

# MTH 441 Lab Assignment 2

Viral Chitlangia

## P1

```
data.File1 <- read.csv("rocket_propellant.csv")
model1 <- lm(Shear.Strength..psi. ~ Age.of.Propellant..weeks., data = data.File1)
model1$coefficients
```

```
##              (Intercept) Age.of.Propellant..weeks.
##              2627.82236                -37.15359
```

## P2

```
data.File2 <- read.csv("delivery_times.csv")
model2 <- lm(Delivery_Time ~ Number_of_Cases_x1 + Distance_x2, data = data.File2)
model2$coefficients
```

```
##          (Intercept) Number_of_Cases_x1      Distance_x2
##          2.76356503      1.11355896      0.02421374
```

```
X <- cbind(numeric(length(data.File1$Observation)) + 1, data.File1)
colnames(X)[1] <- "One"
X <- X[, c(1, 4)]
X <- as.matrix(X, nrow = length(X$'One'), ncol = 2)
```

## P3

```
# 1
variance <- sum(model1$residuals^2)/(length(data.File1[,2]) - 2)
variance
```

```
## [1] 9236.381
```

```
# 2
varVec <- variance * solve(t(X) %*% X)
Var <- diag(varVec)
t <- model1$coefficients/Var
t
```

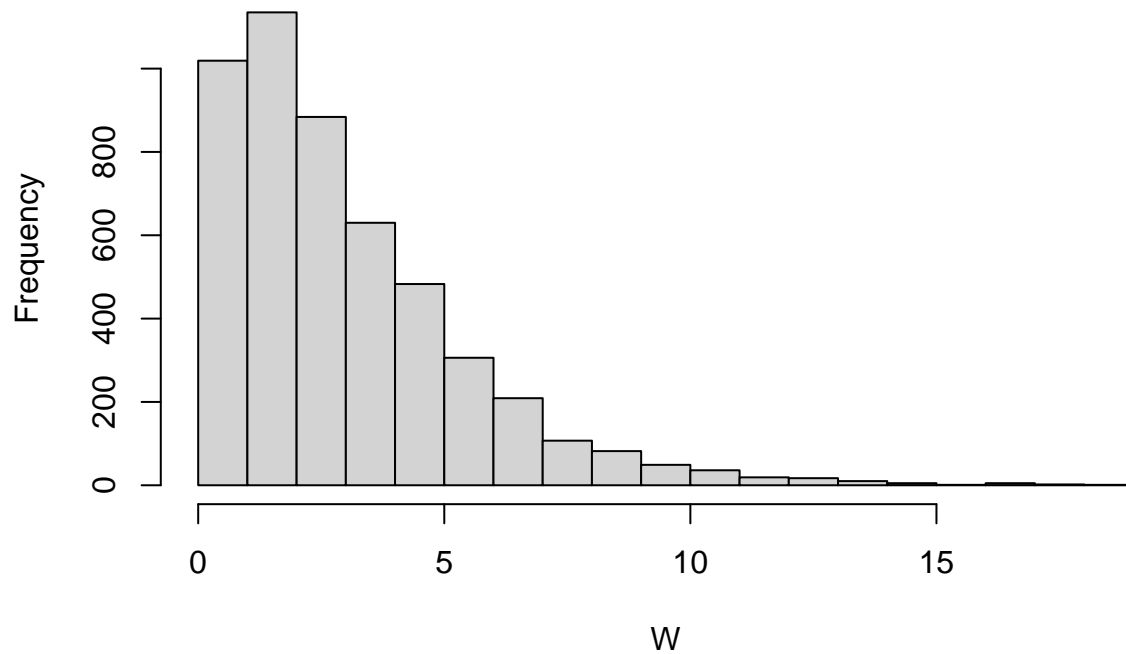
```
##              (Intercept) Age.of.Propellant..weeks.
##              1.346070                -4.451165
```

## P4

```
# 1
W <- vector()
for (i in 1:5000) {
  W <- append(W, rnorm(1)^2 + rnorm(1)^2 + rnorm(1)^2)
}
```

```
# 2
hist(W)
```

## Histogram of W



```
m <- mean(W)
v <- var(W)

# 3
theoretical.mean <- 3
theoretical.variance <- 6
```

```
# 4
m - theoretical.mean
```

```
## [1] -0.01272887
```

```
v - theoretical.variance
```

```
## [1] -0.03559363
```

## P5

```
# 1
X <- matrix(rnorm(40, 5, 3), nrow = 8, ncol = 5)
Px <- X %*% solve(t(X) %*% X) %*% t(X)
if (norm(Px - Px%*%Px, type = "2") < 1e-6) {
  print("Px is Idempotent")
}
```

```
## [1] "Px is Idempotent"
```

```
# 2
S <- NULL
for (i in 1:5000) {
  Y <- NULL
  for (j in 1:8) {
```

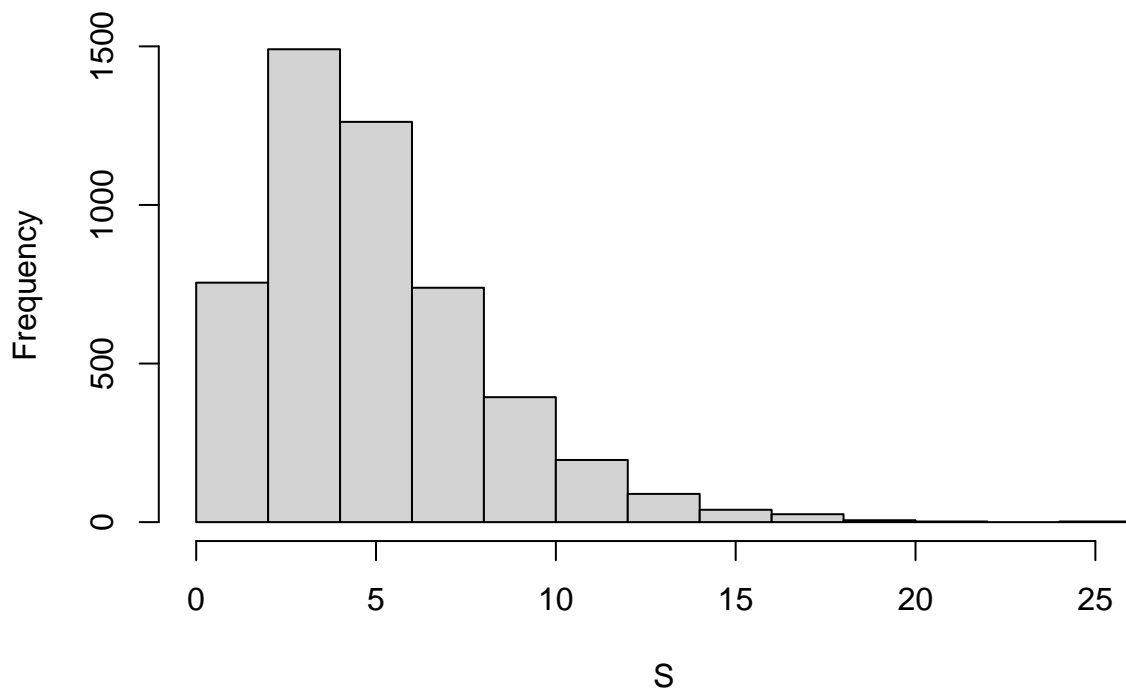
```

    Y <- c(Y, rnorm(1))
  }
  Y <- as.matrix(Y, 1, 8)
  S <- c(S, t(Y) %*% Px %*% Y)
}

# 3
hist(S)

```

**Histogram of S**



```

m <- mean(S)
v <- var(S)
m

```

```
## [1] 4.965037
```

```
v
```

```
## [1] 9.686088
```

```

# 4
theoretical.mean <- 5
theoretical.variance <- 10

```

```

# 5
m - theoretical.mean

```

```
## [1] -0.0349634
```

```
v - theoretical.variance
```

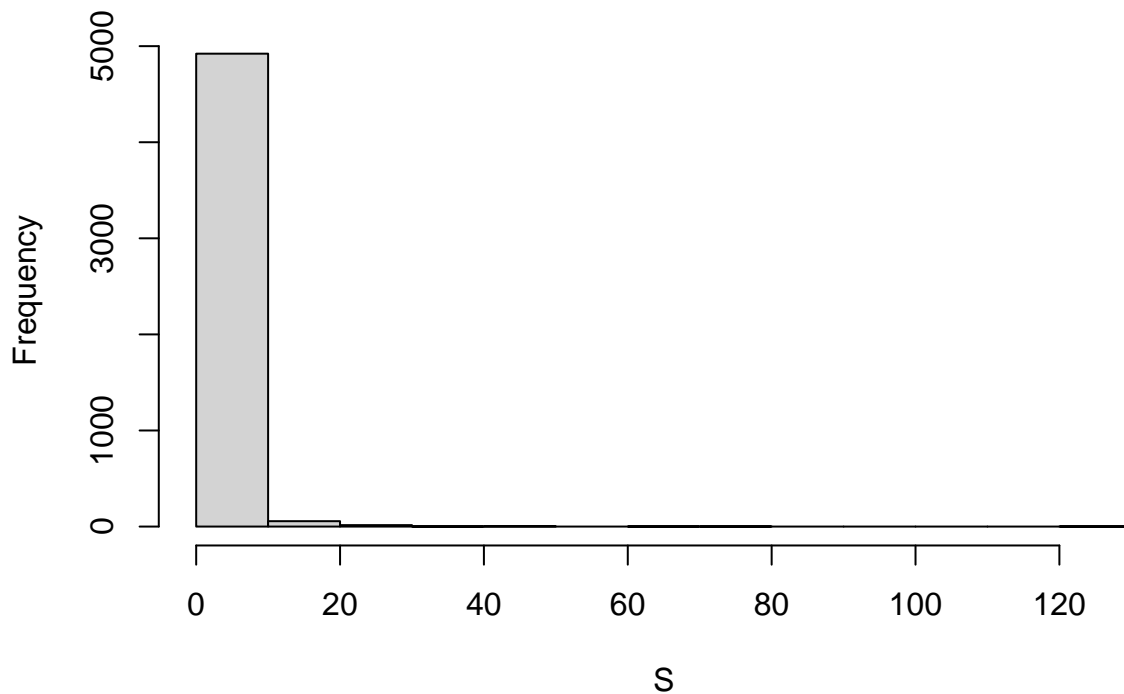
```
## [1] -0.3139115
```

## P6

```
# 2
P.X.1 = Px
P.X.2 = diag(1, 8, 8) - P.X.1
df1 <- qr(P.X.2)$rank
df2 <- qr(P.X.1)$rank
S <- NULL
for (i in 1:5000) {
  Y <- NULL
  for (j in 1:8) {
    Y <- c(Y, rnorm(1))
  }
  Y <- as.matrix(Y, 1, 8)
  S <- c(S, as.vector(t(Y) %*% P.X.2 %*% Y / df1) / (t(Y) %*% P.X.1 %*% Y / df2))
}

# 3
hist(S)
```

Histogram of S



```
m <- mean(S)
v <- var(S)
m

## [1] 1.729102
v

## [1] 12.34148
```

```

# 4
theoretical.mean <- 5/3
theoretical.variance <- (2 * (5^2) * (3 + 5 - 2)) / (3 * (5 - 2)^2 * (5 - 4))
theoretical.mean

## [1] 1.666667
theoretical.variance

## [1] 11.11111
# 5
m - theoretical.mean

## [1] 0.06243544
v - theoretical.variance

## [1] 1.230372

```