Project 2 STAT 355

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1 Part 1

1000 random samples of size 40 were generated from normal distribution with mean $\mu = 3$ and standard deviation $\sigma = 2$.

1.1 Output

Actual:	Theoretical
μ : 3	μ : 3
$E(\overline{X}): 2.9903$	$E(\overline{X})$: 2.9903
σ : 2	σ : 2
$\sigma_{\overline{X}}$: 0.311	$\sigma_{\overline{X}}$: 0.316

1.2 Distribution

Distribution of the data was plotted with a histogram using ggplot2 in Figure 1.

```
# plot a histogram
ggplot() + aes(generatedData) +
    geom_histogram(binwidth=1, color="black", fill="white") + labs(y="Count")
```

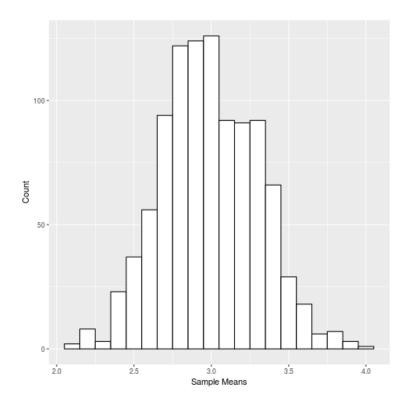


Figure 1: Histogram of the Generated Data

```
# main.R
# This file contains the implementation of the functions in the Project 2
# NOTE: THIS SCRIPT WAS COMPILED ON A LINUX MACHINE - SOME STATEMENTS MAY THROW
# WARNINGS OR ERRORS IN OTHER SYSTEMS
library(ggplot2) # for generating high quality plots
outputTemplate <- "\\subsection{Output}</pre>
   \\begin{table}[h]
       \\centering
       \\begin{tabular*}{200pt}{@{\\extracolsep{\\fill}} c c}
       \\textbf{Actual}: & \\textbf{Theoretical}: \\\\
       $\\mu$: %1.0f & $\\mu$: %1.0f \\\
       E(\\langle X\}\): %.4f & E(\\langle X\}\): %.4f \
       $\\sigma$: %1.0f & $\\sigma$: %1.0f \\\
       \star \ textsubscript
(\ ): %.3f & \ textsubscript
(\ ): %.3f \\
       \\end{tabular*}
   \\end{table}
    ----- Part 1 -----
# initialize parameters for normal distribution
N <- 40 # size
mu <- 3 # mean
sigma <- 2 # standard deviation
sampMeans <- c() # initialize empty array</pre>
# generate 1000 samples
for (i in 1:1000){
   generatedData <- rnorm(N, mu, sigma)</pre>
   # store the sample means in vector
   sampMeans[i] = mean(generatedData)
# save output
sink("part1.tex", append=FALSE, split=FALSE)
cat(
   sprintf(
       outputTemplate,
       mu, mu,
       mean(sampMeans), mean(sampMeans),
       sigma, sigma,
       sd(sampMeans), sigma/sqrt(N)
)
sink()
png(filename="figures/hist1.png")
# plot a histogram of the data
ggplot() + aes(sampMeans) +
   geom_histogram(binwidth=0.1, color="black", fill="white") +
   labs(y="Count", x="Sample Means")
dev.off()
# ------ Part 2 ------
# initialize parameters for binomial distribution
```

```
N <- 15
n <- 10
p <- 0.15
sampMeans <- c()</pre>
for (i in 1:1000){
    generatedData <- rbinom(N, n, p)</pre>
    sampMeans[i] = mean(generatedData)
# save output
sink("part2.out", append=FALSE, split=FALSE)
# save output
sink("part2.tex", append=FALSE, split=FALSE)
cat(
    sprintf(
        outputTemplate,
        n*p, n*p,
        mean(sampMeans), mean(sampMeans),
        n*p*(1-p), n*p*(1-p),
        sd(sampMeans), sd(sampMeans)
)
sink()
png(filename="figures/hist2.png")
\# plot a histogram of the data
ggplot() + aes(sampMeans) +
    geom_histogram(binwidth=0.2, color="black", fill="white") +
    labs(y="Count", x="Sample Means")
dev.off()
# ------ Part 3 ------
\hbox{\tt\# initialize parameters for binomial distribution}\\
N <- 120
n <- 10
p <- 0.15
sampMeans <- c()</pre>
for (i in 1:1000){
    generatedData <- rbinom(N, n, p)</pre>
    sampMeans[i] = mean(generatedData)
# save output
sink("part3.out", append=FALSE, split=FALSE)
# save output
sink("part3.tex", append=FALSE, split=FALSE)
cat(
    sprintf(
        outputTemplate,
       n*p, n*p,
        mean(sampMeans), mean(sampMeans),
        n*p*(1-p), n*p*(1-p),
        sd(sampMeans), sd(sampMeans)
)
sink()
png(filename="figures/hist3.png")
# plot a histogram of the data
```

```
ggplot() + aes(sampMeans) +
    geom_histogram(binwidth=0.3, color="black", fill="white") +
    labs(y="Count", x="Sample Means")

dev.off()
```