

week7 exercise

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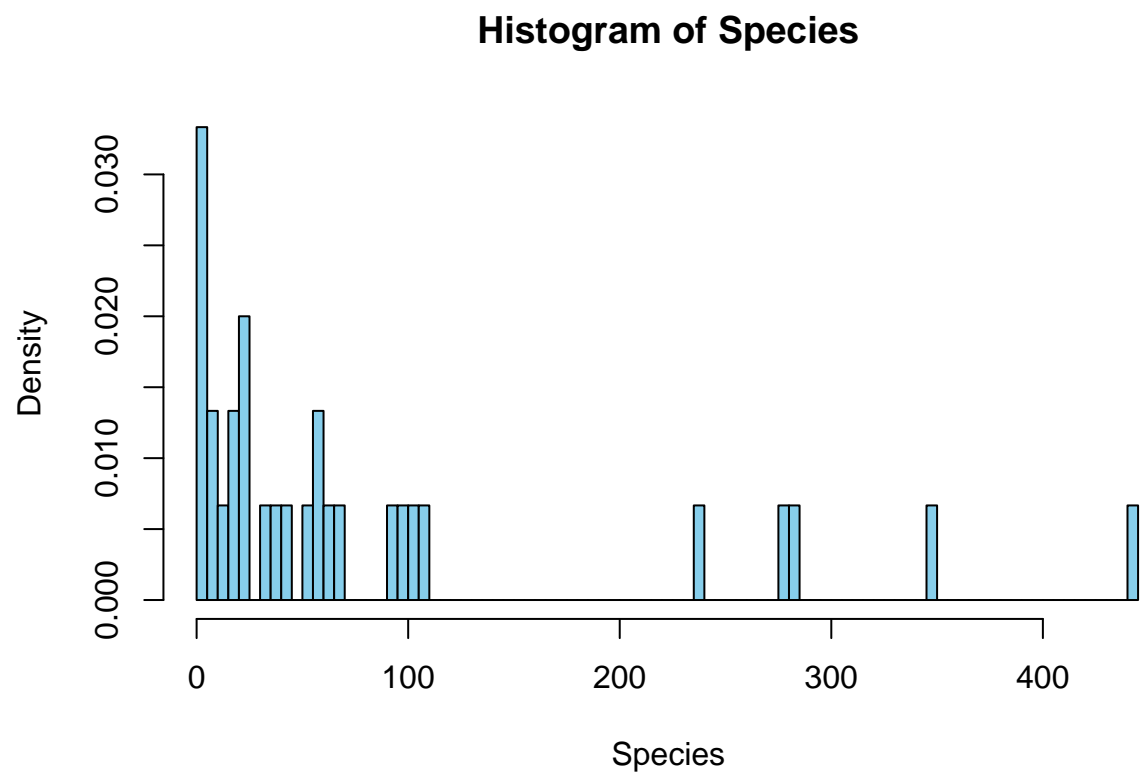
2023/11/16

```
library(faraway)
library(MASS)
```

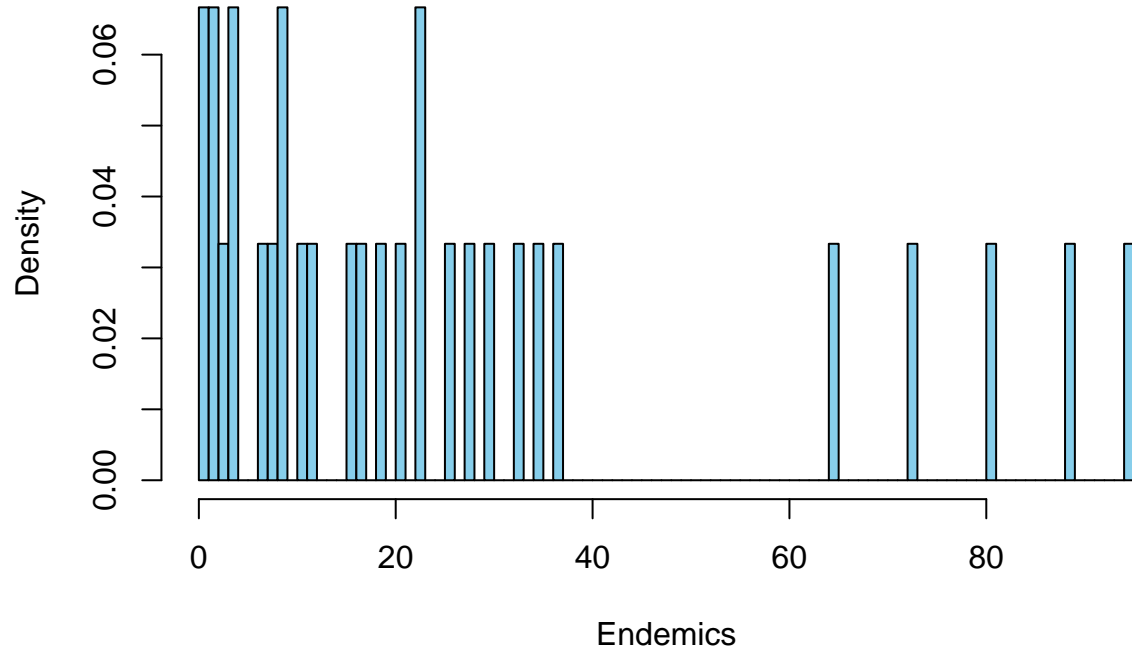
Exercise 2

```
f = function(df) {
  num_col = sapply(df, is.numeric)
  char_col = sapply(df, is.character)
  fac_col = sapply(df, is.factor)
  type = num_col | char_col | fac_col
  if (length(type[type]) == 0) {
    return("This dataframe contains no numeric, character or factor values")
  } else {
    num_df = df[, num_col]
    if (ncol(num_df) > 0) {
      for (i in 1:ncol(num_df)) {
        i_name = colnames(num_df)[i]
        hist(num_df[, i], breaks = 100, xlab = i_name,
             xlim = c(min(num_df[, i]), max(num_df[, i])), col = "skyblue", main = paste0("Histogram of ",
             i_name), probability = TRUE)
      }
    }
    char_df = df[, char_col | fac_col]
    if (ncol(char_df) > 0) {
      for (i in 1:ncol(char_df)) {
        i_name = colnames(char_df)[i]
        count = aggregate(char_df[, 1, drop = FALSE],
                          by = list(name = char_df[, i]), FUN = length)
        colnames(count)[2] = "count"
        barplot(height = count$count, names.arg = count$name,
                xlab = i_name, col = "lightgreen", ylim = c(0,
                max(count$count)), main = paste0("Barplot of ",
                i_name))
      }
    }
  }
}
```

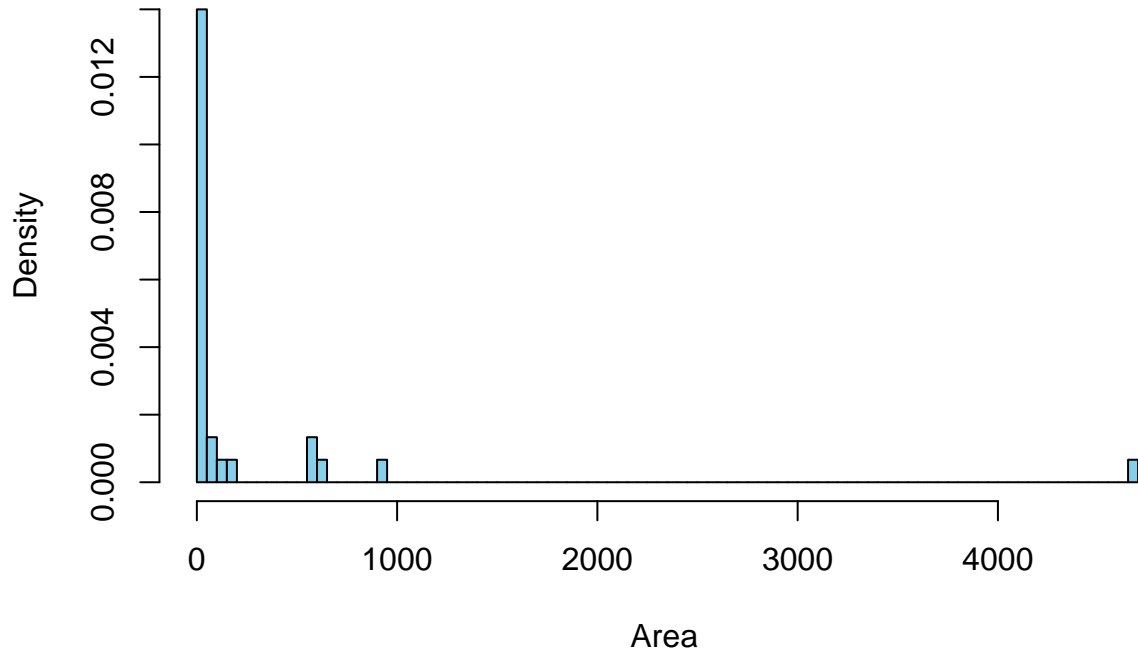
f(gala)



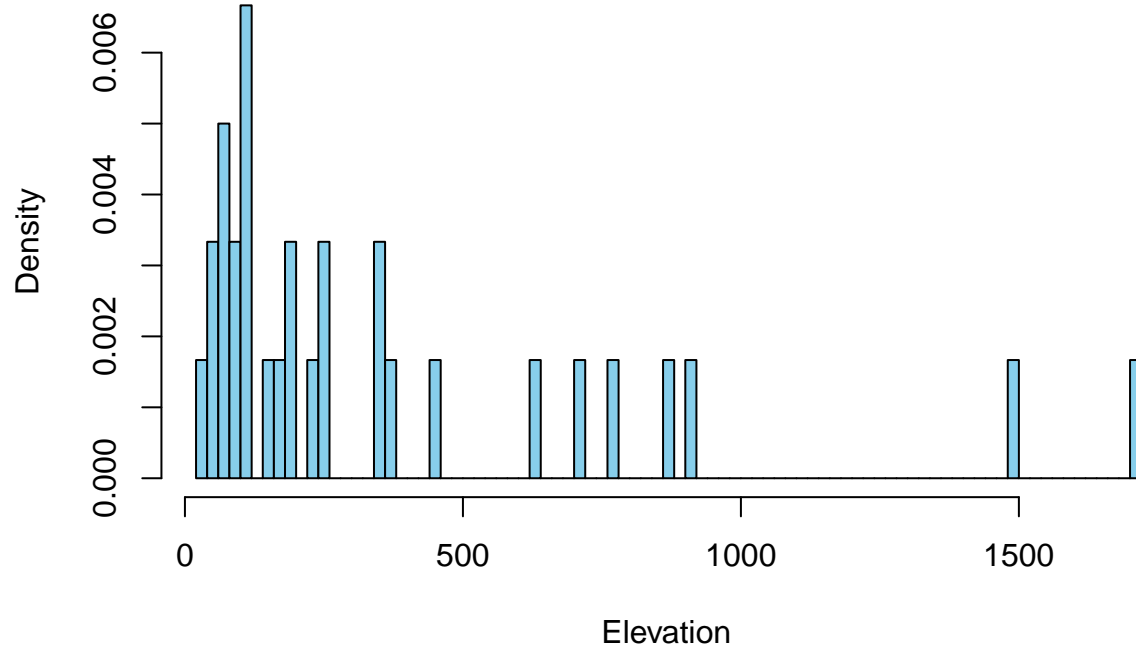
Histogram of Endemics



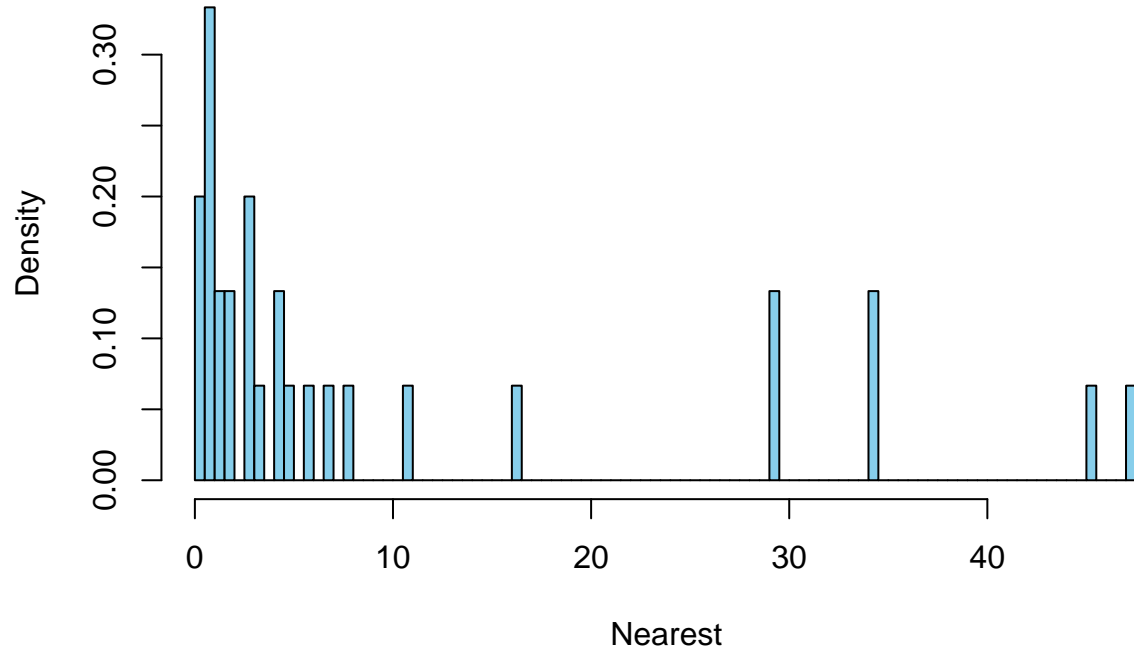
Histogram of Area



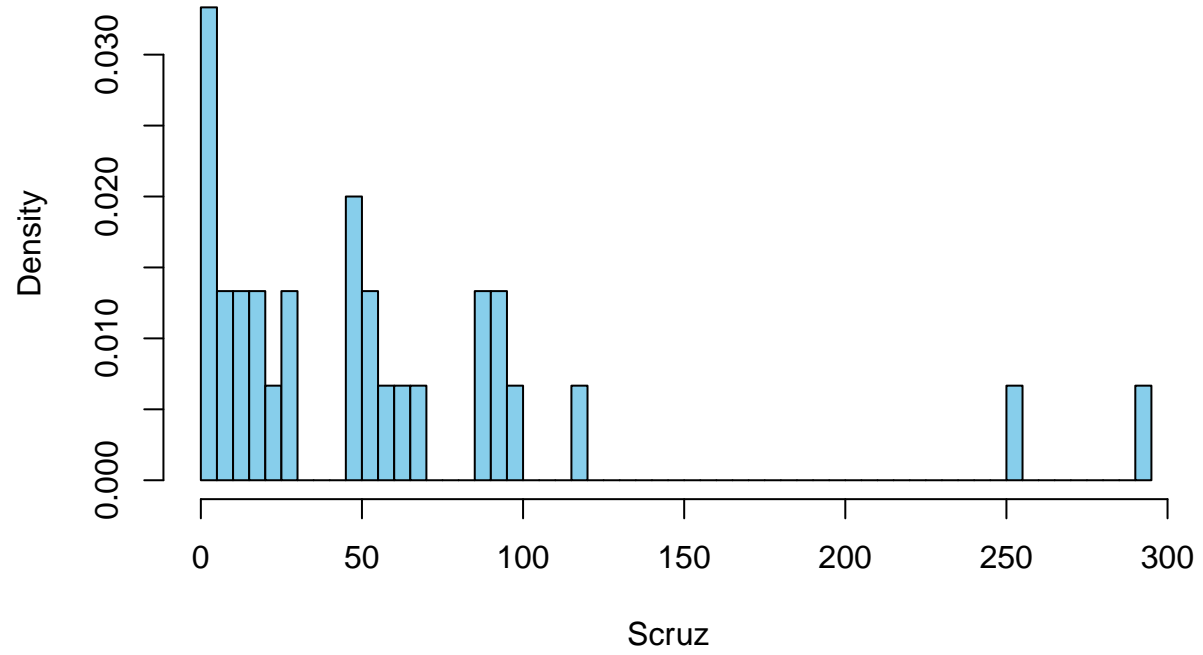
Histogram of Elevation



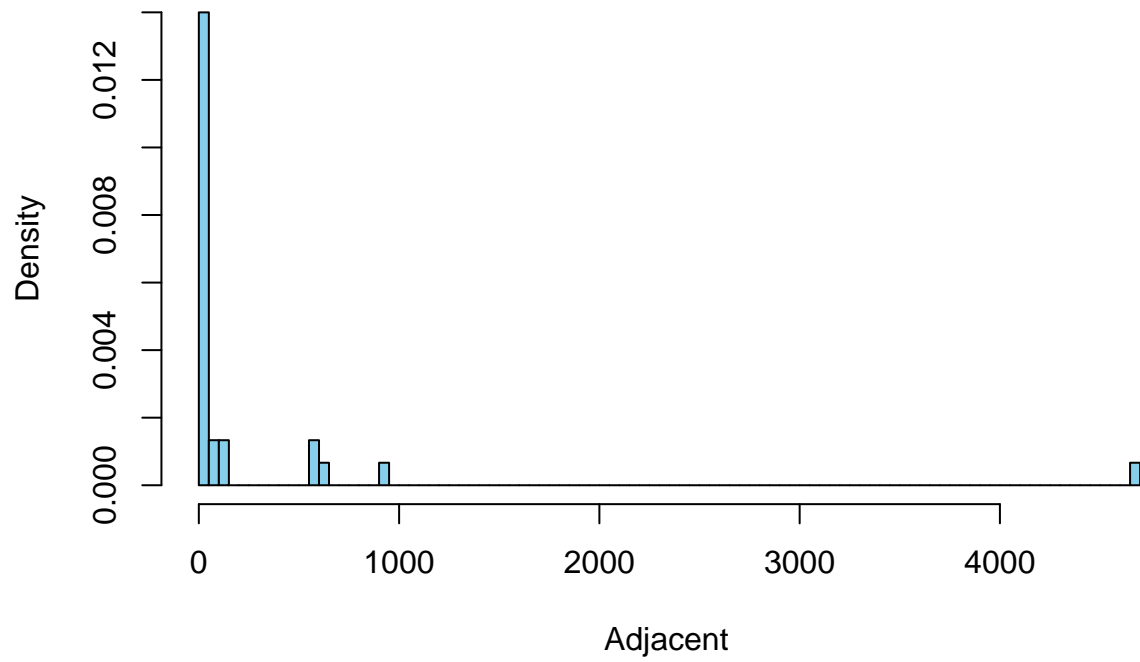
Histogram of Nearest



Histogram of Scruz

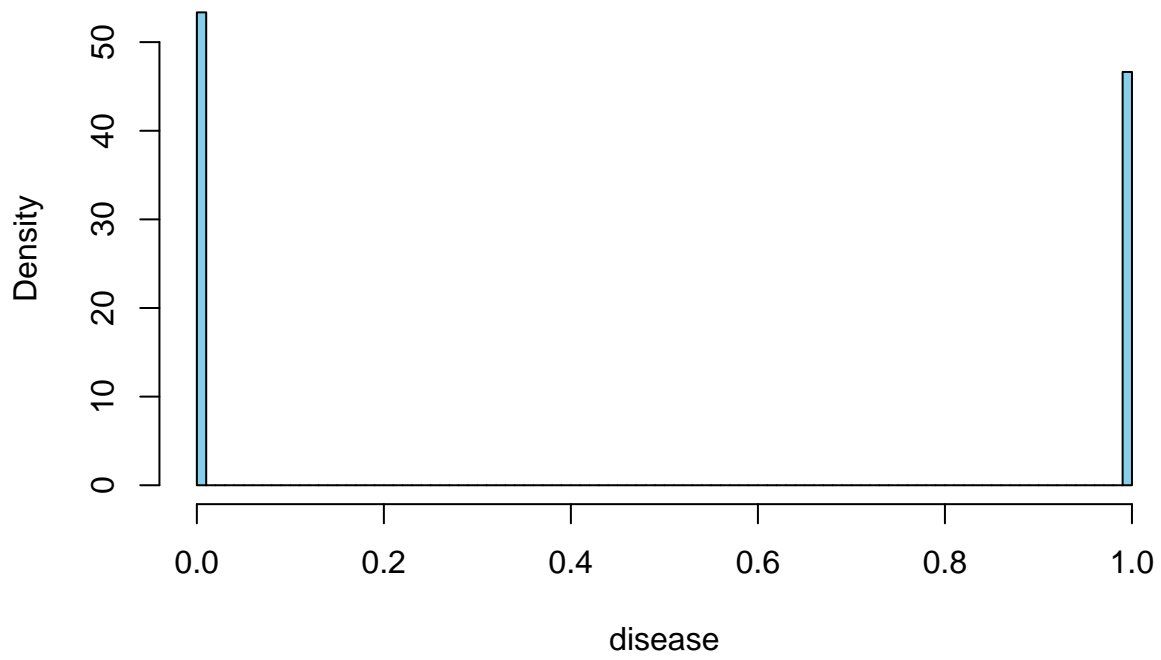


Histogram of Adjacent

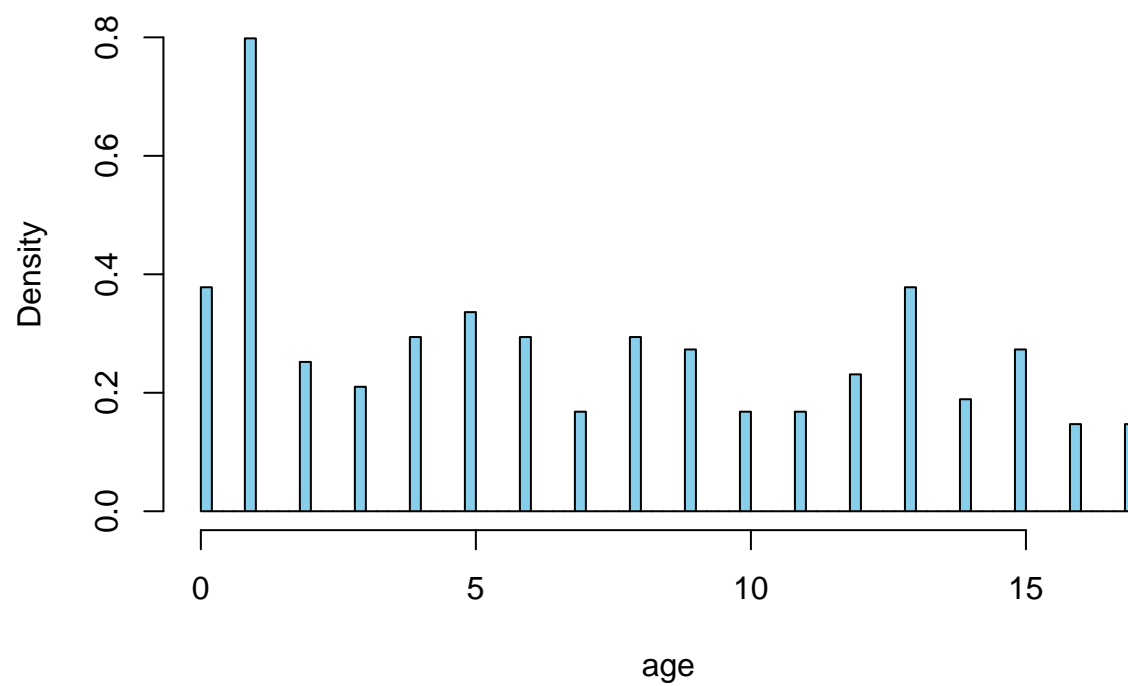


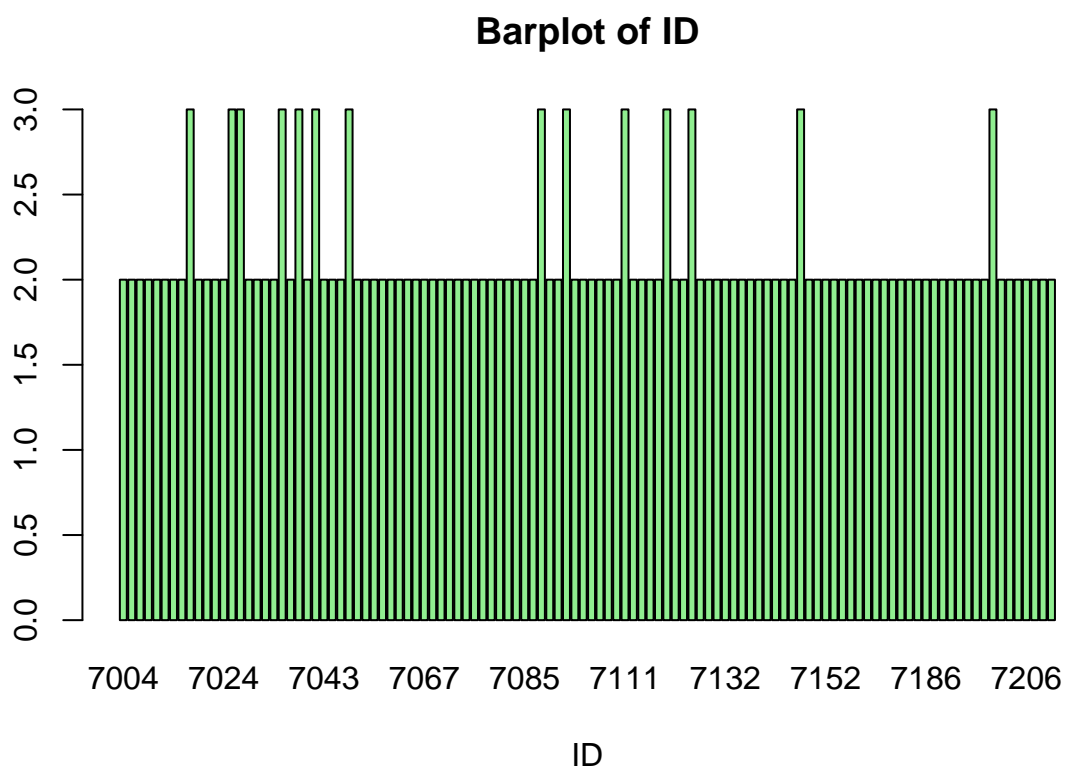
f(amlxray)

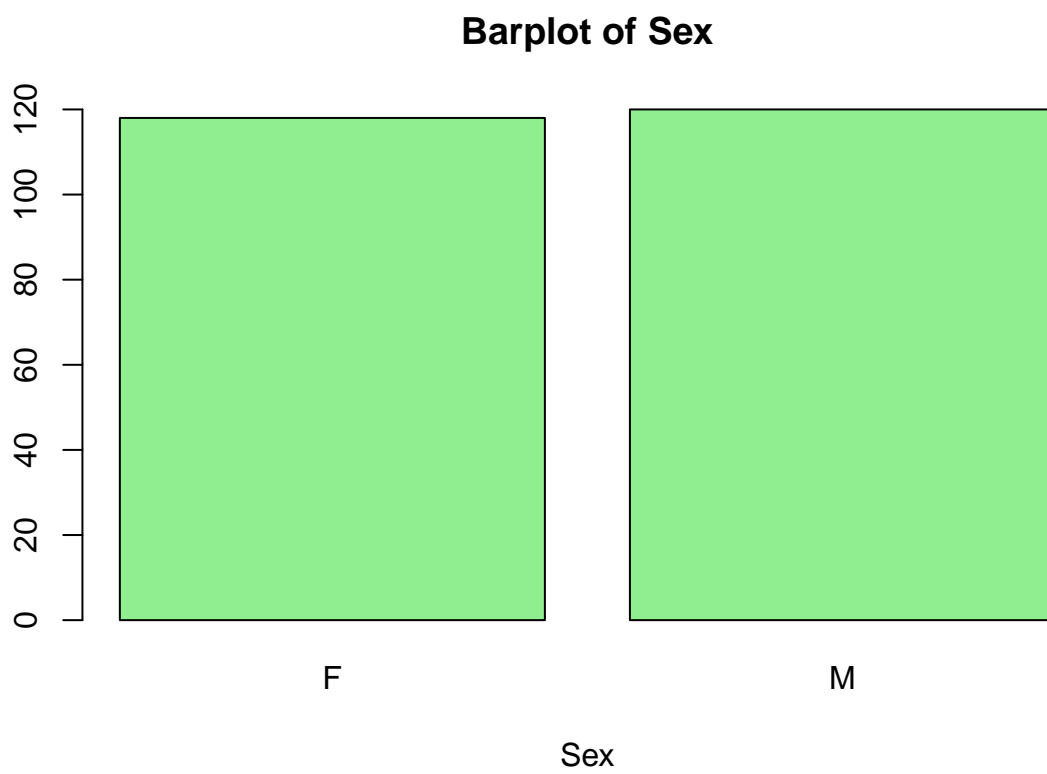
Histogram of disease



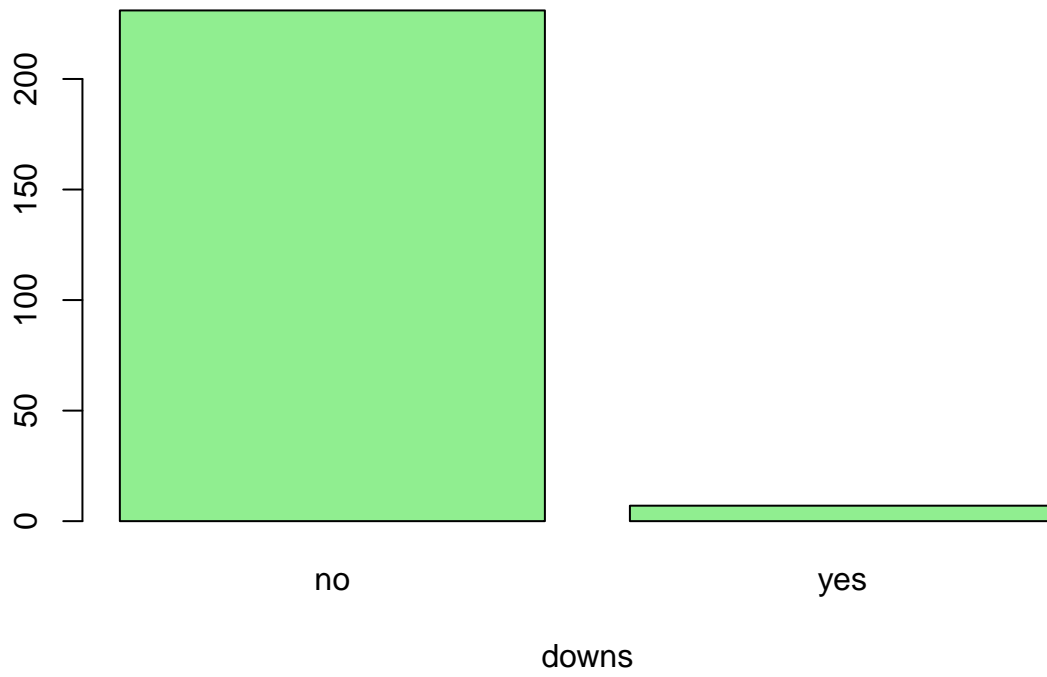
Histogram of age



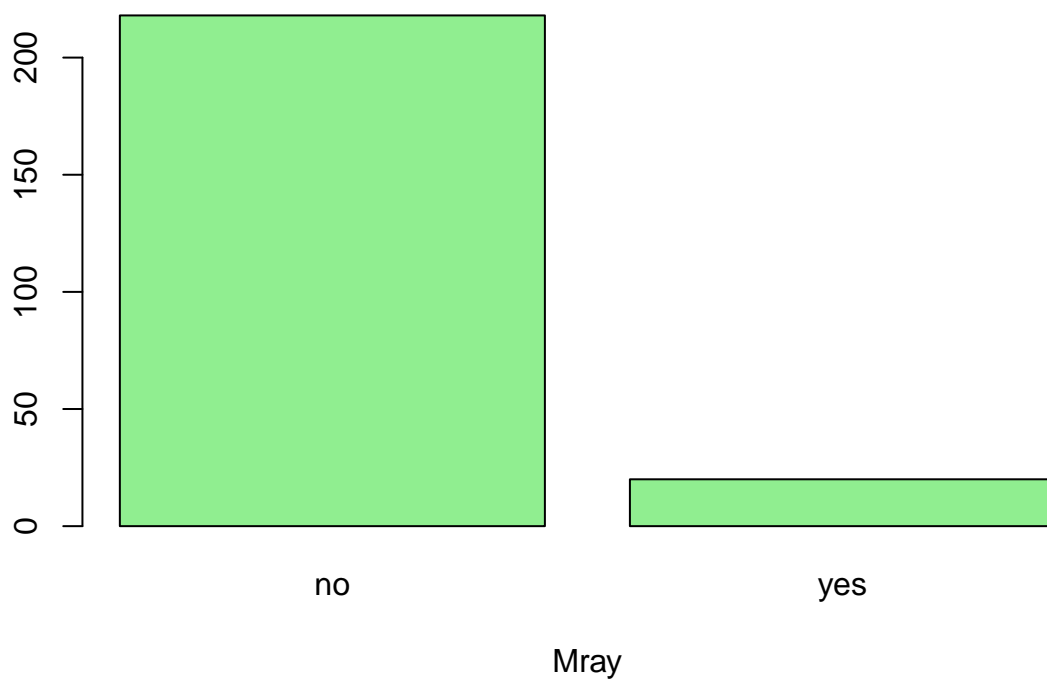


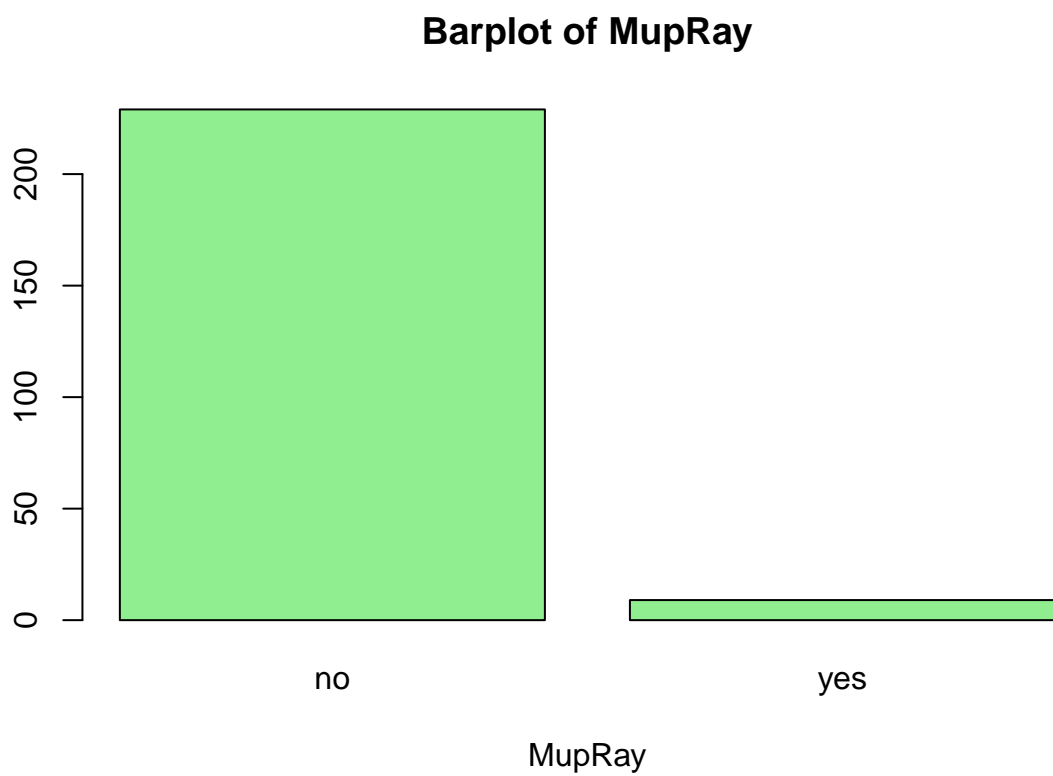


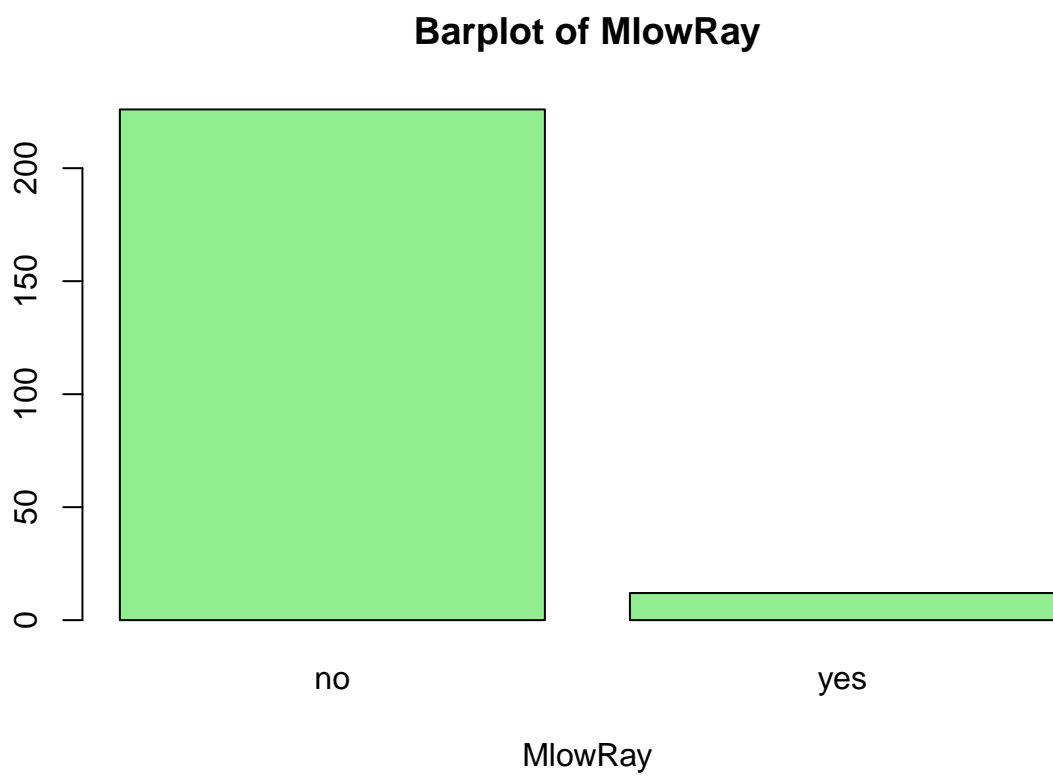
Barplot of downs



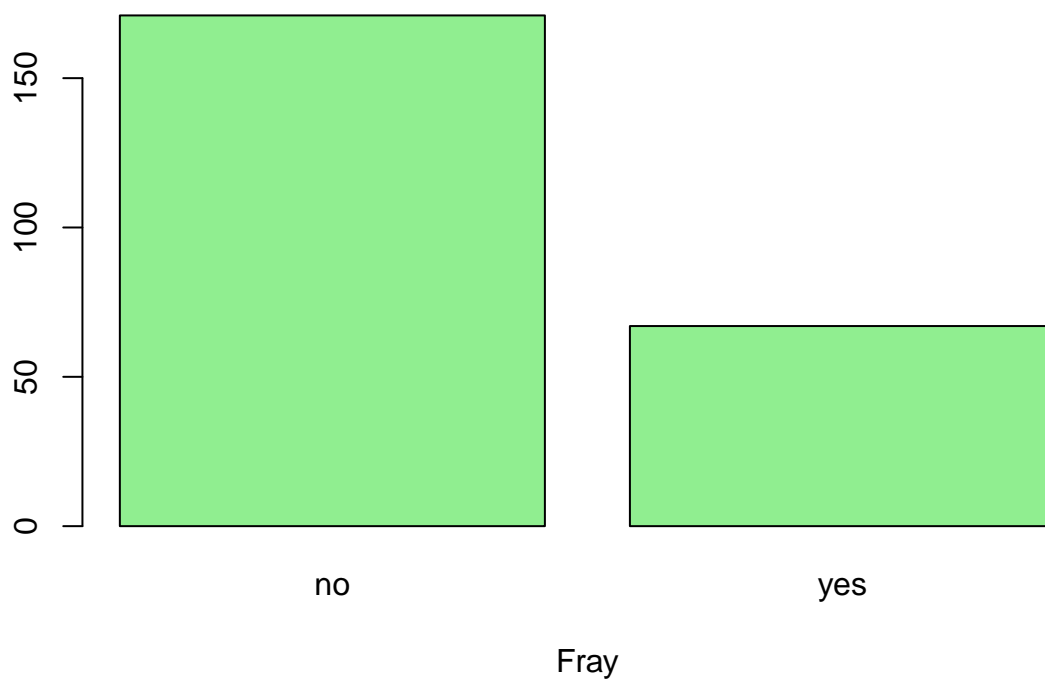
Barplot of Mray



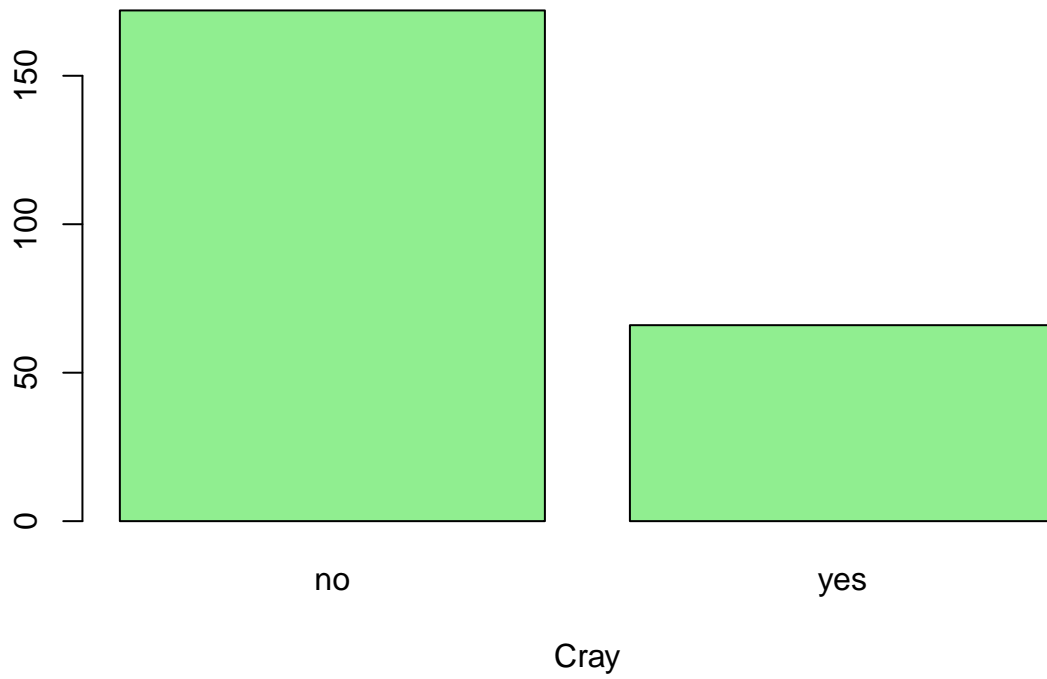




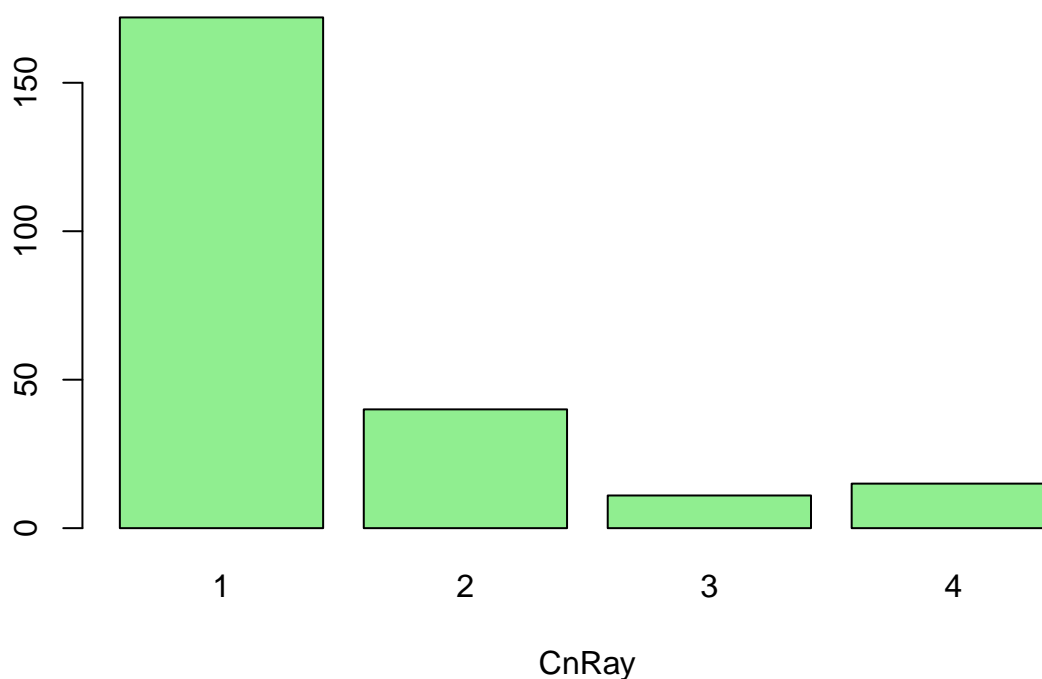
Barplot of Fray



Barplot of Cray



Barplot of CnRay



Exercise 3

1

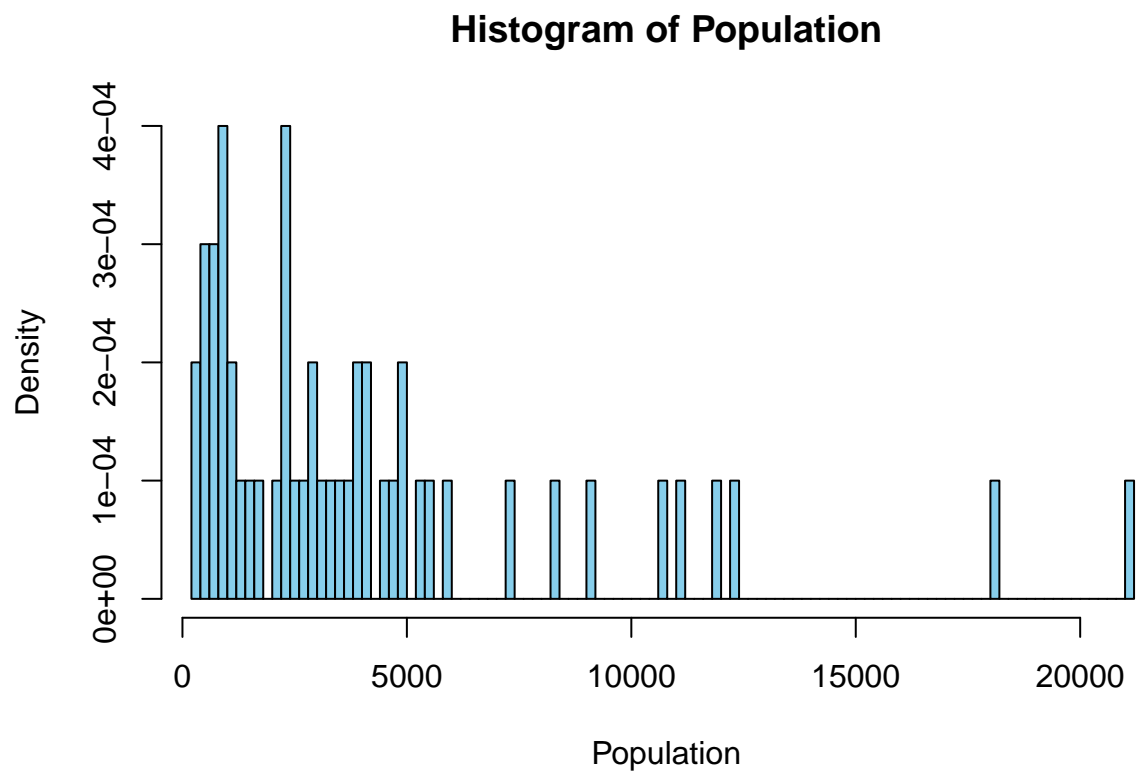
```
mean_v = apply(state.x77, 2, mean)
median_v = apply(state.x77, 2, median)
sd_v = apply(state.x77, 2, sd)
skew_v = apply(state.x77, 2, psych::skew)
sum_sta_df = data.frame(mean = mean_v, median = median_v,
  standard_deviation = sd_v, skewness = skew_v)
sum_sta_df
```

##	mean	median	standard_deviation	skewness
## Population	4246.4200	2838.500	4.464491e+03	1.9222511
## Income	4435.8000	4519.000	6.144699e+02	0.2046903
## Illiteracy	1.1700	0.950	6.095331e-01	0.8185809
## Life Exp	70.8786	70.675	1.342394e+00	-0.1534995
## Murder	7.3780	6.850	3.691540e+00	0.1293391
## HS Grad	53.1080	53.250	8.076998e+00	-0.3192442
## Frost	104.4600	114.500	5.198085e+01	-0.3663767
## Area	70735.8800	54277.000	8.532730e+04	4.0983574

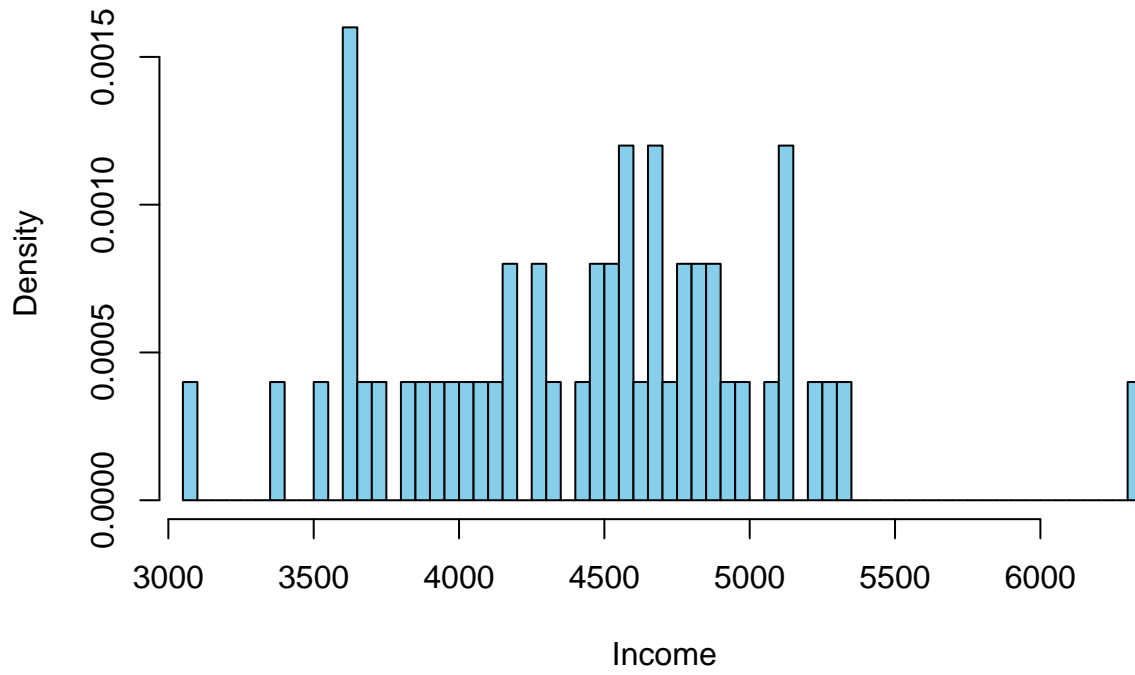
```

cns = colnames(state.x77)
for (i in 1:ncol(state.x77)) {
  i_name = cns[i]
  hist(state.x77[, i], breaks = 100, xlab = i_name, xlim = c(min(state.x77[,
    i]), max(state.x77[, i])), col = "skyblue", main = paste0("Histogram of ",
    i_name), probability = TRUE, plot = TRUE)
}

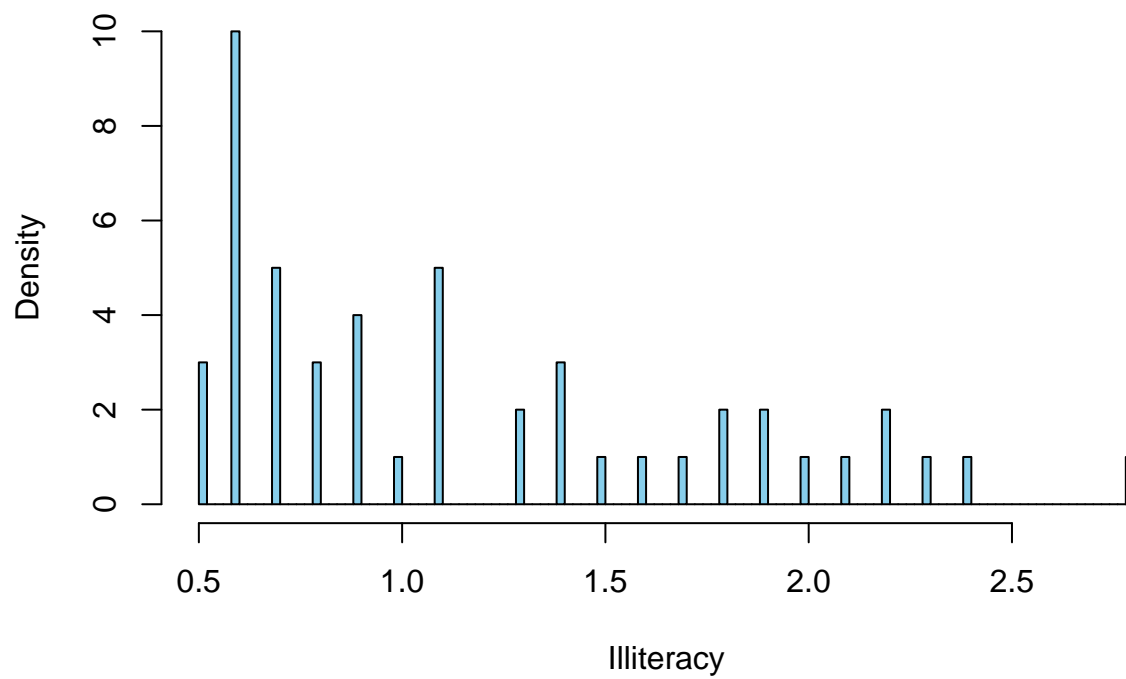
```

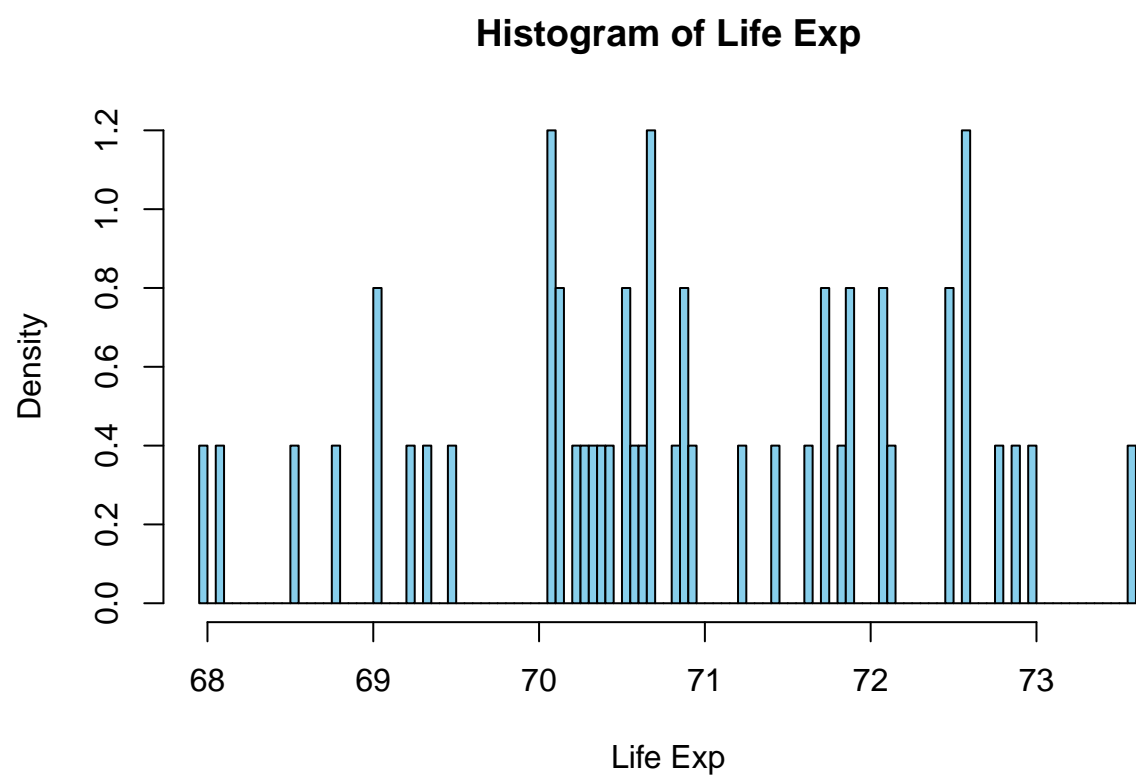


Histogram of Income

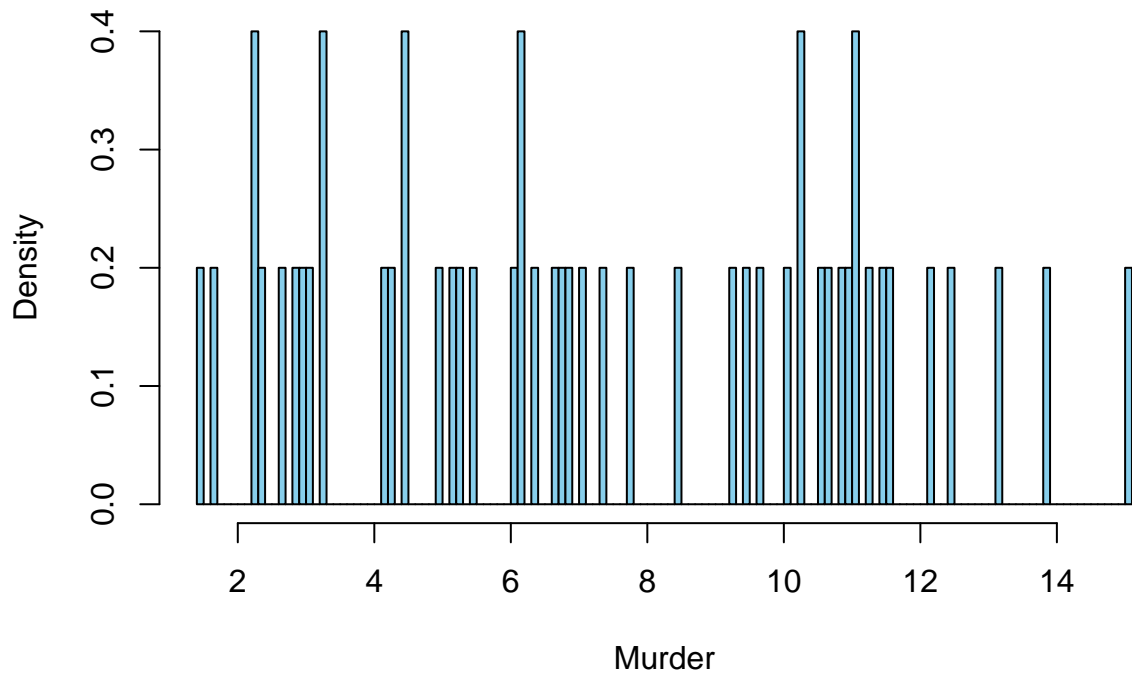


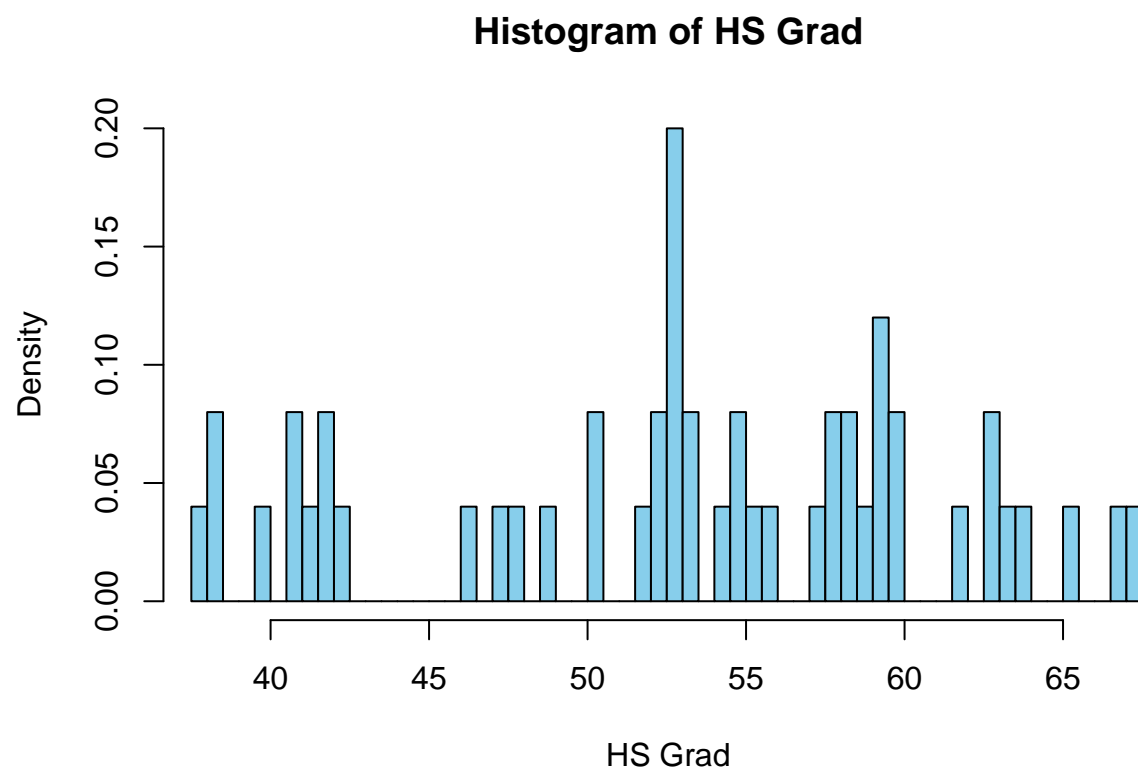
Histogram of Illiteracy

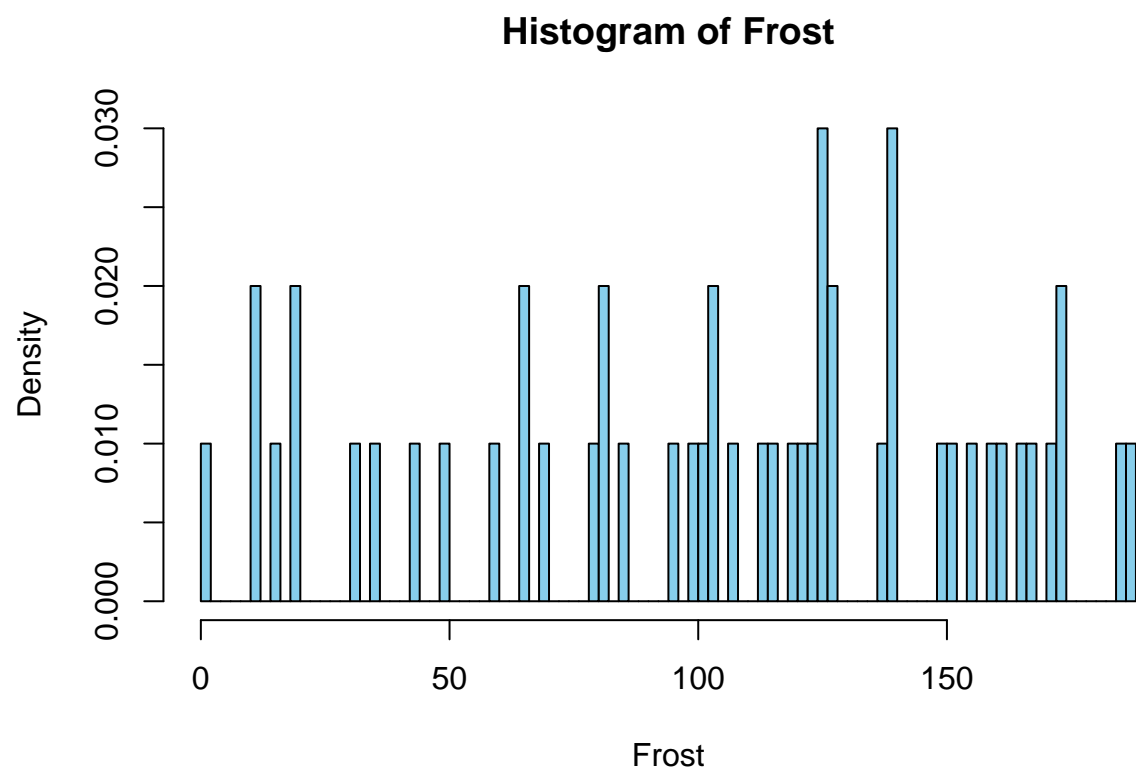


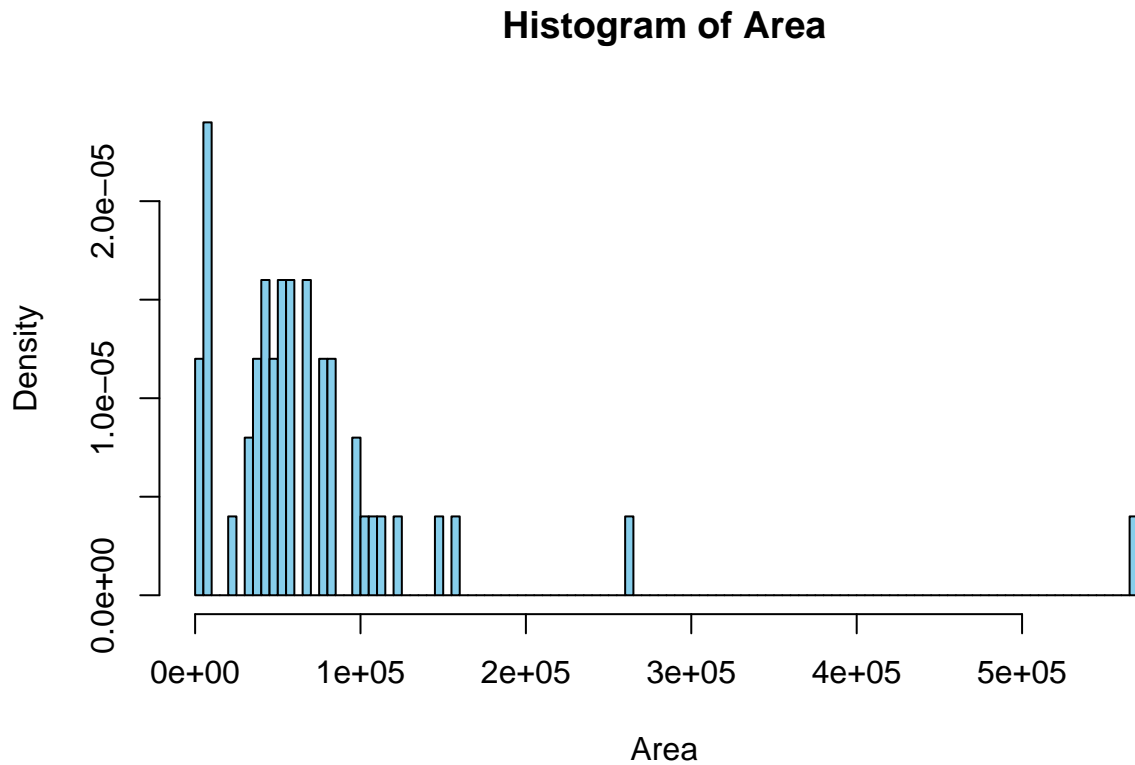


Histogram of Murder







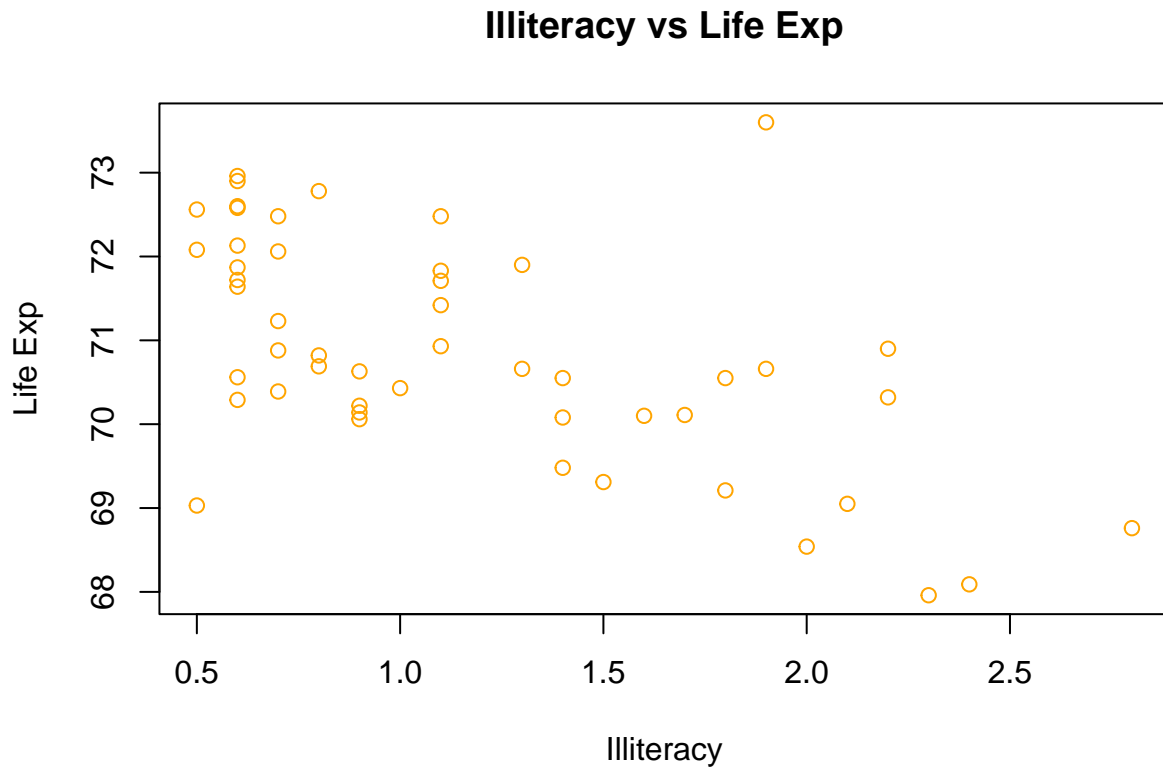


3

The Income, Life Exp and Murder are almost symmetric. Population, Illiteracy and Area are positive skewed. HS Grad and Frost are negative skewed.

4

```
plot(x = state.x77[, "Illiteracy"], y = state.x77[, "Life Exp"],  
     type = "p", col = "orange", xlab = "Illiteracy", ylab = "Life Exp",  
     main = "Illiteracy vs Life Exp")
```



With the increase of Illiteracy rate, the life expectancy decreases.

5

```
cor.test(x = state.x77[, "Illiteracy"], y = state.x77[,
  "Life Exp"], alternative = "two.sided", method = "pearson")
```

```
##
## Pearson's product-moment correlation
##
## data: state.x77[, "Illiteracy"] and state.x77[, "Life Exp"]
## t = -5.0427, df = 48, p-value = 6.969e-06
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.7448226 -0.3708811
## sample estimates:
## cor
## -0.5884779
```

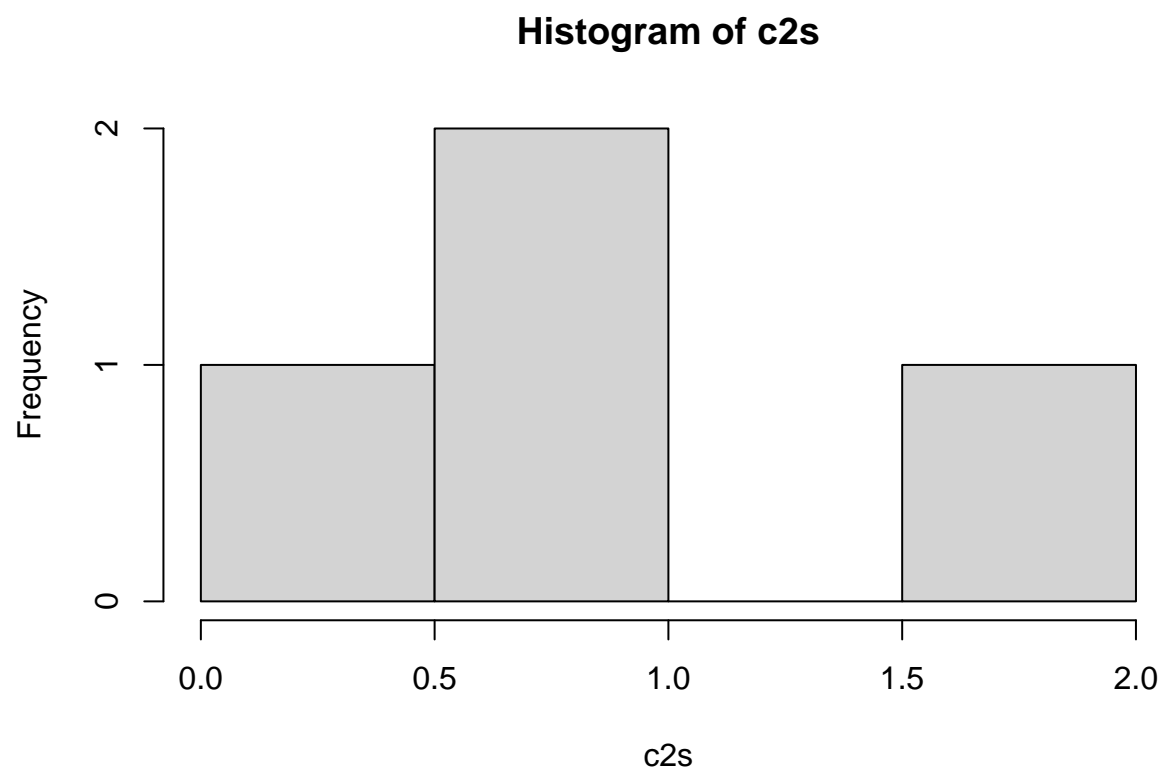
Lower Illiteracy rate means higher education level, which will improve people's life quality and increase life expectancy.

Exercise 4

```
c1 = c(0, 1)
c2 = expand.grid(c1, c1)
c2s = rowSums(c2)
```

1

```
hist(c2s, breaks = 3, main = "Histogram of c2s")
```



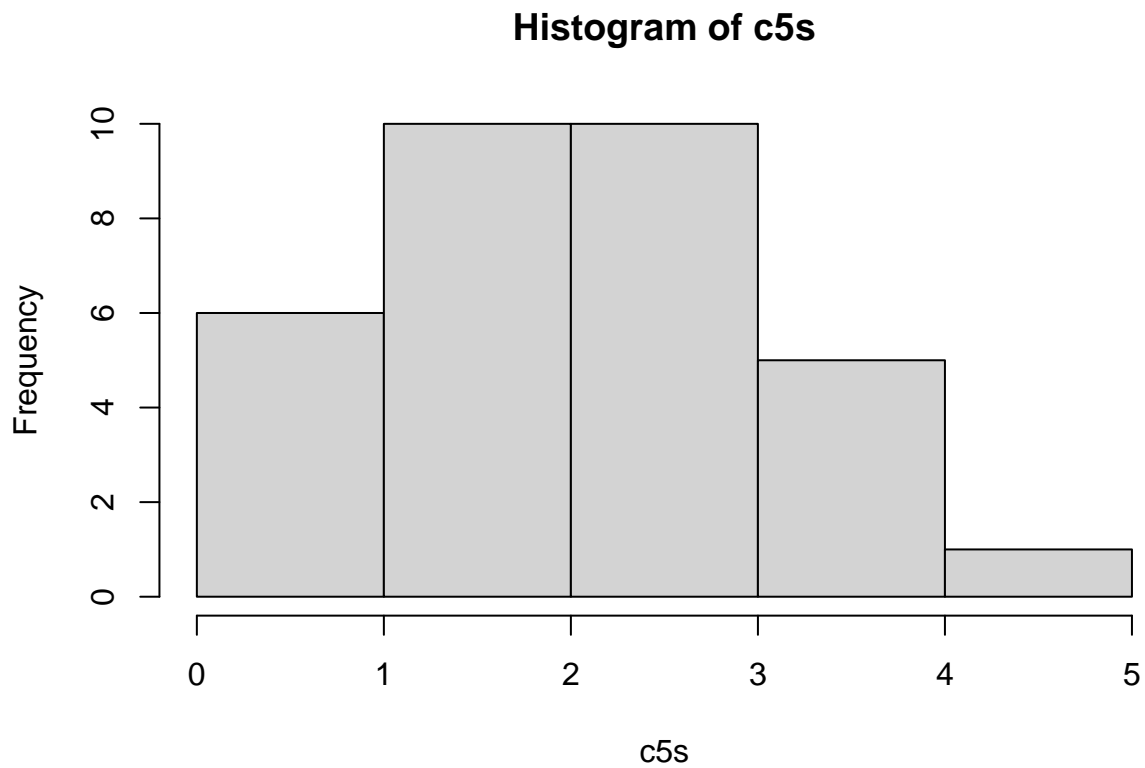
```
dim(c2)
```

```
## [1] 4 2
```

`expand.grid()` lists all combinations from given vectors. `rowSums()` lists the sum of each combination from given vectors.

2

```
c5 = expand.grid(c1, c1, c1, c1, c1)
c5s = rowSums(c5)
hist(c5s, breaks = 6, main = "Histogram of c5s")
```



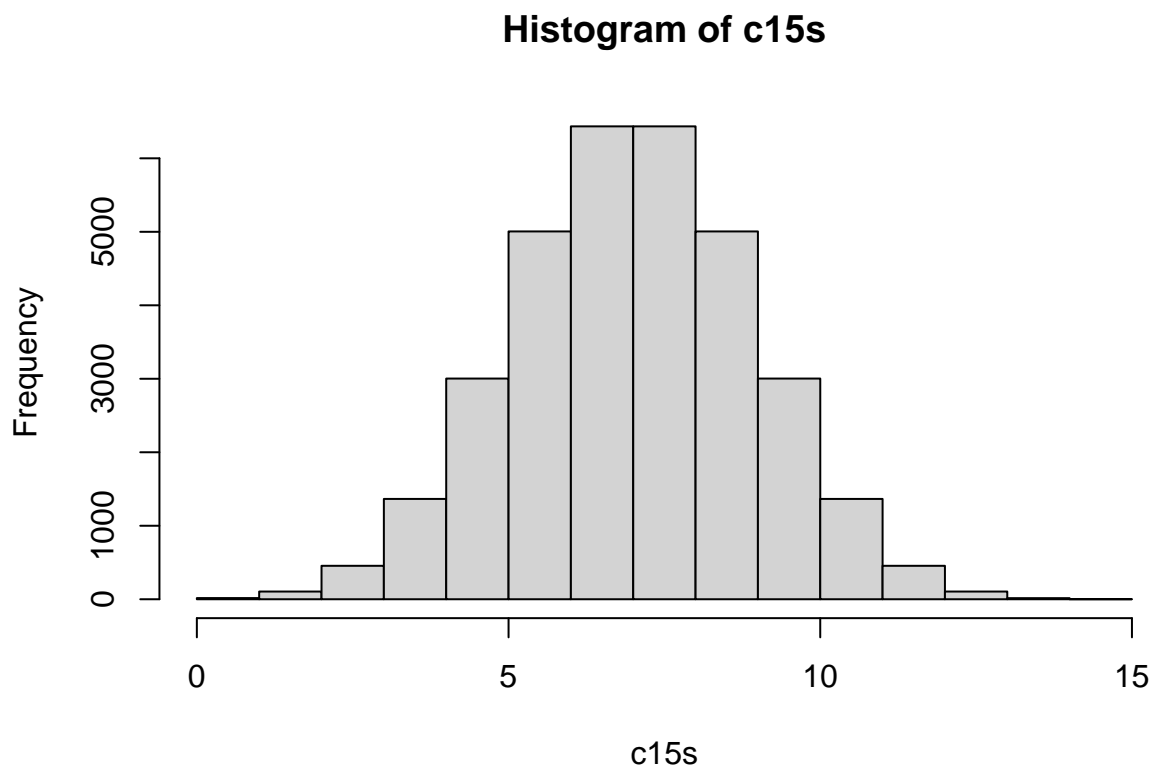
```
dim(c5)
```

```
## [1] 32 5
```

The distribution of c5s gets a peak in center.

3

```
c15 = expand.grid(replicate(15, c1, simplify = F))
c15s = rowSums(c15)
hist(c15s, breaks = 16, main = "Histogram of c15s")
```



```
dim(c15) == c(2^15, 15)
```

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

```
4
```

```
sta_ls = list(range = range(c15s), mean = mean(c15s), std = sd(c15s))
sta_ls
```

```
## $range
## [1] 0 15
##
## $mean
## [1] 7.5
##
## $std
## [1] 1.936521
```

```
c(sta_ls$mean, 15 * 0.5)
```

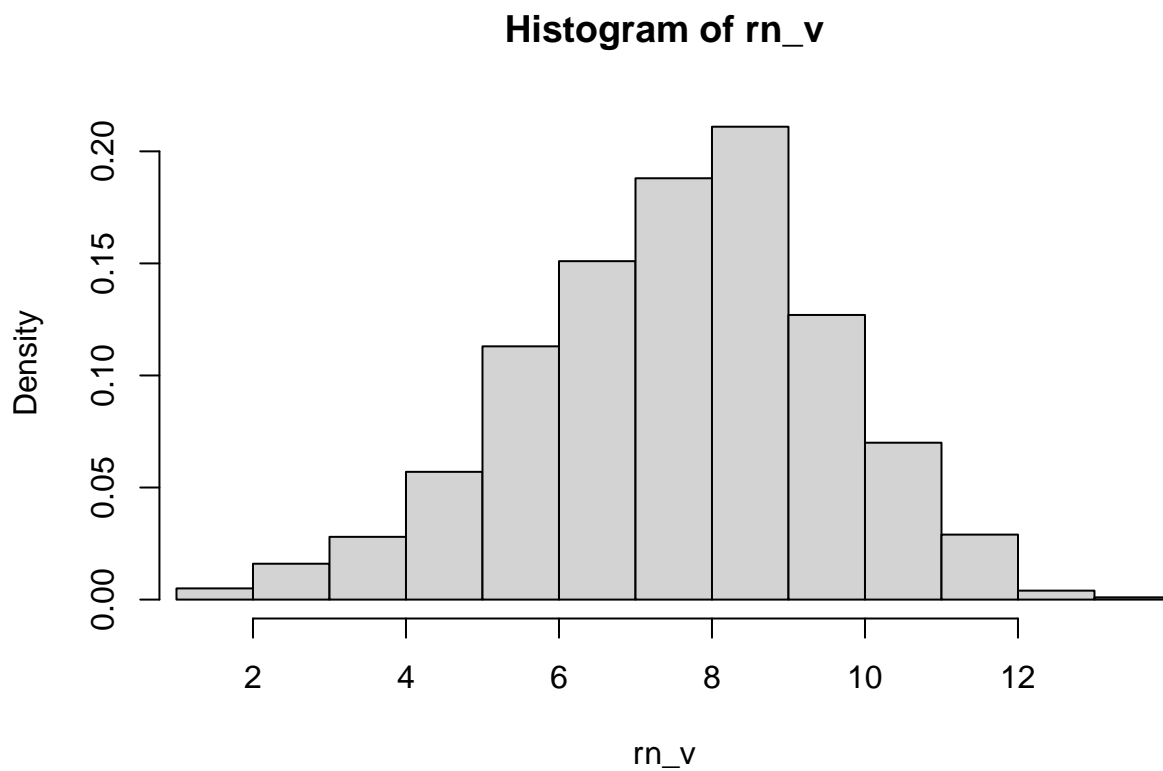
```
## [1] 7.5 7.5
```

```
c(sta_ls$std, (15 * 0.5 * 0.5)^0.5)
```

```
## [1] 1.936521 1.936492
```

5

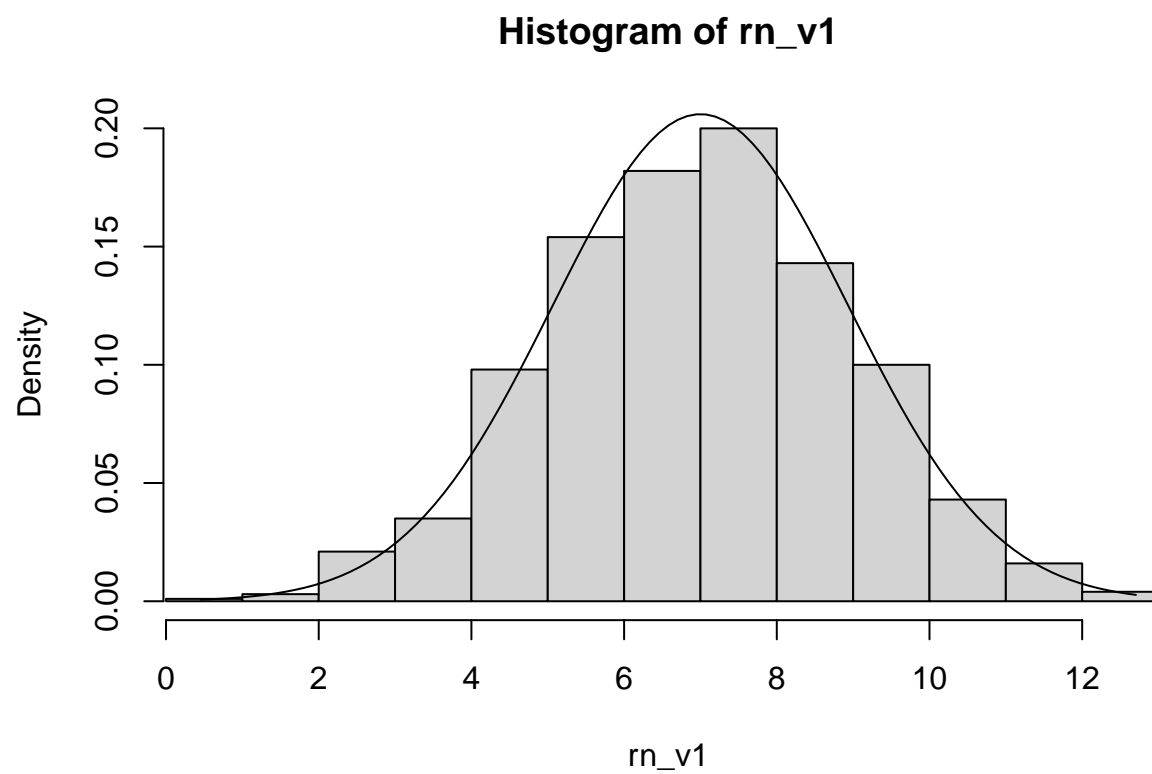
```
rn_v = rnorm(1000, mean = sta_ls$mean, sd = sta_ls$std)
hist(rn_v, breaks = 16, xlim = c(min(rn_v), max(rn_v)),
     main = "Histogram of rn_v", probability = TRUE)
```



The distribution of 10^3 random numbers is similar to the distribution of c15s.

6

```
rn_v1 = rnorm(1000, mean = sta_ls$mean - 0.5, sd = sta_ls$std)
hist(rn_v1, breaks = 16, xlim = c(min(rn_v1), max(rn_v1)),
     main = "Histogram of rn_v1", probability = TRUE)
curve(expr = dnorm(x, sta_ls$mean - 0.5, sta_ls$std), add = TRUE)
```

Exercise 5

```
fib = c(1, 1)
i = 2
fib_sum = sum(fib)
while (fib_sum < 22000) {
  i = i + 1
  fib[i] = fib[i - 2] + fib[i - 1]
  fib_sum = sum(fib)
}
length(fib)
```

```
## [1] 21
```