

Assignment 1: R Competency — Laryngoscope

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Data Exploration

- This dataset comes from a study that tested whether using a video laryngoscope (camera) made intubation easier compared to the regular direct method.
- The patients were adults who needed intubation during surgery.
- The main treatment variable is **Randomization**:
 - 0 - direct laryngoscopy
 - 1 - video laryngoscopy
- The main outcome variables are:
 - **ease** (10 = easy intubation, 100 = difficult intubation)
 - **intubation_overall_S_F** (whether the intubation was a success or a failure)
 - **total_intubation_time** (how long the intubation took, measured in seconds)
 - **sore_throat** (whether the patient had a sore throat afterward)
- The dataset has **99 patients** (rows) and **22 variables** (columns).
- Some variables use numbers instead of words for coding, for example:
 - **gender** is coded as 0 = male, 1 = female
 - **ease** is coded as 10 = easy, 100 = difficult

```
?laryngoscope
```

```
names(laryngoscope)
```

```
## [1] "age" "gender"
## [3] "asa" "BMI"
## [5] "Mallampati" "Randomization"
## [7] "attempt1_time" "attempt1_S_F"
## [9] "attempt2_time" "attempt2_assigned_method"
## [11] "attempt2_S_F" "attempt3_time"
## [13] "attempt3_assigned_method" "attempt3_S_F"
## [15] "attempts" "failures"
## [17] "total_intubation_time" "intubation_overall_S_F"
## [19] "bleeding" "ease"
## [21] "sore_throat" "view"
```

The dataset has 22 variables.

```
unique(laryngoscope$gender)
```

```
## [1] 0 1
```

Gender is coded as 0 and 1, instead of “Male”/“Female.”

Data Analysis

```
table(laryngoscope$gender)
```

```
##
## 0 1
## 78 21
```

The output shows 78 males and 21 females, meaning about one-fifth of the sample were female.

```
# pick only females (1) and under 45
young_females <- laryngoscope[laryngoscope$gender == 1 & laryngoscope$age < 45, ]

# see the rows
young_females
```

There are 3 women younger than 45 in the dataset.

```
which(laryngoscope$gender == 1 & laryngoscope$age < 45)
```

```
## [1] 13 57 88
```

```
# check what "ease" values look like
unique(young_females$ease)
```

```
## [1] 10 100
```

```
# count them
table(young_females$ease)
```

```
##
## 10 100
##  1  2
```

Among these 3 women, 1 had an “easy” intubation (10), while 2 had a “difficult” intubation (100).

```
# keep only females in groups 0 and 1
females_groups <- laryngoscope[laryngoscope$gender == 1 & laryngoscope$Randomization %in% c(0,1), ]
```

```
# look at group 0 separately
quantile(females_groups$BMI[females_groups$Randomization == 0],
         probs = c(0.25, 0.5, 0.75), na.rm = TRUE)
```

```
##      25%      50%      75%
## 37.840 41.820 43.785
```

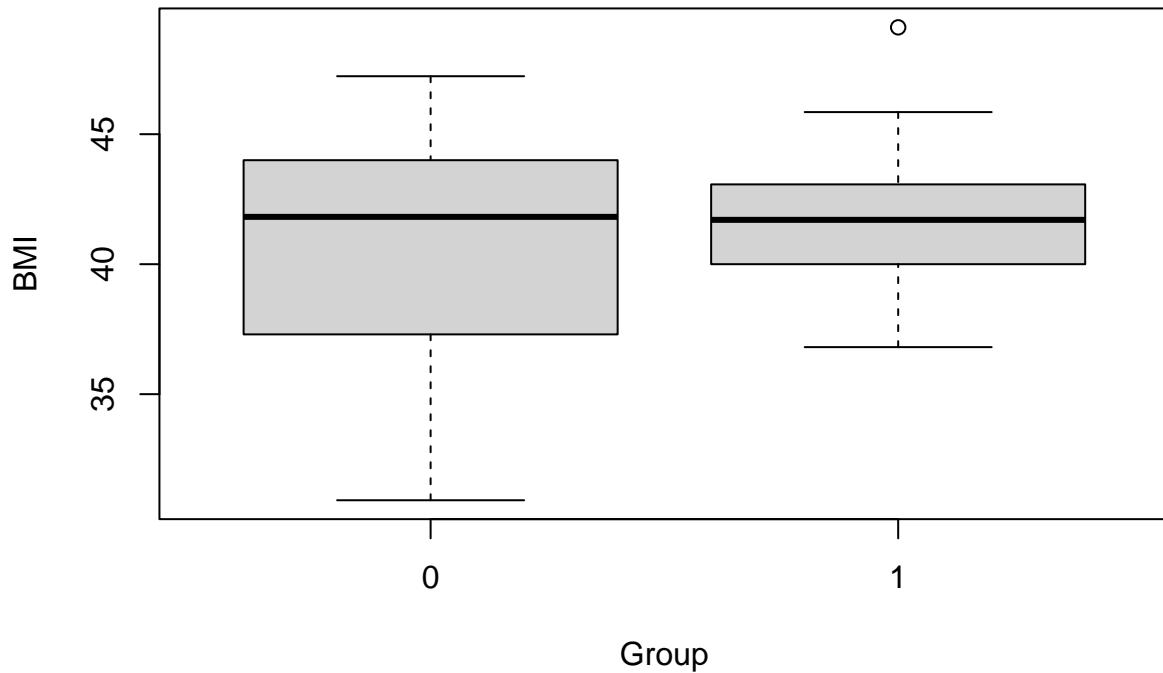
```
# look at group 1 separately
quantile(females_groups$BMI[females_groups$Randomization == 1],
         probs = c(0.25, 0.5, 0.75), na.rm = TRUE)
```

```
##      25%      50%      75%
## 40.3500 41.7050 42.8025
```

Median BMI in group 0 was ~41.8, with most values between ~37.8 and 43.8. Group 0 is more spread out than Group 1 because the BMI values cover a wider range. This means that BMIs among females in Group 0 were more variable, while those in Group 1 were more consistent and clustered around the median.

```
# simple boxplot
boxplot(BMI ~ Randomization, data = females_groups,
        main = "BMI of Females by Group",
        xlab = "Group", ylab = "BMI")
```

BMI of Females by Group

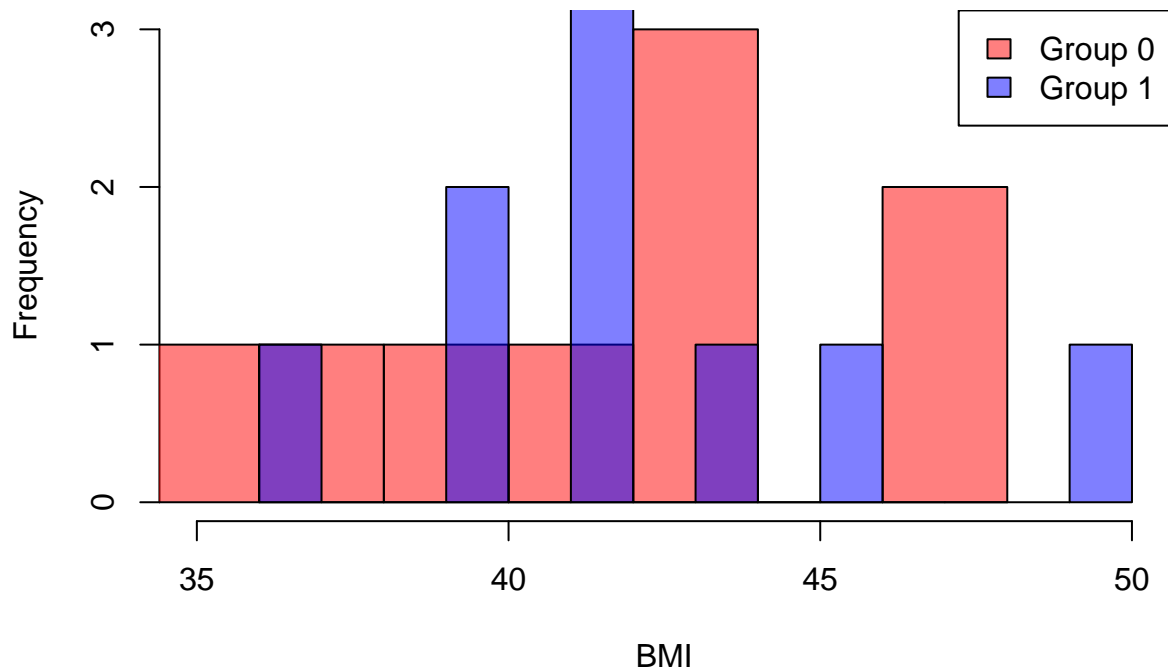


```
hist(females_groups$BMI[females_groups$Randomization == 0],
     main = "BMI Distribution for Females (Group 0 vs Group 1)",
     xlab = "BMI", col = rgb(1,0,0,0.5), xlim = c(35,50), breaks = 10)

hist(females_groups$BMI[females_groups$Randomization == 1],
     col = rgb(0,0,1,0.5), add = TRUE, breaks = 10)

legend("topright", legend = c("Group 0", "Group 1"),
     fill = c(rgb(1,0,0,0.5), rgb(0,0,1,0.5)))
```

BMI Distribution for Females (Group 0 vs Group 1)



The overlapping histograms show that BMI values for Groups 0 and 1 follow almost the same pattern, confirming the boxplot result that the two groups have very similar distributions.

Executive Summary

This project looked at the *laryngoscope* dataset, which tracks patients in a study comparing video versus direct laryngoscopy. There are 99 patients total: 78 men (coded as 0) and 21 women (coded as 1). When I filtered for younger women under 45, I found three patients. Out of these, two had noted an “easy” intubation, while one had a more difficult time.

I also compared BMI values for women in groups 0 and 1. The medians were very close (around 41 for group 0 and 42 for group 1), and the middle ranges overlapped substantially. Group 0 was a bit more spread out, with BMIs covering a wider range, while Group 1’s BMIs were more tightly clustered around the median. This means that BMI was distributed almost the same way in both groups overall, and while there were potentially a few individual outliers, there was no meaningful difference between the groups in terms of BMI.

Overall, the results show that most younger female patients in the dataset had an easy intubation, and there wasn’t much difference in BMI between the two treatment groups. It’s important to note that the subgroup of younger women was very small ($n = 3$), so these results cannot be generalized. Larger samples would be needed to confirm patterns in this age group.

AI Statement: I used AI to figure out RStudio and formatting and rmd format. I used prompts like: “How do I add a new code cell in RStudio?”, “What is the”{r}?” “. I, however, tried as much as possible to write my own code to learn R and I did not use GPT to generate code directly, only to tell me why it’s producing errors and why the formatting is wrong.