

STOR 565 Spring 2018 Homework 1

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Exercise 1. (5 pt) Using the `c`, `rep` or `seq` commands, create the following 6 vectors:

`x1 = (2, .5, 4, 2);`

`x2 = (2, .5, 4, 2, 1, 1, 1, 1);`

`x3 = (1, 0, -1, -2);`

`x4 = ("Hello", " ", "World", "!", "Hello World!");`

Hint. Try `paste` function.

`x5 = (TRUE, TRUE, NA, FALSE);`

Hint. Check `?NA` and `class(NA)` to learn more about the missing value object `NA`.

`x6 = (1, 2, 1, 2, 1, 1, 2, 2).`

```
x1 <- c(2, .5, 4, 2);
x2 <- c(2, .5, 4, 2, rep(1, 4));
x3 <- seq(1, -2, -1);
x4 <- paste("Hello", " ", "World", "!", "Hello World!", sep=" ", collapse = NULL);
x5 <- c(rep(TRUE, 2), NA, TRUE);
x6 <- c(1, 2, 1, 2, rep(1, 2), rep(2, 2));
x1
```

```
## [1] 2.0 0.5 4.0 2.0
```

```
x2
```

```
## [1] 2.0 0.5 4.0 2.0 1.0 1.0 1.0 1.0
```

```
x3
```

```
## [1] 1 0 -1 -2
```

```
x4
```

```
## [1] "Hello World ! Hello World!"
```

```
x5
```

```
## [1] TRUE TRUE NA TRUE
```

```
x6
```

```
## [1] 1 2 1 2 1 1 2 2
```

Exercise 2. (5 pt) Using `matrix`, `rbind` and `cbind`, create

$$\mathbf{X} = \begin{pmatrix} 1 & 2 & 3 & 4 \\ 1 & 0 & -1 & -2 \\ 2 & .5 & 4 & 2 \\ 1 & 1 & 1 & 1 \end{pmatrix}$$

```
A = matrix(c(1,1,2,1), 4, 1)
B = matrix(c(2,0,.5,1), 4, 1)
C = matrix(c(3, -1, 4, 1), 4, 1)
D = matrix(c(4, -2, 2, 1), 4, 1)
```

```
X <- cbind(A, B, C, D)
X
```

```
##      [,1] [,2] [,3] [,4]
## [1,]    1  2.0    3    4
## [2,]    1  0.0   -1   -2
## [3,]    2  0.5    4    2
## [4,]    1  1.0    1    1
```

```
students <- data.frame( id      = c("001", "002", "003"), # ids are characters
                        score_A = c(95, 97, 90),          # scores are numericss
                        score_B = c(80, 75, 84))

students
```

Exercise 3. (5 pt) Applying the conditional selection technique (without using `subset`), extract the record of 003 in `students`.

```
students3 <- students[3,]
students3

##      id score_A score_B
## 3 003        90       84
```

One can also create a matrix or a legitimate list first and then convert it into a `data.frame` as follows.

```
scores <- matrix(c(95, 97, 90, 80, 75, 84), 3, 2)
scores <- data.frame(scores)
colnames(scores) <- c("score_A", "score_B")
id <- c("001", "002", "003")
students1 <- cbind(id, scores)
students2 <- data.frame( list( id      = c("001", "002", "003"),
                              score_A = c(95, 97, 90),
                              score_B = c(80, 75, 84))
                        )
```

Exercise 4. (10 pt) Create a `data.frame` object to display the calendar for Jan 2018 as follows.

```
## Sun Mon Tue Wed Thu Fri Sat
##      NY   2   3   4   5   6
##    7   8   9  10  11  12  13
##  MLK  15  16  17  18  19  20
##   21  22  23  24  25  26  27
##   28  29  30  31
calendar <- matrix(c(" ", 7, "MLK", 21, 28, "NY", 8,
                    15, 22, 29, 2, 9, 16, 23, 30, 3,
                    10, 17, 24, 31, 4, 11, 18, 25, " ",
                    5, 12, 19, 26, " ", 6, 13, 20, 27, " "), 5, 7)
calendar <- data.frame(calendar)
colnames(calendar) <- c("Sun", "Mon", "Tue", "Wed", "Thu", "Fri", "Sat")
print(calendar, row.names=FALSE)

## Sun Mon Tue Wed Thu Fri Sat
##      NY   2   3   4   5   6
##    7   8   9  10  11  12  13
##  MLK  15  16  17  18  19  20
##   21  22  23  24  25  26  27
##   28  29  30  31
```

Factors

Factor is a special data structure in **R** in representing categorical variables and facilitating the data labels and subgroups. It's basically a character vector that keeps track of its distinct values called levels. Consider the longitudinal layout of the previous score table.

```
id      <- rep(c("001","002","003"), 2)
subj    <- rep(c("A","B"), each = 3)
score   <- c(95, 97, 90, 80, 75, 84)
students3 <- data.frame(id, subj, score) # try cbind(id, subj, score) to see the difference

# students3$id and students3$subj are automatically formatted as factors
class(students3$id)
levels(students3$id)

class(students3$subj)
levels(students3$subj)

# combine student 003 with 002 via level rename
students4 <- students3 # work on a copy in case of direct modification of students3
levels(students4$id)[3] <- "002"
levels(students4$id)
students4
```

Exercise 5. (5 pt) Create a factor variable `grade` in `students3`, where the `score` variable is divided into [90,100], [80,89) and [0,79) corresponding to A, B and C in `grade` respectively.

```
grades <- cut(students3$score, breaks = c(100, 89, 79, 0), labels=c("C","B","A"))
grades
```

```
## [1] A A A B C B
## Levels: C B A
```

```
students3 <- data.frame(id, subj, score, grades)
students3
```

```
##   id subj score grades
## 1 001   A    95      A
## 2 002   A    97      A
## 3 003   A    90      A
## 4 001   B    80      B
## 5 002   B    75      C
## 6 003   B    84      B
```

Operations and Functions

Exercise 6. (5 pt) Without using the `var` and `scale` functions, compute the sample covariance `X.var` of the data matrix `X` as in **Exercise 2**.

```
k <- ncol(X) #number of variables
n <- nrow(X) #number of subjects
X_mean <- matrix(data=1,nrow=n) %*% cbind(mean(A),mean(B),mean(C),mean(D)) #creates means for each column
H <- X - X_mean #creates a difference matrix
X.var <- (n-1)^-1 * t(H) %*% H #creates covariance matrix
X.var
```

```
##           [,1]           [,2]           [,3]           [,4]
```

```
## [1,] 0.250 -0.1250000 0.7500000 0.250
## [2,] -0.125 0.7291667 0.9583333 1.875
## [3,] 0.750 0.9583333 4.9166667 4.750
## [4,] 0.250 1.8750000 4.7500000 6.250
```

```
cov(X)
```

```
##      [,1]      [,2]      [,3] [,4]
## [1,] 0.250 -0.1250000 0.7500000 0.250
## [2,] -0.125 0.7291667 0.9583333 1.875
## [3,] 0.750 0.9583333 4.9166667 4.750
## [4,] 0.250 1.8750000 4.7500000 6.250
```

Exercise 7. (10 pt) Create a variable `score.mean` in `students3`, taking value as the mean score among students who have the same `subj` value.

```
A_mean <- mean(score[students3$subj == "A"])
B_mean <- mean(score[students3$subj == "B"])
score.mean <- c(A_mean, B_mean)
score.mean
```

```
## [1] 94.00000 79.66667
```

Writing your own functions

Exercise 8. (15 pt) Write a function `bisect(f, lower, upper, tol = 1e-6)` to find the root of the univariate function `f` on the interval `[lower, upper]` with precision tolerance $\leq \text{tol}$ (defaulted to be 10^{-6}) via bisection, which returns a list consisting of `root`, `f.root` (`f` evaluated at `root`), `iter` (number of iterations) and `estim.prec` (estimated precision). Apply it to the function

$$f(x) = x^3 - x - 1$$

on `[1, 2]` with precision tolerance 10^{-6} . Compare it with the built-in function `uniroot`.

```
bisection <- function(f, a, b, n = 1000, tol = 1e-6) {
  if (!(f(a) < 0) && (f(b) > 0)) {
    stop('signs of f(a) and f(b) differ')
  } else if ((f(a) > 0) && (f(b) < 0)) {
    stop('signs of f(a) and f(b) differ')
  }

  for (i in 1:n) {
    c <- (a + b) / 2 # Calculate midpoint
    if ((f(c) == 0) || ((b - a) / 2) < tol) {
      return(c)
    }
    ifelse(sign(f(c)) == sign(f(a)),
           a <- c,
           b <- c)
  }
  print('Too many iterations')
}

f <- function(x) {x^3 - x - 1} #function
a <- 1 #lower
b <- 2 #upper
tol <- 1e-6 #tolerance
```

	V1	V2	V3	V4
V1		-0.29	0.68	0.2
V2			0.51	0.88
V3				0.86
V4				

Table 1

```
n <- 10000 #iterations
bisection(f, a, b, n, tol)
```

```
## [1] 1.324718
```

Exercise 9. (15 pt) Without using `cor`, compute the sample correlation matrix `X.cor` from `X.var` in **Exercise 6**. Output `X.cor` to a text file “`X_cor.txt`” which displays as in Table 1. Then input “`X_cor.txt`” in **R** and reproduce the correlation matrix `X.cor1`.

Hint. 1) Functions `round` and `lower.tri`; 2) the NA trick; 3) options `sep = "\t", col.names = NA`.

```
##      V1      V2      V3      V4
## V1  1.00 -0.29  0.68  0.20
## V2 -0.29  1.00  0.51  0.88
## V3  0.68  0.20  1.00  0.86
## V4  0.51  0.88  0.86  1.00
value <- sqrt(diag(X.var))
value1 <- solve(diag(value))
X.cor <- value1 %*% X.var %*% value1
X.cor
```

```
##      [,1]      [,2]      [,3]      [,4]
## [1,] 1.0000000 -0.2927700 0.6764814 0.2000000
## [2,] -0.2927700 1.0000000 0.5061367 0.8783101
## [3,] 0.6764814 0.5061367 1.0000000 0.8568765
## [4,] 0.2000000 0.8783101 0.8568765 1.0000000
```

#Output as text file

```
write.table(X.cor, file="X_cor.txt", sep="\t", col.names=NA, quote=FALSE)
X.cor1 <- read.table("X_cor.txt", header=FALSE, sep = "\t")
X.cor1
```

```
##      V1      V2      V3      V4
## 1 NA      V1      V2      V3
## 2 1      1 -0.29277002188456 0.676481425202546
## 3 2 -0.29277002188456      1 0.506136675356011
## 4 3 0.676481425202546 0.506136675356011      1
## 5 4      0.2 0.87831006565368 0.856876471923225
##      V5
## 1      V4
## 2      0.2
## 3 0.87831006565368
## 4 0.856876471923225
## 5      1
```

```
cov2cor(X.var)
```

```
##      [,1]      [,2]      [,3]      [,4]
## [1,] 1.0000000 -0.2927700 0.6764814 0.2000000
```

```
## [2,] -0.2927700  1.0000000 0.5061367 0.8783101
## [3,]  0.6764814  0.5061367 1.0000000 0.8568765
## [4,]  0.2000000  0.8783101 0.8568765 1.0000000
```

Exercise 10. (10 pt) `iris` is a built-in dataset in **R**. Check `?iris` for more information. Randomly divide `iris` into five subsets `iris1` to `iris5` stratified to `iris$Species` (namely, the proportion of `iris$Species` among different levels remains identical across all subsets).

```
set.seed(5)
irisnew <- split(iris, sample(1:5))
iris1 <- irisnew[1]
iris2 <- irisnew[2]
iris3 <- irisnew[3]
iris4 <- irisnew[4]
iris5 <- irisnew[5]
iris.5fold <- list(iris1, iris2, iris3, iris4, iris5)
iris.5fold
```

```
## [[1]]
## [[1]]$`1`
##      Sepal.Length Sepal.Width Petal.Length Petal.Width   Species
## 4              4.6          3.1          1.5          0.2    setosa
## 9              4.4          2.9          1.4          0.2    setosa
## 14             4.3          3.0          1.1          0.1    setosa
## 19             5.7          3.8          1.7          0.3    setosa
## 24             5.1          3.3          1.7          0.5    setosa
## 29             5.2          3.4          1.4          0.2    setosa
## 34             5.5          4.2          1.4          0.2    setosa
## 39             4.4          3.0          1.3          0.2    setosa
## 44             5.0          3.5          1.6          0.6    setosa
## 49             5.3          3.7          1.5          0.2    setosa
## 54             5.5          2.3          4.0          1.3 versicolor
## 59             6.6          2.9          4.6          1.3 versicolor
## 64             6.1          2.9          4.7          1.4 versicolor
## 69             6.2          2.2          4.5          1.5 versicolor
## 74             6.1          2.8          4.7          1.2 versicolor
## 79             6.0          2.9          4.5          1.5 versicolor
## 84             6.0          2.7          5.1          1.6 versicolor
## 89             5.6          3.0          4.1          1.3 versicolor
## 94             5.0          2.3          3.3          1.0 versicolor
## 99             5.1          2.5          3.0          1.1 versicolor
## 104            6.3          2.9          5.6          1.8  virginica
## 109            6.7          2.5          5.8          1.8  virginica
## 114            5.7          2.5          5.0          2.0  virginica
## 119            7.7          2.6          6.9          2.3  virginica
## 124            6.3          2.7          4.9          1.8  virginica
## 129            6.4          2.8          5.6          2.1  virginica
## 134            6.3          2.8          5.1          1.5  virginica
## 139            6.0          3.0          4.8          1.8  virginica
## 144            6.8          3.2          5.9          2.3  virginica
## 149            6.2          3.4          5.4          2.3  virginica
##
##
## [[2]]
## [[2]]$`2`
```

```

##      Sepal.Length Sepal.Width Petal.Length Petal.Width  Species
## 1           5.1           3.5           1.4           0.2    setosa
## 6           5.4           3.9           1.7           0.4    setosa
## 11          5.4           3.7           1.5           0.2    setosa
## 16          5.7           4.4           1.5           0.4    setosa
## 21          5.4           3.4           1.7           0.2    setosa
## 26          5.0           3.0           1.6           0.2    setosa
## 31          4.8           3.1           1.6           0.2    setosa
## 36          5.0           3.2           1.2           0.2    setosa
## 41          5.0           3.5           1.3           0.3    setosa
## 46          4.8           3.0           1.4           0.3    setosa
## 51          7.0           3.2           4.7           1.4 versicolor
## 56          5.7           2.8           4.5           1.3 versicolor
## 61          5.0           2.0           3.5           1.0 versicolor
## 66          6.7           3.1           4.4           1.4 versicolor
## 71          5.9           3.2           4.8           1.8 versicolor
## 76          6.6           3.0           4.4           1.4 versicolor
## 81          5.5           2.4           3.8           1.1 versicolor
## 86          6.0           3.4           4.5           1.6 versicolor
## 91          5.5           2.6           4.4           1.2 versicolor
## 96          5.7           3.0           4.2           1.2 versicolor
## 101         6.3           3.3           6.0           2.5  virginica
## 106         7.6           3.0           6.6           2.1  virginica
## 111         6.5           3.2           5.1           2.0  virginica
## 116         6.4           3.2           5.3           2.3  virginica
## 121         6.9           3.2           5.7           2.3  virginica
## 126         7.2           3.2           6.0           1.8  virginica
## 131         7.4           2.8           6.1           1.9  virginica
## 136         7.7           3.0           6.1           2.3  virginica
## 141         6.7           3.1           5.6           2.4  virginica
## 146         6.7           3.0           5.2           2.3  virginica
##
##
## [[3]]
## [[3]]$`3`
##      Sepal.Length Sepal.Width Petal.Length Petal.Width  Species
## 2           4.9           3.0           1.4           0.2    setosa
## 7           4.6           3.4           1.4           0.3    setosa
## 12          4.8           3.4           1.6           0.2    setosa
## 17          5.4           3.9           1.3           0.4    setosa
## 22          5.1           3.7           1.5           0.4    setosa
## 27          5.0           3.4           1.6           0.4    setosa
## 32          5.4           3.4           1.5           0.4    setosa
## 37          5.5           3.5           1.3           0.2    setosa
## 42          4.5           2.3           1.3           0.3    setosa
## 47          5.1           3.8           1.6           0.2    setosa
## 52          6.4           3.2           4.5           1.5 versicolor
## 57          6.3           3.3           4.7           1.6 versicolor
## 62          5.9           3.0           4.2           1.5 versicolor
## 67          5.6           3.0           4.5           1.5 versicolor
## 72          6.1           2.8           4.0           1.3 versicolor
## 77          6.8           2.8           4.8           1.4 versicolor
## 82          5.5           2.4           3.7           1.0 versicolor
## 87          6.7           3.1           4.7           1.5 versicolor

```

```

## 92          6.1          3.0          4.6          1.4 versicolor
## 97          5.7          2.9          4.2          1.3 versicolor
## 102         5.8          2.7          5.1          1.9 virginica
## 107         4.9          2.5          4.5          1.7 virginica
## 112         6.4          2.7          5.3          1.9 virginica
## 117         6.5          3.0          5.5          1.8 virginica
## 122         5.6          2.8          4.9          2.0 virginica
## 127         6.2          2.8          4.8          1.8 virginica
## 132         7.9          3.8          6.4          2.0 virginica
## 137         6.3          3.4          5.6          2.4 virginica
## 142         6.9          3.1          5.1          2.3 virginica
## 147         6.3          2.5          5.0          1.9 virginica
##
##
## [[4]]
## [[4]]$`4`
##      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 3              4.7          3.2          1.3          0.2      setosa
## 8              5.0          3.4          1.5          0.2      setosa
## 13             4.8          3.0          1.4          0.1      setosa
## 18             5.1          3.5          1.4          0.3      setosa
## 23             4.6          3.6          1.0          0.2      setosa
## 28             5.2          3.5          1.5          0.2      setosa
## 33             5.2          4.1          1.5          0.1      setosa
## 38             4.9          3.6          1.4          0.1      setosa
## 43             4.4          3.2          1.3          0.2      setosa
## 48             4.6          3.2          1.4          0.2      setosa
## 53             6.9          3.1          4.9          1.5      versicolor
## 58             4.9          2.4          3.3          1.0      versicolor
## 63             6.0          2.2          4.0          1.0      versicolor
## 68             5.8          2.7          4.1          1.0      versicolor
## 73             6.3          2.5          4.9          1.5      versicolor
## 78             6.7          3.0          5.0          1.7      versicolor
## 83             5.8          2.7          3.9          1.2      versicolor
## 88             6.3          2.3          4.4          1.3      versicolor
## 93             5.8          2.6          4.0          1.2      versicolor
## 98             6.2          2.9          4.3          1.3      versicolor
## 103            7.1          3.0          5.9          2.1      virginica
## 108            7.3          2.9          6.3          1.8      virginica
## 113            6.8          3.0          5.5          2.1      virginica
## 118            7.7          3.8          6.7          2.2      virginica
## 123            7.7          2.8          6.7          2.0      virginica
## 128            6.1          3.0          4.9          1.8      virginica
## 133            6.4          2.8          5.6          2.2      virginica
## 138            6.4          3.1          5.5          1.8      virginica
## 143            5.8          2.7          5.1          1.9      virginica
## 148            6.5          3.0          5.2          2.0      virginica
##
##
## [[5]]
## [[5]]$`5`
##      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 5              5.0          3.6          1.4          0.2      setosa
## 10             4.9          3.1          1.5          0.1      setosa

```


## 15	5.8	4.0	1.2	0.2	setosa
## 20	5.1	3.8	1.5	0.3	setosa
## 25	4.8	3.4	1.9	0.2	setosa
## 30	4.7	3.2	1.6	0.2	setosa
## 35	4.9	3.1	1.5	0.2	setosa
## 40	5.1	3.4	1.5	0.2	setosa
## 45	5.1	3.8	1.9	0.4	setosa
## 50	5.0	3.3	1.4	0.2	setosa
## 55	6.5	2.8	4.6	1.5	versicolor
## 60	5.2	2.7	3.9	1.4	versicolor
## 65	5.6	2.9	3.6	1.3	versicolor
## 70	5.6	2.5	3.9	1.1	versicolor
## 75	6.4	2.9	4.3	1.3	versicolor
## 80	5.7	2.6	3.5	1.0	versicolor
## 85	5.4	3.0	4.5	1.5	versicolor
## 90	5.5	2.5	4.0	1.3	versicolor
## 95	5.6	2.7	4.2	1.3	versicolor
## 100	5.7	2.8	4.1	1.3	versicolor
## 105	6.5	3.0	5.8	2.2	virginica
## 110	7.2	3.6	6.1	2.5	virginica
## 115	5.8	2.8	5.1	2.4	virginica
## 120	6.0	2.2	5.0	1.5	virginica
## 125	6.7	3.3	5.7	2.1	virginica
## 130	7.2	3.0	5.8	1.6	virginica
## 135	6.1	2.6	5.6	1.4	virginica
## 140	6.9	3.1	5.4	2.1	virginica
## 145	6.7	3.3	5.7	2.5	virginica
## 150	5.9	3.0	5.1	1.8	virginica

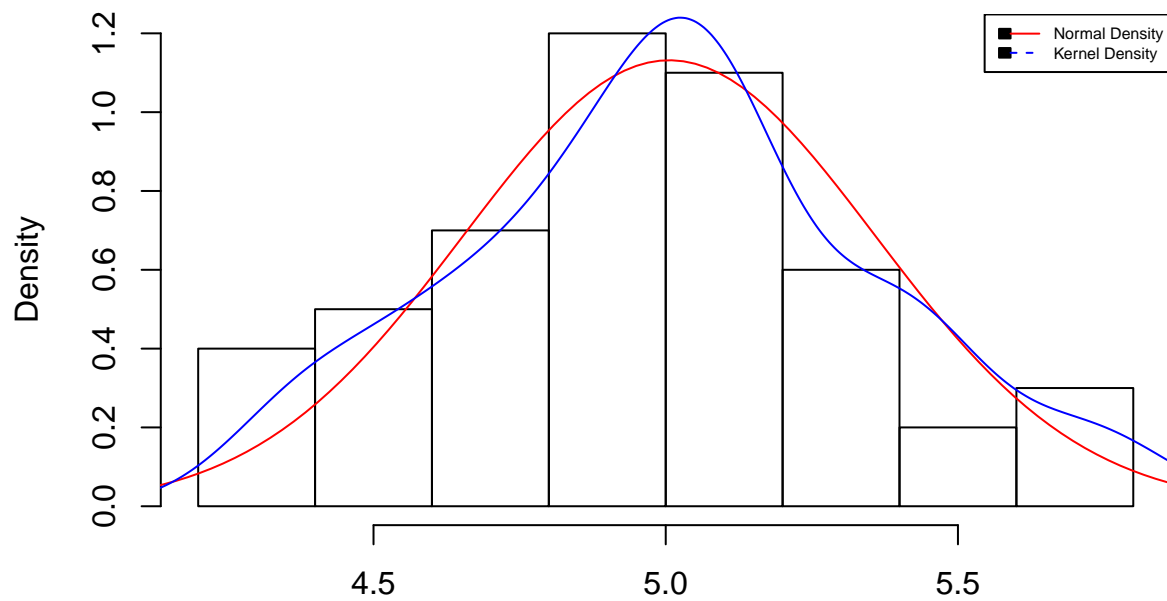
Exercise 11. (15 pt) Reproduce the code that generates the following plot about Sepal.Length in iris.

```
knitr::include_graphics("Iris_Sepal_Length.pdf")
```

Hint. 1) Most decorations are based on defaults in `hist` with `ylim = c(0, 1.3)`; 2) let `h` be the object resulted by `hist` and set `xlim = range(h$breaks)`; 3) set `cex = 0.5` in `legend`; 4) use `curve` function in plotting the normal density with specified parameters.

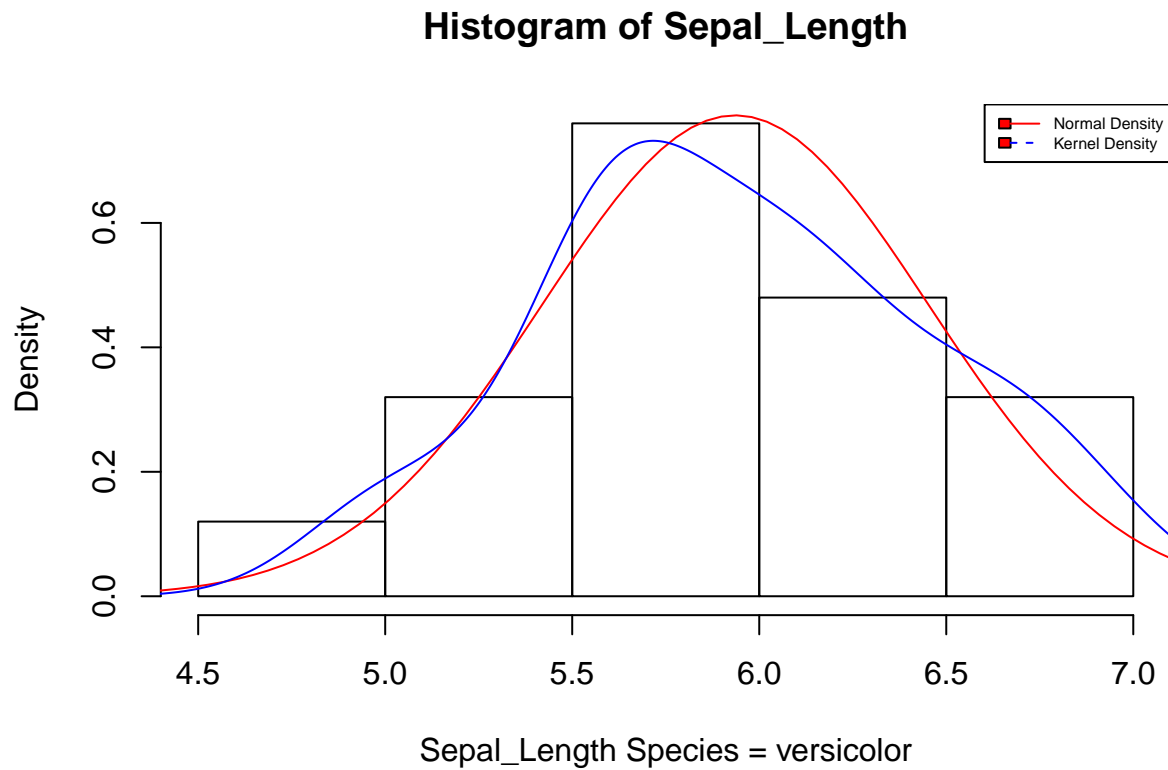
```
setosa_length <- iris$Sepal.Length[iris$Species == "setosa"]
hist(setosa_length, main = "Histogram of Sepal_Length",
     xlab = "Sepal_Length Species = setosa", freq=FALSE)
x <- seq(4, 6, length.out=100)
y <- with(iris, dnorm(x, mean(setosa_length), sd(setosa_length)))
lines(x, y, col="red")
lines(density(setosa_length), col="blue")
legend("topright", 5, 1.3, legend=c("Normal Density", "Kernel Density"),
     col=c("red", "blue"), lty=1:2, cex=0.5)
```

Histogram of Sepal_Length



Sepal_Length Species = setosa

```
versicolor_length <- iris$Sepal.Length[iris$Species == "versicolor"]
hist(versicolor_length, main = "Histogram of Sepal_Length",
     xlab = "Sepal_Length Species = versicolor", freq=FALSE)
x <- seq(4, 8, length.out=100)
y <- with(iris, dnorm(x, mean(versicolor_length), sd(versicolor_length)))
lines(x, y, col="red")
lines(density(versicolor_length), col="blue")
legend("topright", 5, 2, legend=c("Normal Density", "Kernel Density"),
     col=c("red", "blue"), lty=1:2, cex=0.5)
```



```
virginica_length <- iris$Sepal.Length[iris$Species == "virginica"]
hist(virginica_length, main = "Histogram of Sepal_Length",
     xlab = "Sepal_Length Species = virginica", freq=FALSE)
x <- seq(4, 8, length.out=100)
y <- with(iris, dnorm(x, mean(virginica_length), sd(virginica_length)))
lines(x, y, col="red")
lines(density(virginica_length), col="blue")
legend("topright", 5, 2, legend=c("Normal Density", "Kernel Density"),
     col=c("red", "blue"), lty=1:2, cex=0.5)
```

Histogram of Sepal_Length

