

## Assignment 6

**All screenshots that are included are from my own R Script file. You can access it with this link:** <https://github.com/kiseraidan/STAT-3010/blob/main/Assignment%206/Assignment6.R>

1. (a).

Code:

```
15 # 1. (a).
16
17 # calc the proportion of failures for each day
18 data$p <- data$Failed / data$Tested
19
20 # calc the average proportion of failures (p-bar)
21 p_bar <- mean(data$p)
22
23 # calc the standard deviation of the proportion of failures
24 p_sd <- sqrt(p_bar * (1 - p_bar) / mean(data$Tested))
25
26 # calc control limits
27 LCL <- max(0, p_bar - 3 * p_sd) # lower control limit, cannot be less than 0
28 UCL <- p_bar + 3 * p_sd # upper control limit
29
30 # print
31 cat("Centerline (p-bar):", p_bar, "\n")
32 cat("Lower Control Limit (LCL):", LCL, "\n")
33 cat("Upper Control Limit (UCL):", UCL, "\n")
```

Output:

```
Centerline (p-bar): 0.01049215
Lower Control Limit (LCL): 0.003958794
Upper Control Limit (UCL): 0.0170255
> |
```

(b).

Yes, there is a sign of an out-of-control condition in the data. On Day 21, the proportion of failed keyboards (approximately 0.0189) exceeds the upper control limit of

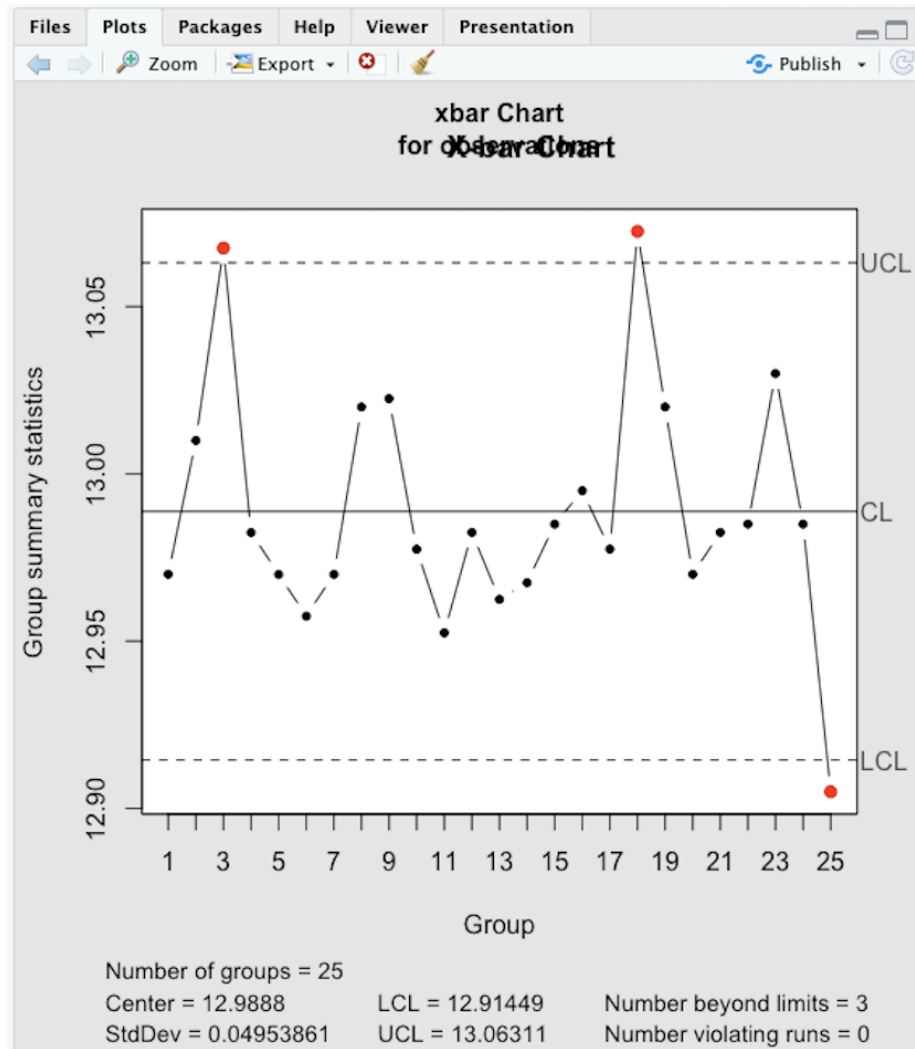
approximately 0.0170. This indicates a potential issue in the manufacturing process on that day that requires further investigation.

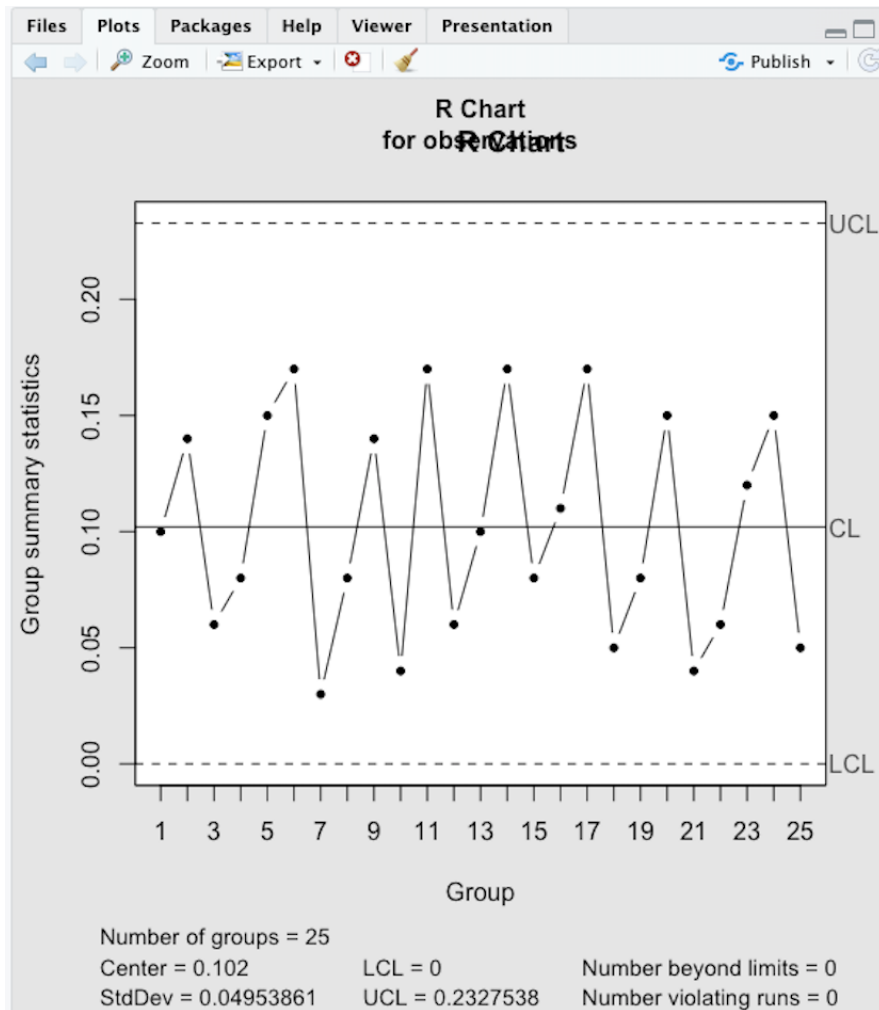
2. (a). & (b).

Code:

```
43 # 2. (a).
44
45 # prepare the matrix of observations for qcc
46 observations <- as.matrix(data[,c("x1", "x2", "x3", "x4")])
47
48 # construct the X-bar chart
49 xbar_chart <- qcc(observations, type="xbar", plot=TRUE)
50 title("X-bar Chart")
51
52 # construct the R chart
53 r_chart <- qcc(observations, type="R", plot=TRUE)
54 title("R Chart")
```

Output:





Report:

Based on these charts, there are no out-of-control signals present, implying that the production process for the bath faucet is under statistical control with respect to the threaded stem diameter.

(c).

There were no out-of-control signals present in the data. All sample means and ranges from the subgroups fell within the calculated control limits. This indicates that the production process of the bath faucet threaded stems is operating under statistical control, and there is no evidence of special cause variation.

3. (a).

Code:

```
56 # 3. (a).
57
58 # given values
59 sigma <- 5 # Population standard deviation
60 k <- 1 # Units from the mean
61
62 # sample sizes
63 n_values <- c(2, 5, 10, 30)
64
65 # function to calculate probability
66 calculate_probability <- function(n) {
67   sigma_x_bar <- sigma / sqrt(n) # standard error
68   z1 <- (k) / sigma_x_bar # z-score for mu + k
69   z2 <- (-k) / sigma_x_bar # z-score for mu - k
70   # area under the curve between z1 and z2
71   probability <- pnorm(z1) - pnorm(z2)
72   return(probability)
73 }
74
75 # calc and print probabilities for each sample size
76 probabilities <- sapply(n_values, calculate_probability)
77 names(probabilities) <- n_values
78 probabilities
```

Output:

```
> probabilities
      2      5     10     30
0.2227026 0.3452792 0.4729107 0.7266783
> |
```

(b).

Code:

```

80 # 3. (b).
81
82 # updated sample sizes
83 n_values_updated <- c(50, 100, 1000)
84
85 # use the same function as before for calculating probability
86 # calculate_probability is already defined from the previous part
87
88 # calc and print probabilities for the updated sample sizes
89 probabilities_updated <- sapply(n_values_updated, calculate_probability)
90 names(probabilities_updated) <- n_values_updated
91 probabilities_updated

```

Output:

```

> probabilities_updated
      50      100     1000
0.8427008 0.9544997 1.0000000
> |

```

(c).

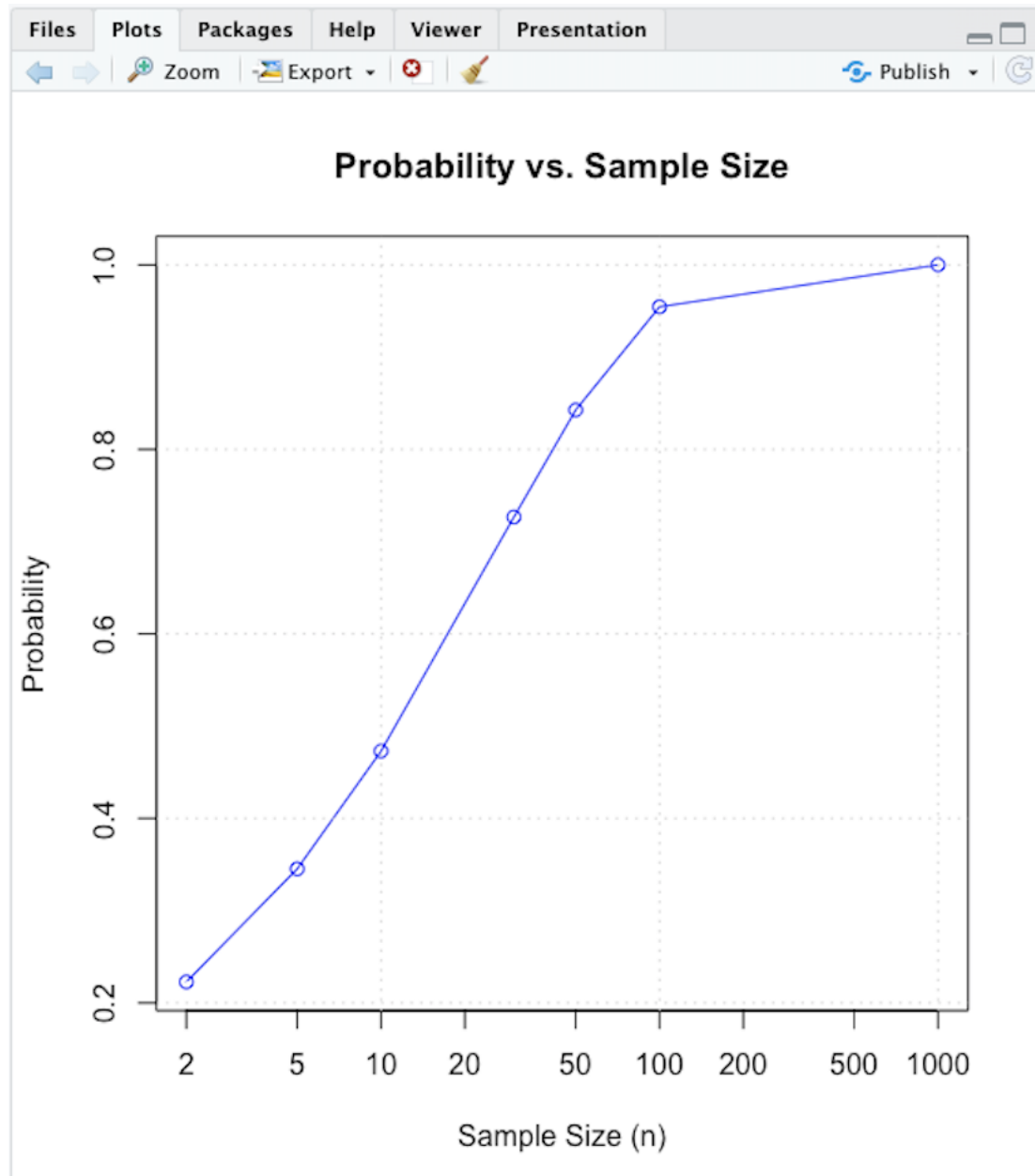
Code:

```

93 # 3. (c).
94
95 # combine all sample sizes and their corresponding probabilities
96 all_n_values <- c(2, 5, 10, 30, 50, 100, 1000)
97 all_probabilities <- c(probabilities, probabilities_updated)
98
99 # plotting
100 plot(all_n_values, all_probabilities, type="o", col="blue",
101       xlab="Sample Size (n)", ylab="Probability",
102       main="Probability vs. Sample Size", log="x")
103 grid()

```

Output:



Report:

- The probability increases with the sample size, showing a clear trend that as you collect more data, the precision of the estimate of the population mean improves. For smaller sample sizes, the probability is relatively low, indicating more variability around the population mean. As the sample size grows, this variability decreases, and the probability that the sample mean is close to the population mean increases significantly.
- For very large sample sizes (e.g.,  $n=1000$ ), the probability approaches 1, indicating that with enough data, the sample mean is almost certain to be within 1 unit of the population mean. This is a direct consequence of the Central Limit Theorem, which states that the

distribution of the sample mean becomes increasingly normal (and narrower) as the sample size increases, regardless of the population's distribution.

- The x-axis is on a logarithmic scale to accommodate the wide range of sample sizes and illustrate how the probability increases sharply with small increases in sample size initially, and then more gradually as the sample size becomes very large.