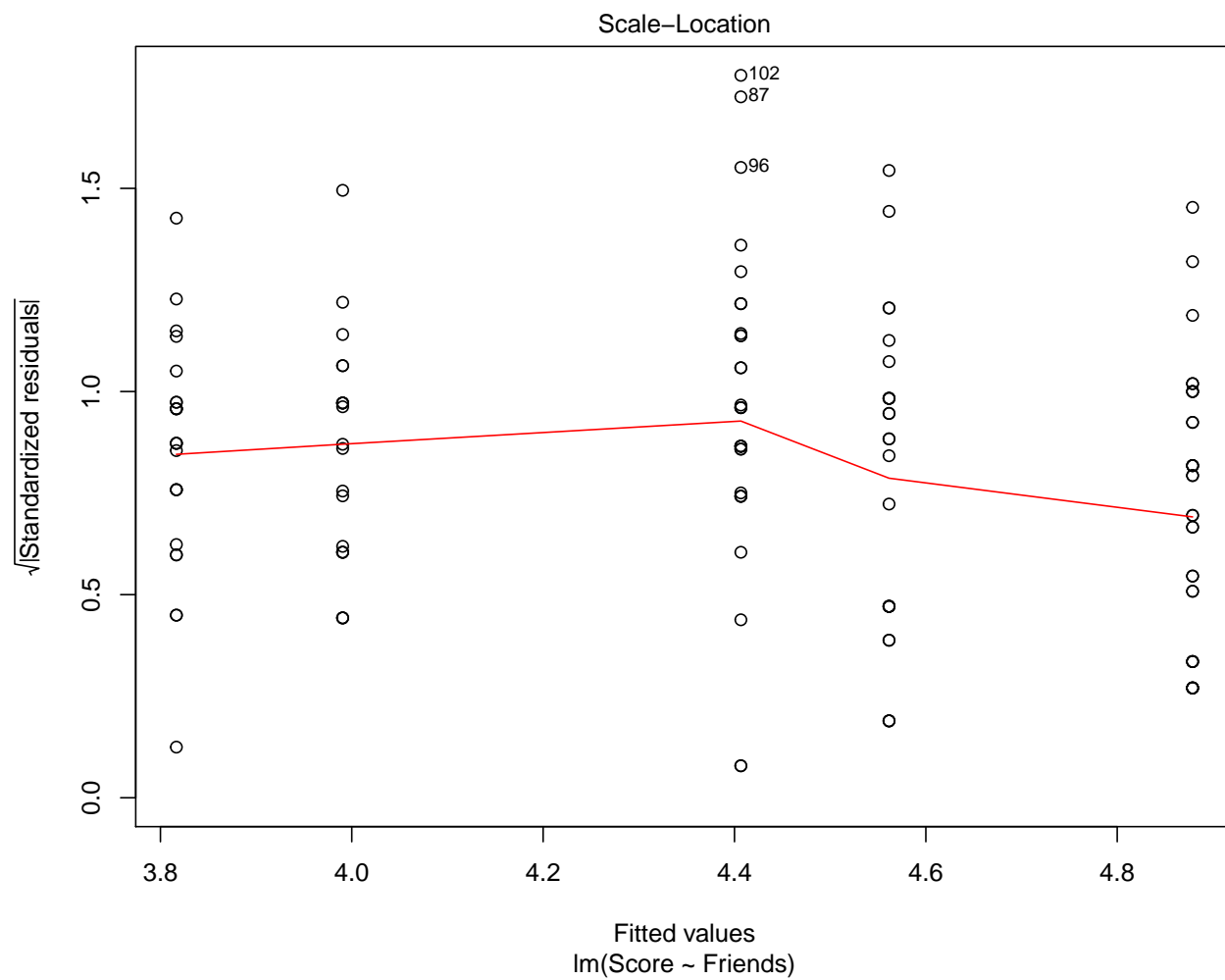
```
## ---
```

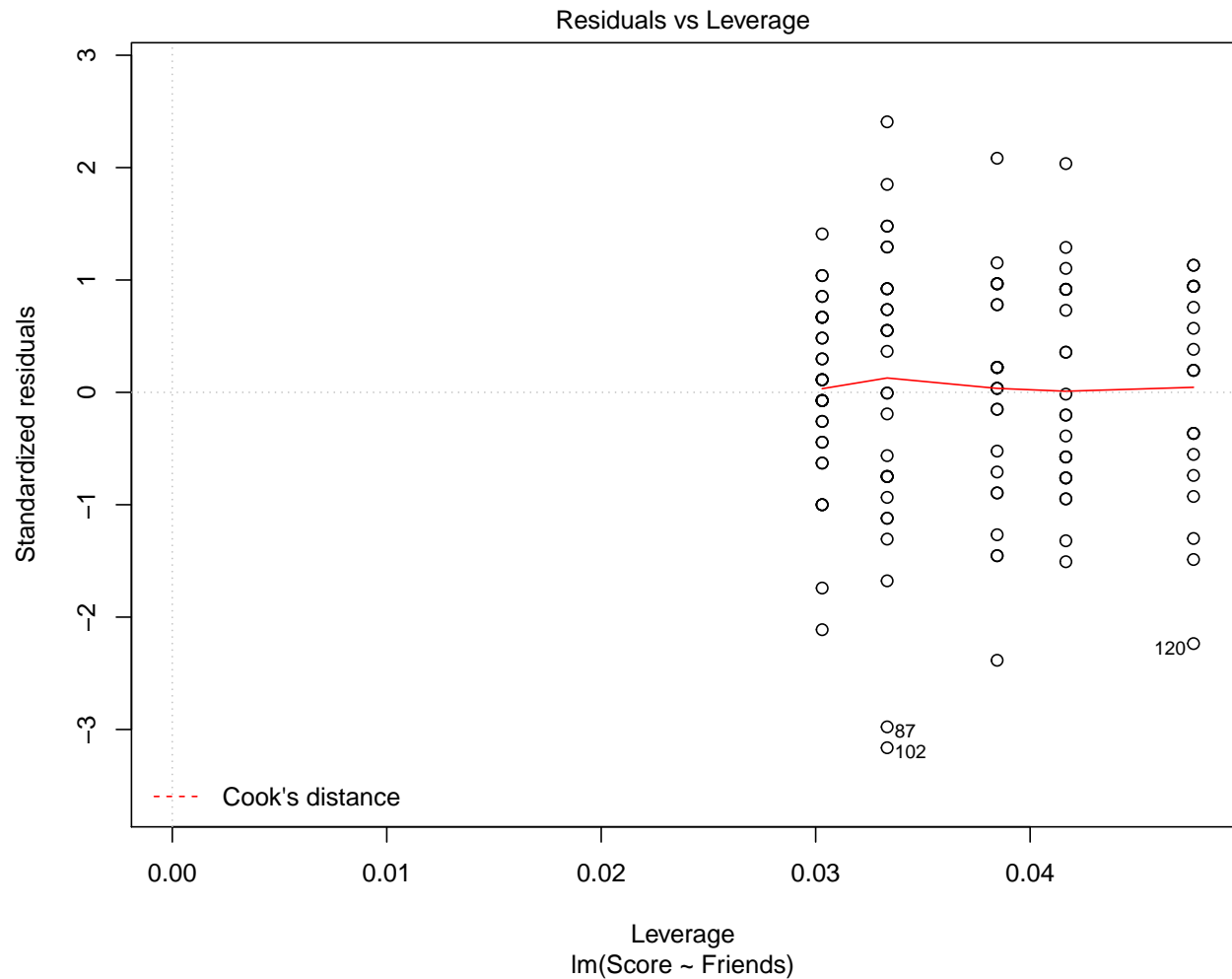
```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
plot(lm)
```









```
aov <- aov(Score ~ Friends, dat)
aov
```

```
## Call:
## aov(formula = Score ~ Friends, data = dat)
##
## Terms:
##             Friends Residuals
## Sum of Squares 19.89023 154.86679
## Deg. of Freedom      4      129
##
## Residual standard error: 1.095681
## Estimated effects may be unbalanced
```

Fisher's LSD

```
library(agricolae)
LSD_none <- LSD.test(aov, "Friends", p.adj = "none")
LSD_none$groups
```

```
##           Score groups
## 302 4.878788      a
## 502 4.561538     ab
```

```
## 702 4.406667    abc
## 902 3.990476    bc
## 102 3.816667    c
```

```
LSD_bon <- LSD.test(aov, "Friends", p.adj = "bonferroni")
LSD_bon$groups
```

```
##          Score groups
## 302 4.878788      a
## 502 4.561538     ab
## 702 4.406667     ab
## 902 3.990476      b
## 102 3.816667      b
```

Tukey's HSD

```
HSD <- TukeyHSD(aov, conf.level = 0.95)
HSD$Friends
```

##		diff	lwr	upr	p adj
##	302-102	1.0621212	0.2488644	1.87537798	0.003889635
##	502-102	0.7448718	-0.1132433	1.60298691	0.121456224
##	702-102	0.5900000	-0.2402014	1.42020143	0.288431585
##	902-102	0.1738095	-0.7320145	1.07963355	0.984016816
##	502-302	-0.3172494	-1.1121910	0.47769215	0.804080046
##	702-302	-0.4721212	-1.2368466	0.29260420	0.432633745
##	902-302	-0.8883117	-1.7345313	-0.04209203	0.034535577
##	702-502	-0.1548718	-0.9671402	0.65739661	0.984391504
##	902-502	-0.5710623	-1.4604793	0.31835479	0.391768065
##	902-702	-0.4161905	-1.2787075	0.44632652	0.669927748