pst-semester-project

December 30, 2024

```
[10]: data("case0402", package = "Sleuth2")

# Setting seed for reproducibility.
set.seed(123)

[11]: # Loading libraries
library(knitr)
library(crayon)
library(ggplot2)

cat("\n")
cat(bold("Displaying the first few rows of data:"))
kable(head(case0402), format = "simple")
cat("\n")
```

Displaying the first few rows of data:

```
Time Treatmt Censor
----- 68 Modified 0
70 Modified 0
73 Modified 0
75 Modified 0
77 Modified 0
80 Modified 0
```

```
[12]: # Displaying the structure of the data
cat("\n")
cat(bold("Data Structure:\n"))
str(case0402)

# Table of the first rows of data in a nicer format
cat(bold("\nData Sample:\n"))
```

```
kable(head(case0402), format = "pipe", align = "c") # The 'pipe' format_
 ⇔improves readability
# Summary overview of the data
cat(bold("\nSummary Overview of Data:\n"))
summary table <- as.data.frame(summary(case0402)) # Convert to a table</pre>
kable(summary_table, format = "pipe", align = "l") # Table formatting
cat(bold("\nVisualization of Graphs:\n"))
# Boxplot by groups
ggplot(case0402, aes(x = Treatmt, y = Time, fill = Treatmt)) +
  geom_boxplot() +
 scale fill manual(values = c("Modified" = "lightgreen", "Conventional" = __

¬"skyblue")) +

 labs(title = "Boxplot of Time by Groups",
      x = "Group",
      y = "Time") +
  theme_minimal()
# Histograms for individual groups
ggplot(case0402, aes(x = Time, fill = Treatmt)) +
  geom_histogram(position = "dodge", bins = 10, alpha = 0.7) +
 scale fill manual(values = c("Modified" = "lightgreen", "Conventional" = __

¬"skyblue")) +

 labs(title = "Histogram of Time by Groups",
      x = "Time".
      y = "Count") +
  theme minimal()
# Density plot by groups
ggplot(case0402, aes(x = Time, color = Treatmt)) +
  geom_density(linewidth = 1) + # Using linewidth instead of size
 scale_color_manual(values = c("Modified" = "darkgreen", "Conventional" = __

¬"steelblue")) +

 labs(title = "Density of Time by Groups",
      x = "Time",
      y = "Density") +
  theme minimal()
```

Data Structure:

```
'data.frame': 28 obs. of 3 variables:

$ Time : num 68 70 73 75 77 80 80 132 148 155 ...

$ Treatmt: Factor w/ 2 levels "Modified", "Conventional": 1 1 1 1 1 1 1 1 1 1 ...
```

\$ Censor : num 0 0 0 0 0 0 0 0 0 0 ...

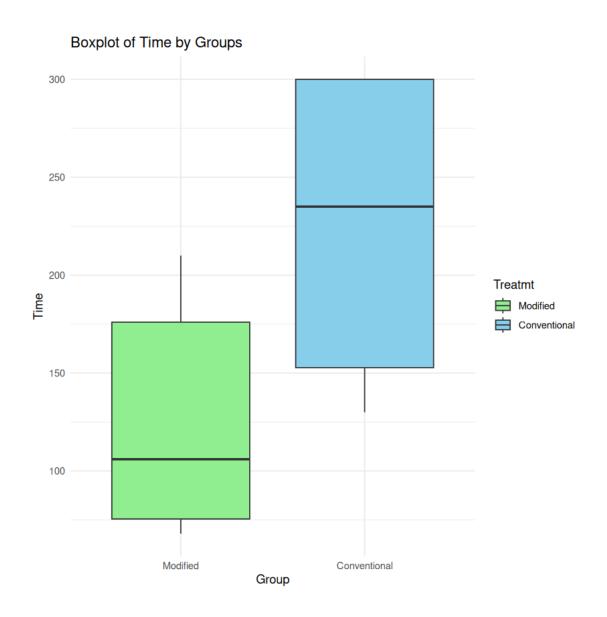
Data Sample:

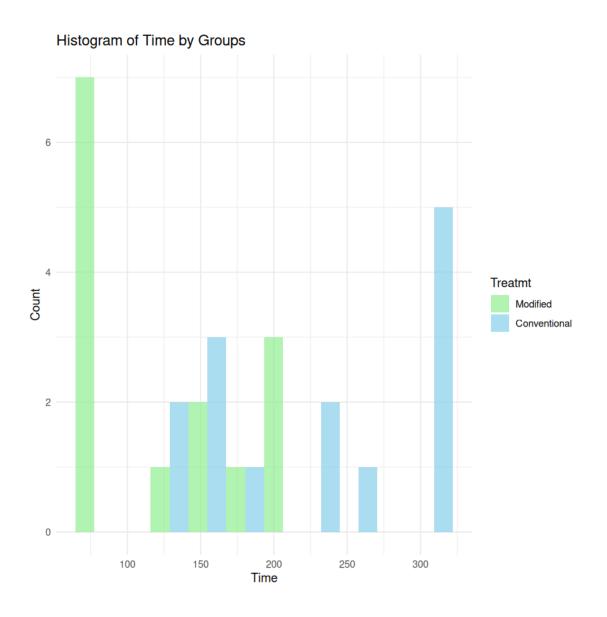
	Time		Treatmt		Censor	-
:		: :	::	1:	:	:
	68		Modified	1	0	-
	70		Modified	1	0	-
	73		Modified		0	-
	75		Modified	1	0	-
1	77		Modified	Ι	0	-
I	80		Modified	Ι	0	١

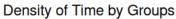
Summary Overview of Data:

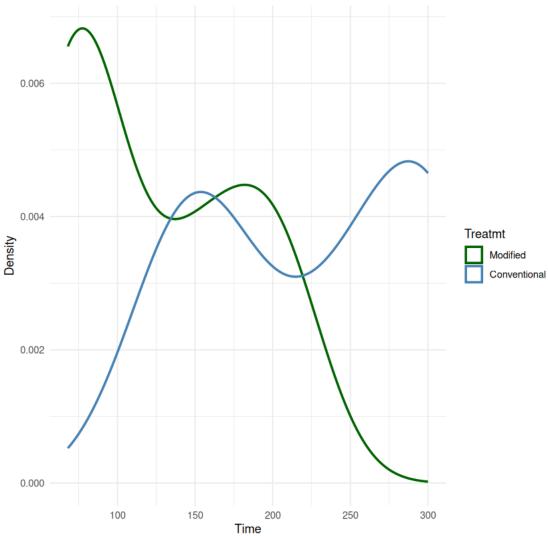
Var1	Var2	Freq
:	- :	- :
1	Time	Min. : 68.0
1	Time	1st Qu.:117.5
1	Time	Median :158.0
1	Time	Mean :174.7
1	Time	3rd Qu.:231.5
1	Time	Max. :300.0
1	Treatmt	Modified :14
1	Treatmt	Conventional:14
1	Treatmt	NA
1	Censor	Min. :0.0000
1	Censor	1st Qu.:0.0000
1	Censor	Median :0.0000

Visualization of Graphs:









```
[13]: library(ggplot2)

# Splitting into two groups based on the 'Treatmt' column
group1 <- subset(case0402, Treatmt == "Modified") # First group
group2 <- subset(case0402, Treatmt == "Conventional") # Second group

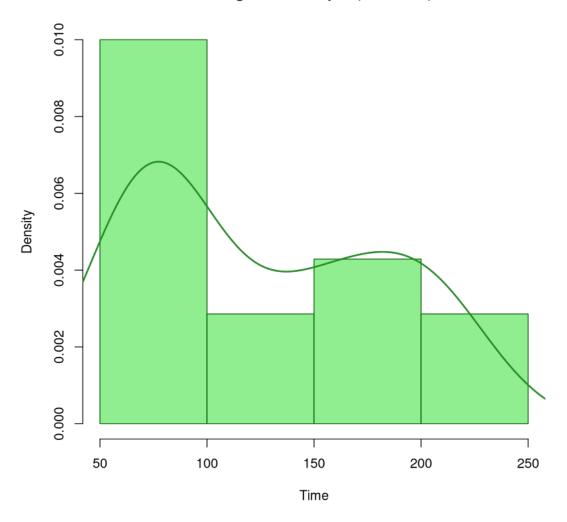
# Calculations for group 1
mean1 <- mean(group1$Time) # Mean
var1 <- var(group1$Time) # Variance
median1 <- median(group1$Time) # Median

# Calculations for group 2
mean2 <- mean(group2$Time)</pre>
```

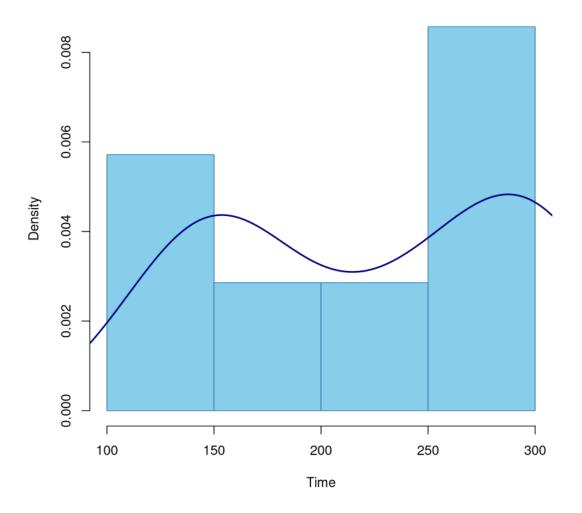
```
var2 <- var(group2$Time)</pre>
      median2 <- median(group2$Time)</pre>
      # Dynamic output with description for group 1
      cat(bold("\nResults for the first group (Modified):"))
      cat("\n- Mean:", mean1, "\n")
      cat("- Variance:", var1, "\n")
      cat("- Median:", median1, "\n\n")
      # Dynamic output with description for group 2
      cat(bold("Results for the second group (Conventional):"))
      cat("\n- Mean:", mean2, "\n")
      cat("- Variance:", var2, "\n")
      cat("- Median:", median2, "\n\n")
     Results for the first group (Modified):
     - Mean: 125.2857
     - Variance: 3203.297
     - Median: 106
     Results for the second group (Conventional):
     - Mean: 224.1429
     - Variance: 4976.901
     - Median: 235
[14]: # Loading the data file
      data("case0402", package = "Sleuth2")
      # Splitting into two groups based on the 'Treatmt' column
      group1 <- subset(case0402, Treatmt == "Modified") # Group 1</pre>
      group2 <- subset(case0402, Treatmt == "Conventional") # Group 2</pre>
      # Plotting histogram for group 1
      hist(
        group1$Time,
        main = "Histogram - Group 1 (Modified)",
        xlab = "Time",
       ylab = "Density",
        col = "lightgreen", # Fill color
        border = "darkgreen", # Border color
        freq = FALSE # Setting for density instead of frequency
      lines(density(group1$Time), col = "forestgreen", lwd = 2) # Density curve
      # Plotting histogram for group 2
```

```
hist(
 group2$Time,
 main = "Histogram - Group 2 (Conventional)",
 xlab = "Time",
 ylab = "Density",
 col = "skyblue", # Fill color
 border = "steelblue", # Border color
 freq = FALSE
lines(density(group2$Time), col = "navy", lwd = 2) # Density curve
# Empirical distribution function for group 1
plot(
 ecdf(group1$Time),
 main = "Empirical Distribution Function - Group 1 (Modified)",
 xlab = "Time",
 ylab = "Probability",
 col = "darkgreen",
 lwd = 2
# Empirical distribution function for group 2
plot(
 ecdf(group2$Time),
 main = "Empirical Distribution Function - Group 2 (Conventional)",
 xlab = "Time",
 ylab = "Probability",
 col = "navy",
 lwd = 2
```

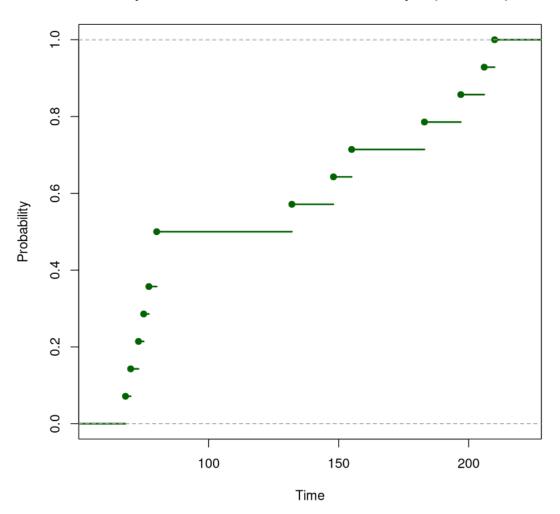
Histogram - Group 1 (Modified)



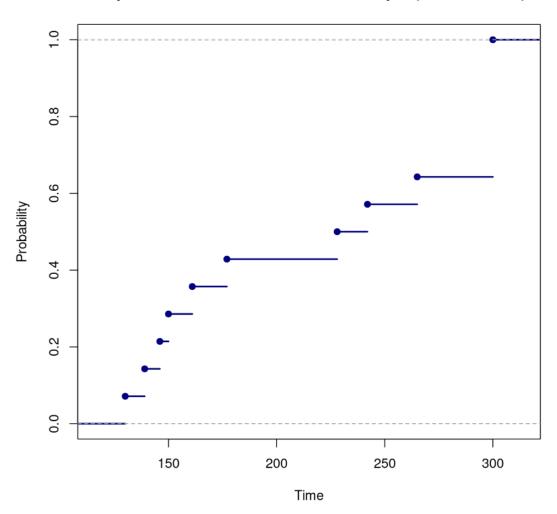
Histogram - Group 2 (Conventional)



Empirical Distribution Function - Group 1 (Modified)



Empirical Distribution Function - Group 2 (Conventional)



```
[19]: cat("\n")
# Function to plot density with histogram
plot_densities <- function(data, group) {
    # Histogram
    hist(
        data,
        freq = FALSE,
        main = paste("Histogram with Densities -", group),
        xlab = "Time",
        ylab = "Density",
        col = if (group == "Group 1 (Modified)") "lightgreen" else "skyblue",
        border = if (group == "Group 1 (Modified)") "darkgreen" else "steelblue"
    )</pre>
```

```
# Parameter estimation
mean_val <- mean(data) # Mean - estimated as the arithmetic average of the
\hookrightarrow data
sd val <- sd(data)
                         # Standard deviation - calculated from the data using
⇔the standard sd() function
lambda_val <- 1 / mean_val # Exponential parameter - estimated as 1/mean
min_val <- min(data)</pre>
                       # Minimum for uniform distribution - obtained as the
→minimum of the data
max val <- max(data)</pre>
                       # Maximum for uniform distribution - obtained as the
→maximum of the data
# Explanation of parameter estimation
cat(bold(paste0("\nExplanation of Parameter Estimation for ", group, ":")), ___
\hookrightarrow"\n")
cat("- Normal Distribution: Mean (arithmetic average) and standard deviation ⊔
⇒calculated directly from the data.\n")
cat("- Exponential Distribution: Lambda (rate) is estimated as 1/mean.\n")
cat("- Uniform Distribution: Min and Max are estimated as the smallest and []
⇒largest values of the data.\n\n")
# Normal density
curve(dnorm(x, mean = mean val, sd = sd val),
      col = "red", lwd = 2, add = TRUE, lty = 1)
# Exponential density
curve(dexp(x, rate = lambda_val),
      col = "blue", lwd = 2, add = TRUE, lty = 2)
# Uniform density
curve(dunif(x, min = min_val, max = max_val),
      col = "purple", lwd = 2, add = TRUE, lty = 3)
# Legend
legend_position <- if (group == "Group 1 (Modified)") "topright" else⊔
legend(legend_position, legend = c("Normal", "Exponential", "Uniform"),
       col = c("red", "blue", "purple"), lty = 1:3, cex = 0.8)
# Displaying parameters
cat(bold(paste0("Explanation of Parameter Estimation for ", group, ":")), u
cat("- Normal: mean =", round(mean_val, 2), "\n")
cat("- Standard Deviation =", round(sd val, 2), "\n")
cat("- Exponential: lambda =", round(lambda_val, 2), "\n")
```

```
cat("- Uniform: min =", round(min val, 2), ", max =", round(max val, 2), "
 \hookrightarrow"\n\n")
  # Discussion for individual groups
  if (group == "Group 1 (Modified)") {
    cat(bold("Discussion and Reflection for Group 1 (Modified):"))
    cat("\nAfter analyzing the graph, it is evident that the data for Group 1_{\sqcup}
 → (Modified) better fits an exponential distribution. The shape of the ...
 \hookrightarrowhistogram and the exponential density curve indicate an asymmetrical_{\sqcup}
 ⇔distribution with a higher number of lower time values.\n\n")
  } else if (group == "Group 2 (Conventional)") {
    cat(bold("Discussion and Reflection:"))
    cat("\nGroup 2 (Conventional):\n")
    cat("After analyzing the graph, it is clear that the data for Group 2_{\sqcup}
 _{\hookrightarrow} (Conventional) do not exhibit a clearly fitting distribution. Although the _{\sqcup}
 ⇔normal distribution (red line) roughly matches the shape of the histogram, ⊔
 \negdifferences at the data extremes suggest deviations from the theoretical\sqcup
 ⇔normal distribution.\n")
    cat("\n")
    \mathtt{cat}("\mathtt{However}, \ \mathtt{the} \ \mathtt{exponential} \ \mathtt{distribution} \ \mathtt{(blue} \ \mathtt{line)} \ \mathtt{and} \ \mathtt{the} \ \mathtt{uniform}_\sqcup
 ⊸distribution (purple line) do not sufficiently approximate the histogram. ⊔
 \hookrightarrowTherefore, the normal distribution appears to be the best approximation, \sqcup
 ⇒albeit with certain reservations. This data group likely requires a more
 \hookrightarrowdetailed analysis or tests for a more accurate identification of the \sqcup
 ⇒appropriate distribution model.\n\n")
 }
}
# Plotting for Group 1
plot_densities(group1$Time, "Group 1 (Modified)")
# Plotting for Group 2
plot_densities(group2$Time, "Group 2 (Conventional)")
```

Explanation of Parameter Estimation for Group 1 (Modified):

- Normal Distribution: Mean (arithmetic average) and standard deviation calculated directly from the data.
- Exponential Distribution: Lambda (rate) is estimated as 1/mean.
- Uniform Distribution: Min and Max are estimated as the smallest and largest values of the data.

Explanation of Parameter Estimation for Group 1 (Modified):

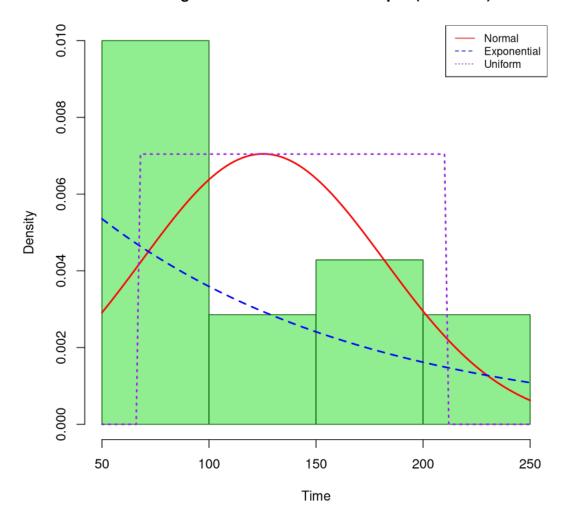
```
Normal: mean = 125.29Standard Deviation = 56.6Exponential: lambda = 0.01
```

- Uniform: min = 68, max = 210

Discussion and Reflection for Group 1 (Modified):

After analyzing the graph, it is evident that the data for Group 1 (Modified) better fits an exponential distribution. The shape of the histogram and the exponential density curve indicate an asymmetrical distribution with a higher number of lower time values.

Histogram with Densities - Group 1 (Modified)



Explanation of Parameter Estimation for Group 2 (Conventional):

- Normal Distribution: Mean (arithmetic average) and standard deviation calculated directly from the data.

- Exponential Distribution: Lambda (rate) is estimated as 1/mean.
- Uniform Distribution: Min and Max are estimated as the smallest and largest values of the data.

Explanation of Parameter Estimation for Group 2 (Conventional):

- Normal: mean = 224.14

- Standard Deviation = 70.55

- Exponential: lambda = 0

- Uniform: min = 130, max = 300

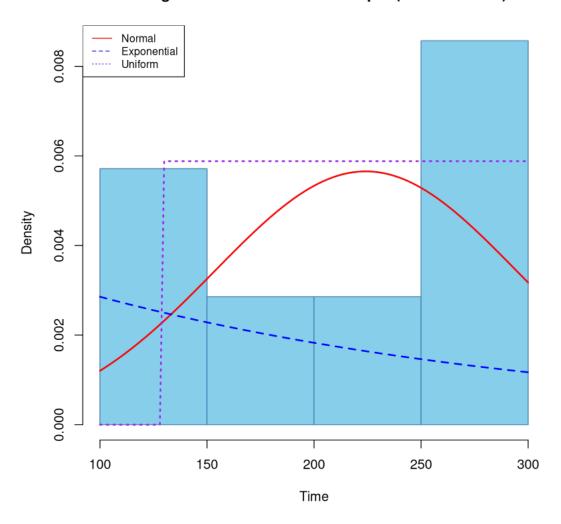
Discussion and Reflection:

Group 2 (Conventional):

After analyzing the graph, it is clear that the data for Group 2 (Conventional) do not exhibit a clearly fitting distribution. Although the normal distribution (red line) roughly matches the shape of the histogram, differences at the data extremes suggest deviations from the theoretical normal distribution.

However, the exponential distribution (blue line) and the uniform distribution (purple line) do not sufficiently approximate the histogram. Therefore, the normal distribution appears to be the best approximation, albeit with certain reservations. This data group likely requires a more detailed analysis or tests for a more accurate identification of the appropriate distribution model.

Histogram with Densities - Group 2 (Conventional)



```
[16]: # Function to generate and compare simulated data
    compare_histograms_improved <- function(data, simulated_data, group,
    distribution_type) {

    # Define common breaks for both histograms
    breaks_common <- seq(
        floor(min(c(data, simulated_data))),
        ceiling(max(c(data, simulated_data))),
        length.out = 15
    )

    # Setting layout for vertically stacked plots</pre>
```

```
layout(matrix(c(1, 2), nrow = 2, ncol = 1, byrow = TRUE)) # Plots stacked
\rightarrowvertically
# Histogram of original data
hist(
  data,
  breaks = breaks common,
  freq = FALSE,
  main = paste("Original Data -", group),
  xlab = "Time",
  ylab = "Density",
  col = "lightgray", # Fill color
  border = "darkgray", # Border color
  ylim = c(0, max(density(data)$y, density(simulated_data)$y))
lines(density(data), col = "black", lwd = 2) # Density of original data
# Histogram of simulated data
hist(
  simulated_data,
  breaks = breaks common,
  freq = FALSE,
  main = paste("Simulated Data -", group),
  xlab = "Time",
  ylab = "Density",
  col = if (distribution_type == "Exponential") "lightgreen" else "skyblue",
  border = if (distribution_type == "Exponential") "darkgreen" else⊔
⇔"steelblue".
  ylim = c(0, max(density(data)$y, density(simulated_data)$y))
lines(density(simulated_data), col = if (distribution_type == "Exponential")__
→"green" else "blue", lwd = 2, lty = 2) # Density of simulated data
# Resetting layout
layout(1)
# Numerical summary
cat("\n")
cat(bold(paste0("Numerical Summary for ", group, ":")), "\n")
cat("\n")
cat(bold("Original Data:\n"))
cat("- Minimum:", round(min(data), 2), "\n")
cat("- First Quartile:", round(quantile(data, 0.25), 2), "\n")
cat("- Median:", round(median(data), 2), "\n")
cat("- Mean:", round(mean(data), 2), "\n")
cat("- Third Quartile:", round(quantile(data, 0.75), 2), "\n")
```

```
cat("- Maximum:", round(max(data), 2), "\n")
  cat("- Standard Deviation:", round(sd(data), 2), "\n")
  cat("\n")
  cat(bold("Simulated Data:\n"))
  cat("- Minimum:", round(min(simulated_data), 2), "\n")
  cat("- First Quartile:", round(quantile(simulated_data, 0.25), 2), "\n")
  cat("- Median:", round(median(simulated_data), 2), "\n")
  cat("- Mean:", round(mean(simulated data), 2), "\n")
  cat("- Third Quartile:", round(quantile(simulated_data, 0.75), 2), "\n")
  cat("- Maximum:", round(max(simulated data), 2), "\n")
  cat("- Standard Deviation:", round(sd(simulated_data), 2), "\n")
  cat("\n")
}
# Generating simulated data and comparison for Group 1 (Modified) - Exponential \Box
 \rightarrow distribution
lambda1 <- 1 / mean(group1$Time)</pre>
simulated_group1 <- rexp(100, rate = lambda1)</pre>
compare histograms improved (group1$Time, simulated group1, "Group 111
→(Modified)", "Exponential")
# Generating simulated data and comparison for Group 2 (Conventional) - Normal
\hookrightarrow distribution
mean2 <- mean(group2$Time)</pre>
sd2 <- sd(group2$Time)</pre>
simulated_group2 <- rnorm(100, mean = mean2, sd = sd2)</pre>
compare histograms improved(group2$Time, simulated group2, "Group 211
⇔(Conventional)", "Normal")
# Discussion
cat(bold("Discussion:"))
cat("\nFor Group 1 (Modified), an exponential simulation was used. The ⊔
 histogram of the simulated data shows a similar skewness and shape to the
 ⇔original data, with the simulated data having a slightly sharper ⊔
 ⇔distribution.\n")
cat("The numerical summary confirms that the mean and median are close, but the _{\sqcup}
 ⇒standard deviation may differ slightly.\n\n")
cat("For Group 2 (Conventional), a normal simulation was used. The histogram of \Box
 \hookrightarrowthe simulated data aligns well with the original data, especially around the \sqcup
 →mean.\n")
cat("The numerical summary shows that the mean and median of the simulated data__
 ⇔are very similar to the original data, confirming the suitability of the⊔
 →normal distribution for this group.\n")
```

Numerical Summary for Group 1 (Modified):

Original Data:

- Minimum: 68

- First Quartile: 75.5

- Median: 106 - Mean: 125.29

- Third Quartile: 176

- Maximum: 210

- Standard Deviation: 56.6

Simulated Data:

- Minimum: 0.58

- First Quartile: 39.35

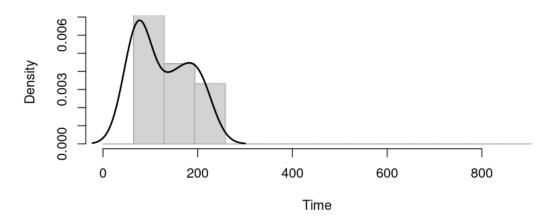
- Median: 106.21 - Mean: 131.01

- Third Quartile: 184.02

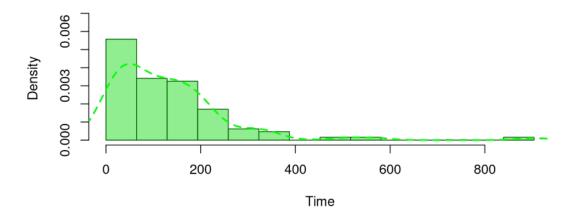
- Maximum: 903.44

- Standard Deviation: 130.12

Original Data - Group 1 (Modified)



Simulated Data - Group 1 (Modified)



Numerical Summary for Group 2 (Conventional):

Original Data:

- Minimum: 130

- First Quartile: 152.75

- Median: 235 - Mean: 224.14

- Third Quartile: 300

- Maximum: 300

- Standard Deviation: 70.55

Simulated Data:

- Minimum: 79.29

- First Quartile: 177.12

- Median: 212.97 - Mean: 222.16

- Third Quartile: 263.09

- Maximum: 452.79

- Standard Deviation: 69.55

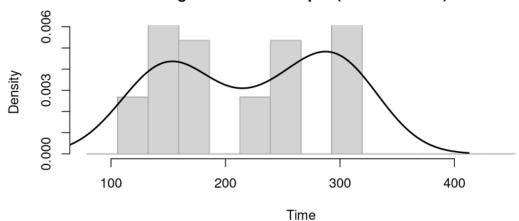
Discussion:

For Group 1 (Modified), an exponential simulation was used. The histogram of the simulated data shows a similar skewness and shape to the original data, with the simulated data having a slightly sharper distribution.

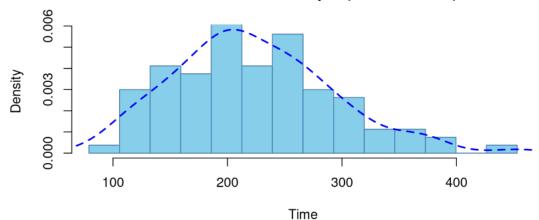
The numerical summary confirms that the mean and median are close, but the standard deviation may differ slightly.

For Group 2 (Conventional), a normal simulation was used. The histogram of the simulated data aligns well with the original data, especially around the mean. The numerical summary shows that the mean and median of the simulated data are very similar to the original data, confirming the suitability of the normal distribution for this group.

Original Data - Group 2 (Conventional)



Simulated Data - Group 2 (Conventional)



```
[17]: cat("\n")
# Function to calculate and display the 95% confidence interval for the mean
calculate_confidence_interval <- function(data, group_name) {
    # Using t.test to obtain the confidence interval
    test <- t.test(data, conf.level = 0.95)

# Displaying the results
cat(bold("Group:", group_name))
cat("\n")
cat("\n")
cat("95% Confidence Interval for the Mean:",
    round(test$conf.int[1], 2), "-", round(test$conf.int[2], 2), "\n\n")
}</pre>
```

```
# Calculating the confidence interval for Group 1 (Modified)
      calculate_confidence_interval(group1$Time, "Group 1 (Modified)")
      # Calculating the confidence interval for Group 2 (Conventional)
      calculate_confidence_interval(group2$Time, "Group 2 (Conventional)")
     Group: Group 1 (Modified)
     95% Confidence Interval for the Mean: 92.61 - 157.96
     Group: Group 2 (Conventional)
     95% Confidence Interval for the Mean: 183.41 - 264.88
[18]: cat("\n")
      # Function to calculate and display the 95% confidence interval for the mean
      calculate_confidence_interval <- function(data, group_name) {</pre>
        # Using t.test to obtain the confidence interval
        test <- t.test(data, conf.level = 0.95)</pre>
        # Displaying the results
        cat(bold("Group:", group_name))
        cat("\n")
        cat("95% Confidence Interval for the Mean:",
            round(test$conf.int[1], 2), "-", round(test$conf.int[2], 2), "\n\n")
      }
      # Calculating the confidence interval for Group 1 (Modified)
      calculate_confidence_interval(group1$Time, "Group 1 (Modified)")
      # Calculating the confidence interval for Group 2 (Conventional)
      calculate confidence interval(group2$Time, "Group 2 (Conventional)")
      cat("\n")
      # Splitting data into groups
      group1 <- subset(case0402, Treatmt == "Modified")</pre>
      group2 <- subset(case0402, Treatmt == "Conventional")</pre>
      # Defining the value of K
      K <- 19 # the day of my birthday
      # Function to perform a two-sample t-test and display the results
      perform_group_comparison <- function(data1, data2) {</pre>
        # Performing a two-sample t-test
        test <- t.test(data1, data2, alternative = "two.sided", var.equal = TRUE)</pre>
```

Displaying the results

```
cat(bold("Group Comparison (Modified vs. Conventional):"))
 cat("\n")
 cat("- Null Hypothesis HO: The means of both groups are equal.\n")
  cat("- Alternative Hypothesis Ha: The means of both groups are different.\n")
  cat("- Test Statistic (t):", round(test$statistic, 3), "\n")
  cat("- P-value:", round(test$p.value, 4), "\n")
 cat("\n")
  # Decision on rejecting HO:
 cat(bold("Decision on Rejecting HO"))
 cat("\n")
 if (test$p.value < 0.05) {</pre>
    cat("We reject HO at the 5% significance level. The means of the groups are

different.\n")
 } else {
    ⇒groups are not different.\n")
 return(test)
}
# Visualization of data using boxplot with consistent colors
ggplot(case0402, aes(x = Treatmt, y = Time, fill = Treatmt)) +
 geom_boxplot(color = c("darkgreen", "steelblue"), lwd = 1.2) + # Borders⊔
 ⇔according to colors
 scale_fill_manual(values = c("Modified" = "lightgreen", "Conventional" = 
 ⇔"skyblue")) + # Consistent fills
 labs(title = "Boxplot of Time by Groups",
       x = "Group",
      y = "Time") +
 theme_minimal()
# Performing the test for both groups
t_test_result <- perform_group_comparison(group1$Time, group2$Time)</pre>
# Practical interpretation of the results
cat("\n")
if (t_test_result$p.value < 0.05) {</pre>
 cat(bold("Interpretation:"))
 cat("\n")
 cat("At the 5% significance level, there is sufficient evidence to state that ⊔
 \hookrightarrowthe average time values between the 'Modified' and 'Conventional' groups_{\sqcup}

differ.\n")
} else {
 cat(bold("Interpretation:"))
 cat("\n")
```

```
Group: Group 1 (Modified)
```

95% Confidence Interval for the Mean: 92.61 - 157.96

Group: Group 2 (Conventional)

95% Confidence Interval for the Mean: 183.41 - 264.88

Group Comparison (Modified vs. Conventional):

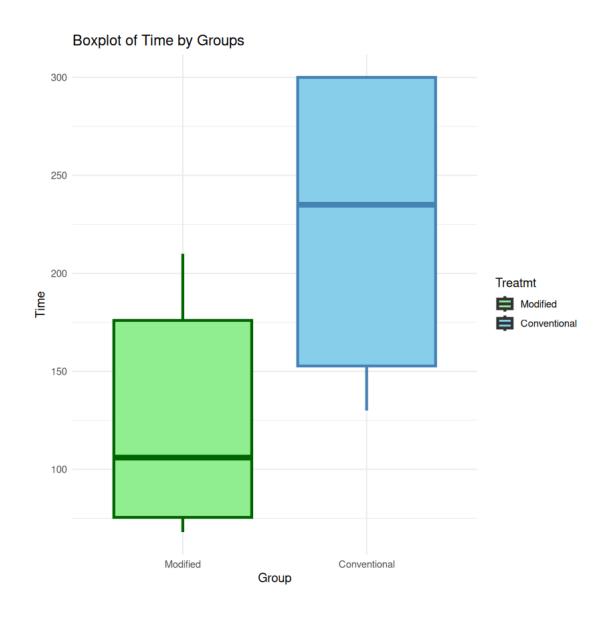
- Null Hypothesis HO: The means of both groups are equal.
- Alternative Hypothesis Ha: The means of both groups are different.
- Test Statistic (t): -4.09
- P-value: 4e-04

Decision on Rejecting HO

We reject HO at the 5% significance level. The means of the groups are different.

Interpretation:

At the 5% significance level, there is sufficient evidence to state that the average time values between the 'Modified' and 'Conventional' groups differ.



[]: