

Homework 01 Solutions

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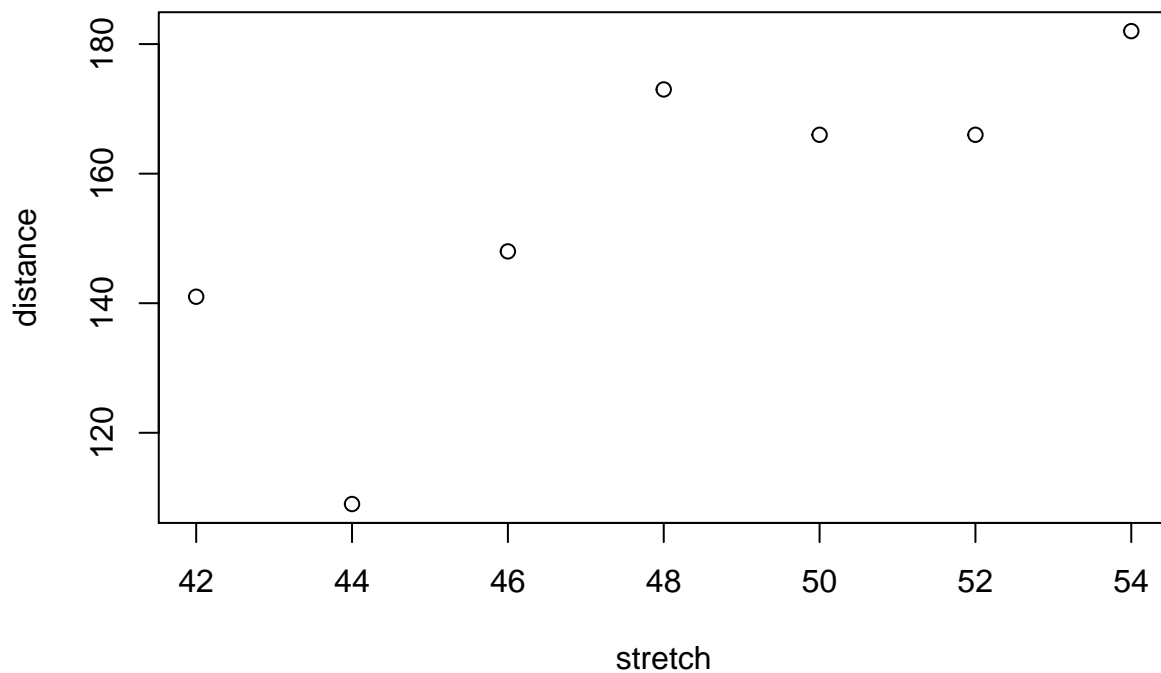
8/30/2018

Chapter 1 Exercises

```
load("../usingR.RData")
```

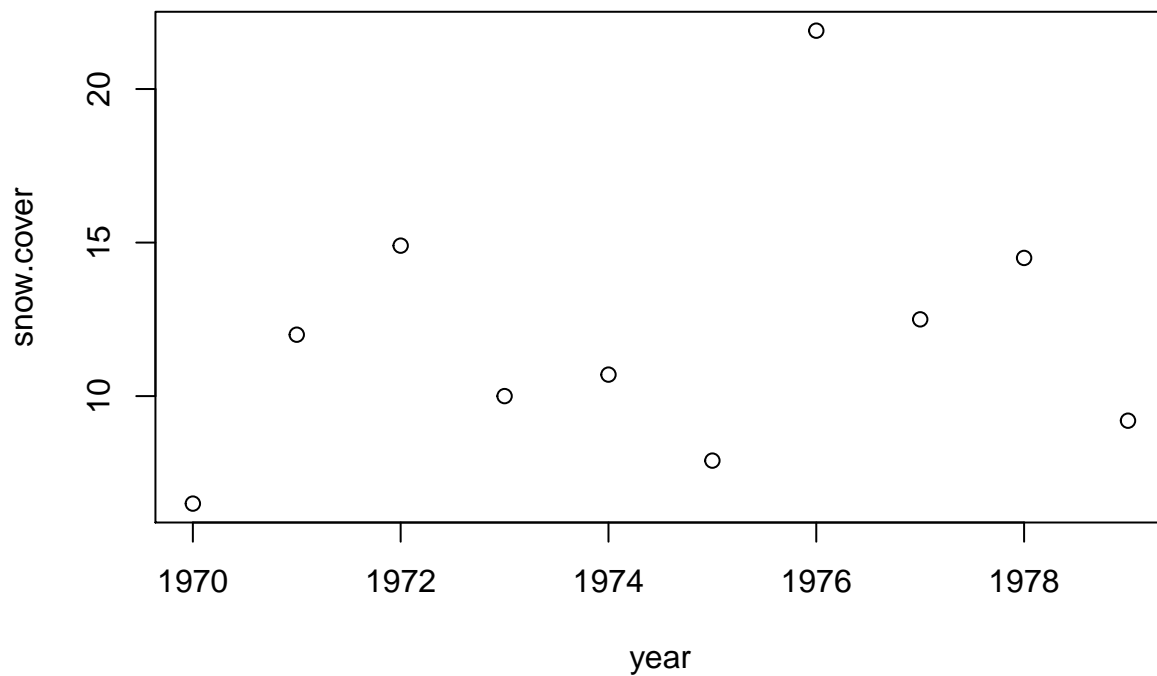
Exercise 1

```
plot(distance ~ stretch, data = elasticband)
```



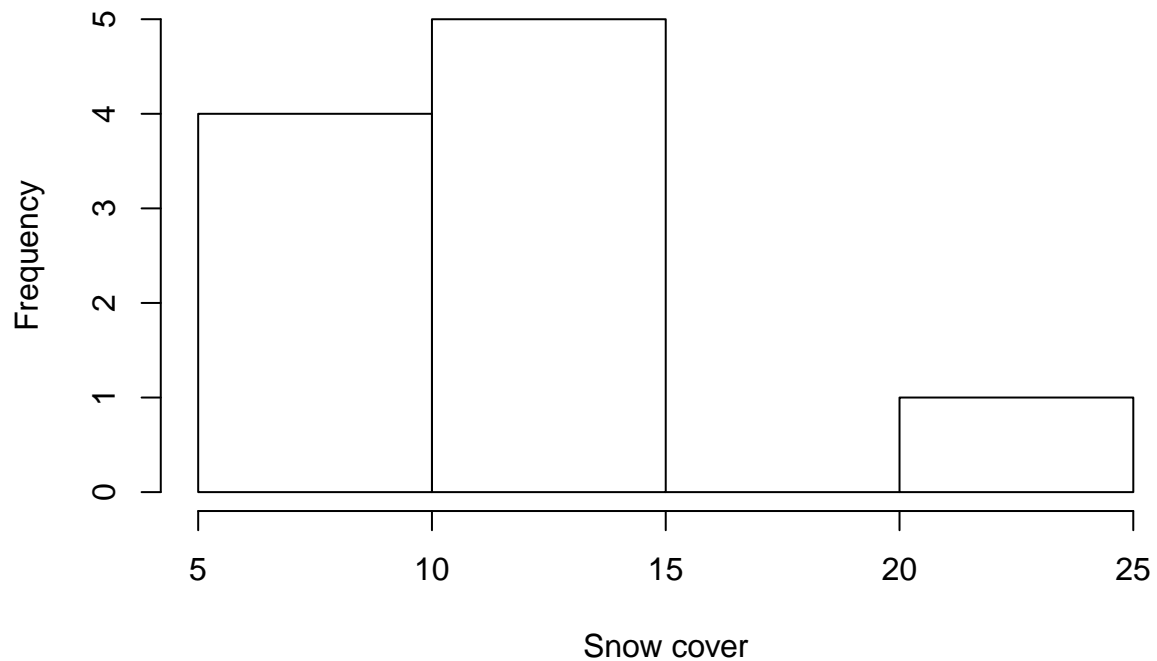
Exercise 2

```
# i.  
snow <- data.frame(year = 1970:1979,  
                  snow.cover = c(6.5, 12, 14.9, 10, 10.7, 7.9, 21.9, 12.5, 14.5, 9.2))  
  
# ii.  
plot(snow.cover ~ year, data = snow)
```

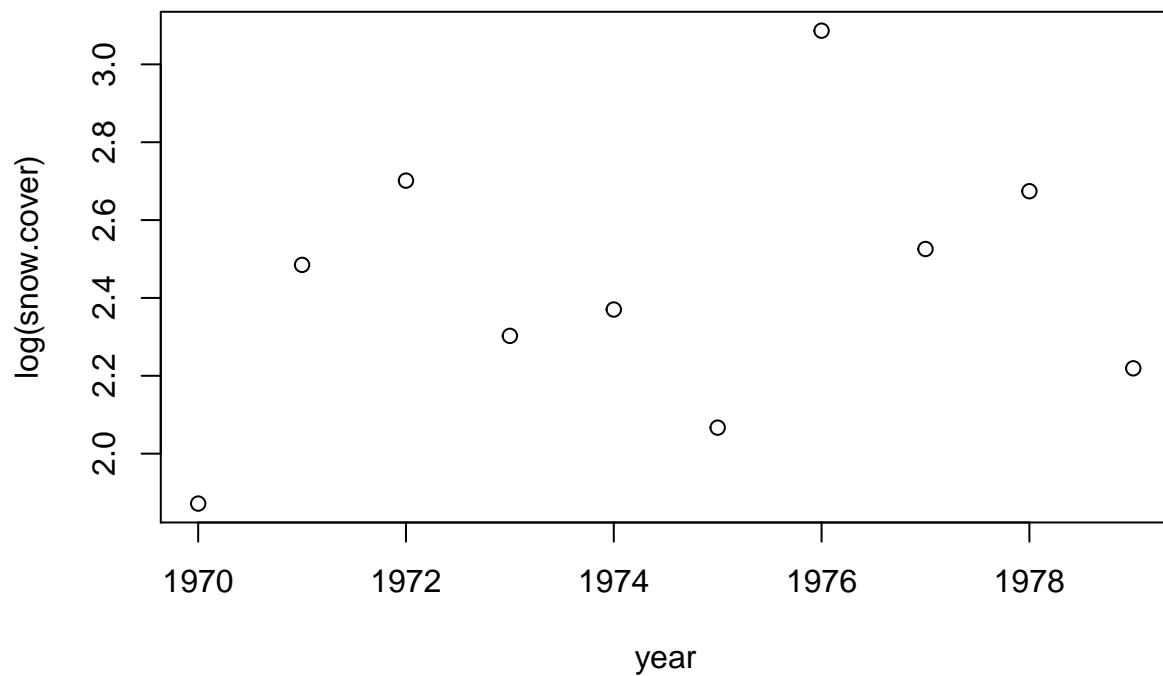


```
# iii.
hist(snow$snow.cover, xlab = "Snow cover", main = "Histogram of snow cover")
```

Histogram of snow cover

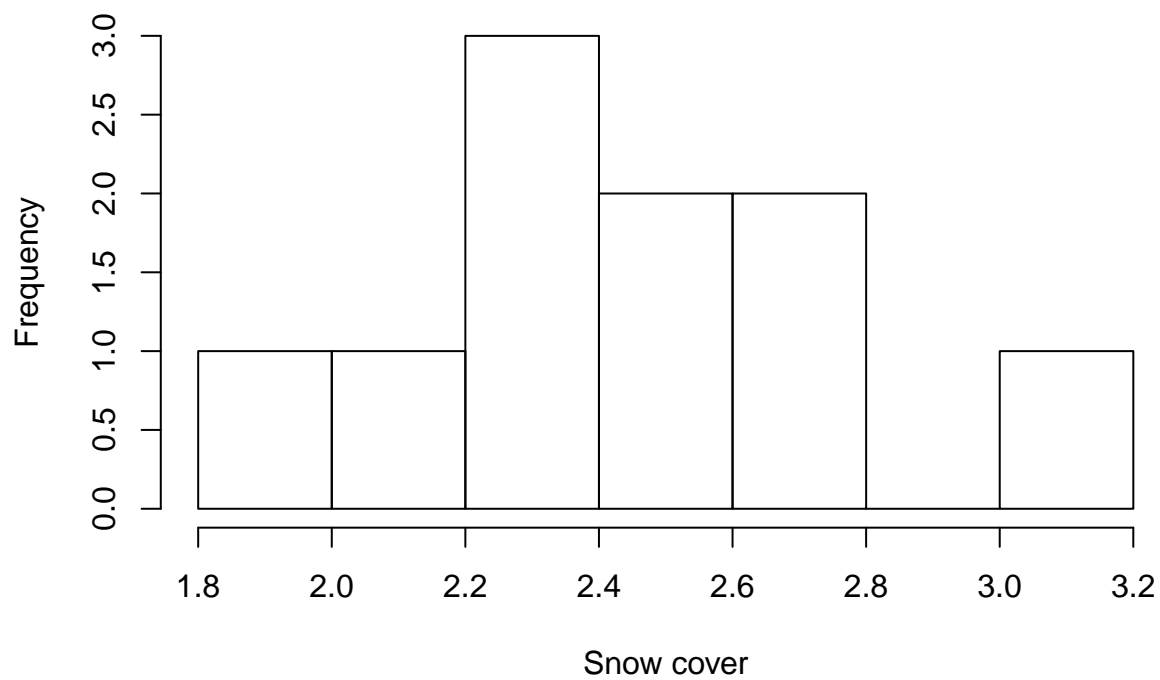


```
# iv.
plot(log(snow.cover) ~ year, data = snow)
```



```
hist(log(snow$snow.cover), xlab = "Snow cover", main = "Histogram of snow cover")
```

Histogram of snow cover



Exercise 3

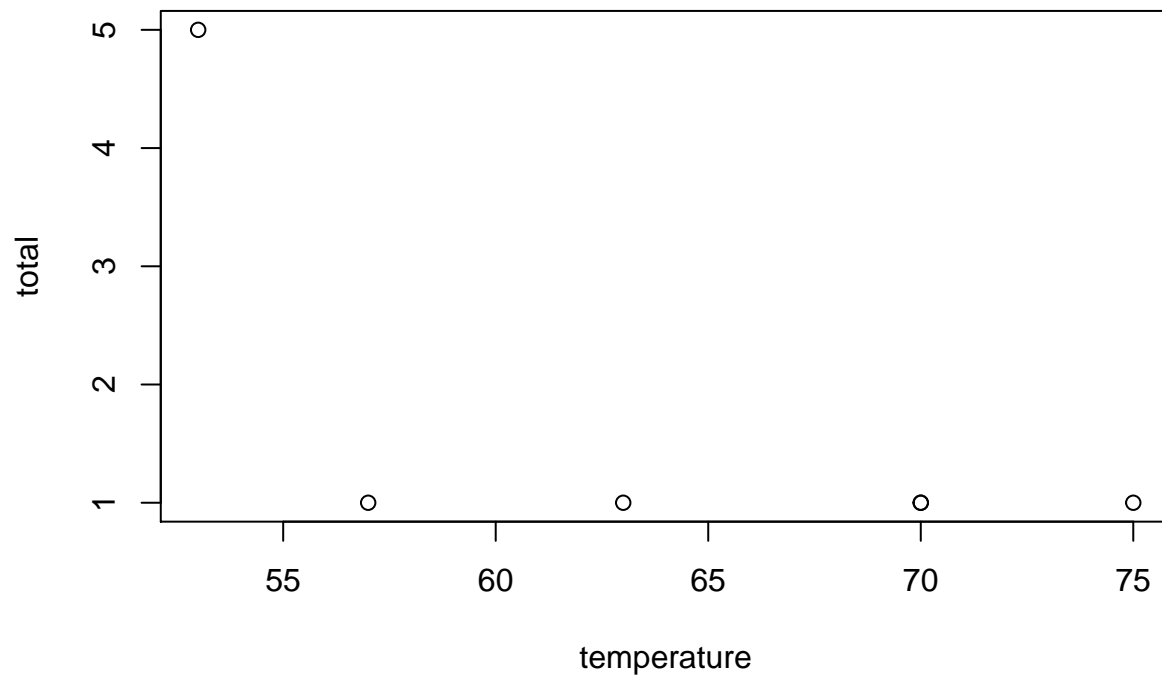
```
orings <- data.frame(temperature = c(53, 57, 63, 70, 70, 75),
                        erosion = c(3, 1, 1, 1, 1, 0),
```

```

blowby = c(2, 0, 0, 0, 0, 2),
total = c(5, 1, 1, 1, 1, 1))

plot(total ~ temperature, data = orings)

```



Chapter 2 Exercises

Exercise 1

```

# 2.1(a)
answer <- 0
for (j in 3:5) {answer <- j + answer}
print(answer)

```

```
## [1] 12
```

```

# 2.1(b)
answer <- 10
for (j in 3:5) {answer <- j + answer}
print(answer)

```

```
## [1] 22
```

```

# 2.1(c)
answer <- 10
for (j in 3:5) {answer <- j*answer}
print(answer)

```

```
## [1] 600
```

Exercise 2

```
?prod  
prod(c(10, 3, 4, 5))
```

```
## [1] 600
```

Exercise 5

```
radii <- 3:20  
spheres <- data.frame(radius = radii, volume = 4*pi*radii^3)  
spheres
```

```
##      radius      volume  
## 1         3    339.2920  
## 2         4    804.2477  
## 3         5   1570.7963  
## 4         6   2714.3361  
## 5         7   4310.2651  
## 6         8   6433.9818  
## 7         9   9160.8842  
## 8        10  12566.3706  
## 9        11  16725.8393  
## 10       12  21714.6884  
## 11       13  27608.3162  
## 12       14  34482.1210  
## 13       15  42411.5008  
## 14       16  51471.8540  
## 15       17  61738.5788  
## 16       18  73287.0734  
## 17       19  86192.7360  
## 18       20 100530.9649
```

Exercise 6

```
# Identify which columns are factors  
factor_columns <- sapply(tinting, is.factor)
```

```
# Get levels of factor columns  
sapply(tinting[factor_columns], levels)
```

```
## $sex  
## [1] "f" "m"  
##  
## $tint  
## [1] "no" "lo" "hi"  
##  
## $target  
## [1] "locon" "hicon"  
##  
## $agegp  
## [1] "Younger" "Older"
```

```
# Check which columns are ordered factors
sapply(tinting, is.ordered)
```

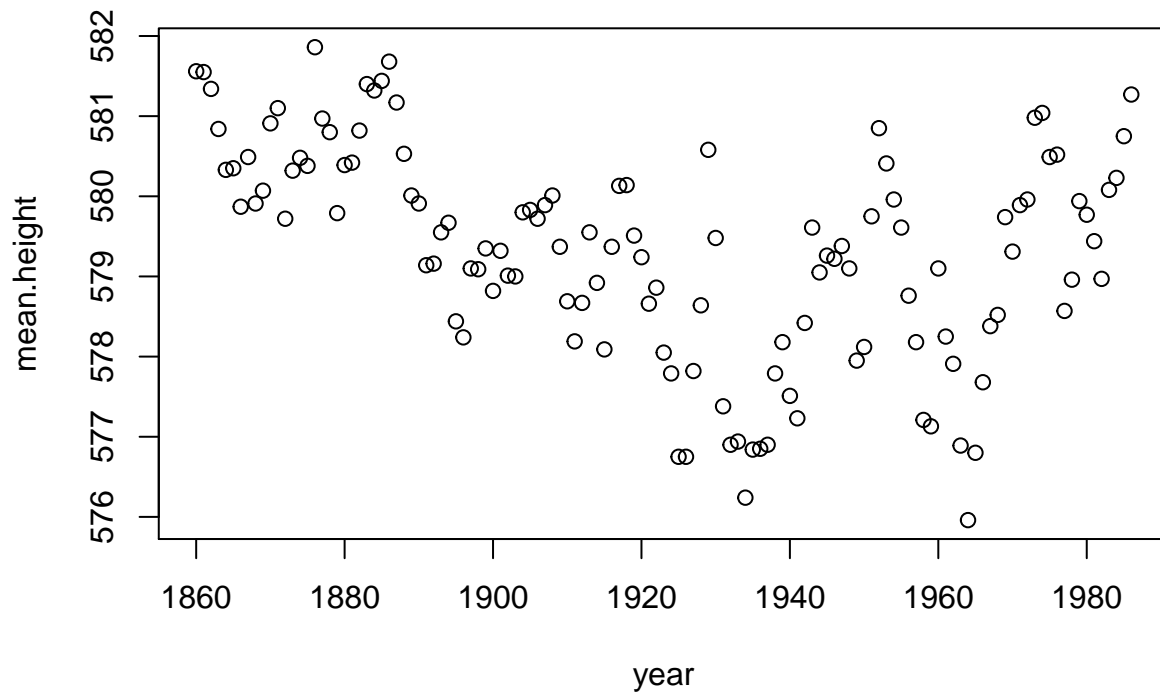
```
##   case    id   age  sex  tint target    it   csoa agegp
## FALSE FALSE FALSE FALSE  TRUE  FALSE FALSE  FALSE FALSE
```

Chapter 3 Exercises

Exercise 1

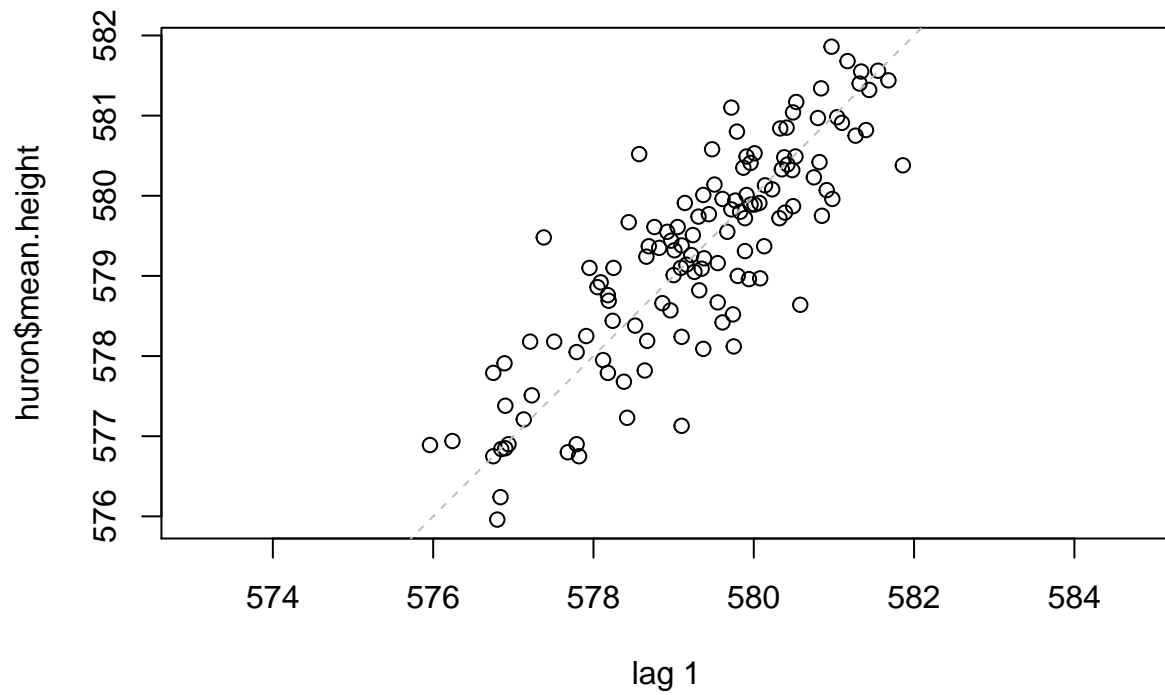
```
# 3.1(a)
plot(mean.height ~ year, data = huron)
```

```
# 3.1(b)
# Note: These labels won't necessarily show up in your saved output, which is fine.
identify(huron$year, huron$mean.height, labels = huron$year)
```



```
## integer(0)
```

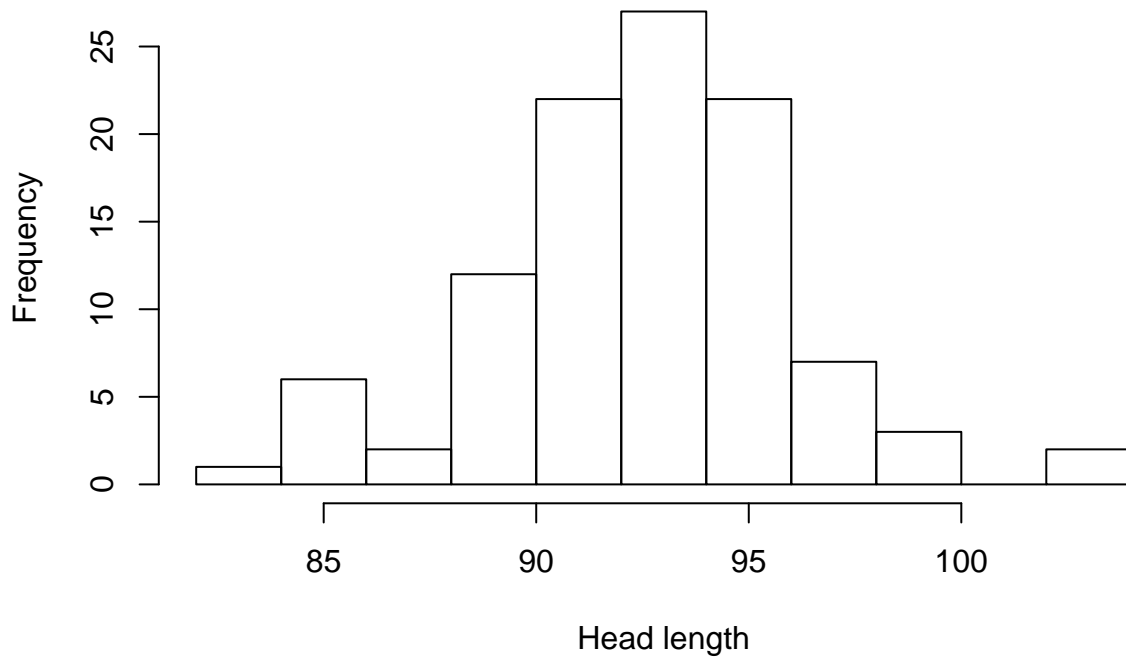
```
# 3.1(c)
# Note: If you read the docs for lag.plot(), with ?lag.plot or help(lag.plot),
# you'll see that you can't change certain graphical parameters, including xlab and ylab.
lag.plot(huron$mean.height, do.lines = FALSE)
```



Exercise 3

```
# 3.3(a)  
hist(possum$hdlngth, xlab = "Head length")
```

Histogram of possum\$hdlngth

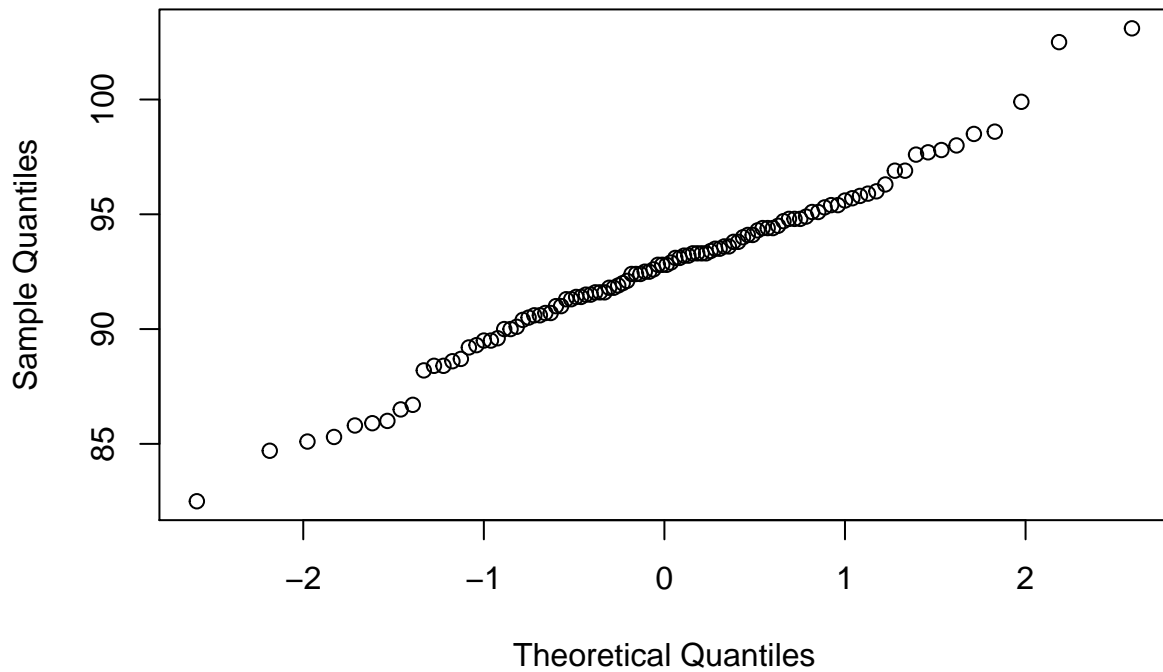


```
# 3.3(b)
# Note: the code in the problem is incorrectly given as stem(qqnorm(possum$hdlength))
stem(possum$hdlength)
```

```
##
## The decimal point is at the |
##
## 82 | 5
## 83 |
## 84 | 7
## 85 | 1389
## 86 | 057
## 87 |
## 88 | 24467
## 89 | 23556
## 90 | 001456677
## 91 | 00334455666889
## 92 | 014445568889
## 93 | 112233334556688
## 94 | 0113444578889
## 95 | 113446789
## 96 | 0399
## 97 | 678
## 98 | 056
## 99 | 9
## 100 |
## 101 |
## 102 | 5
## 103 | 1
```

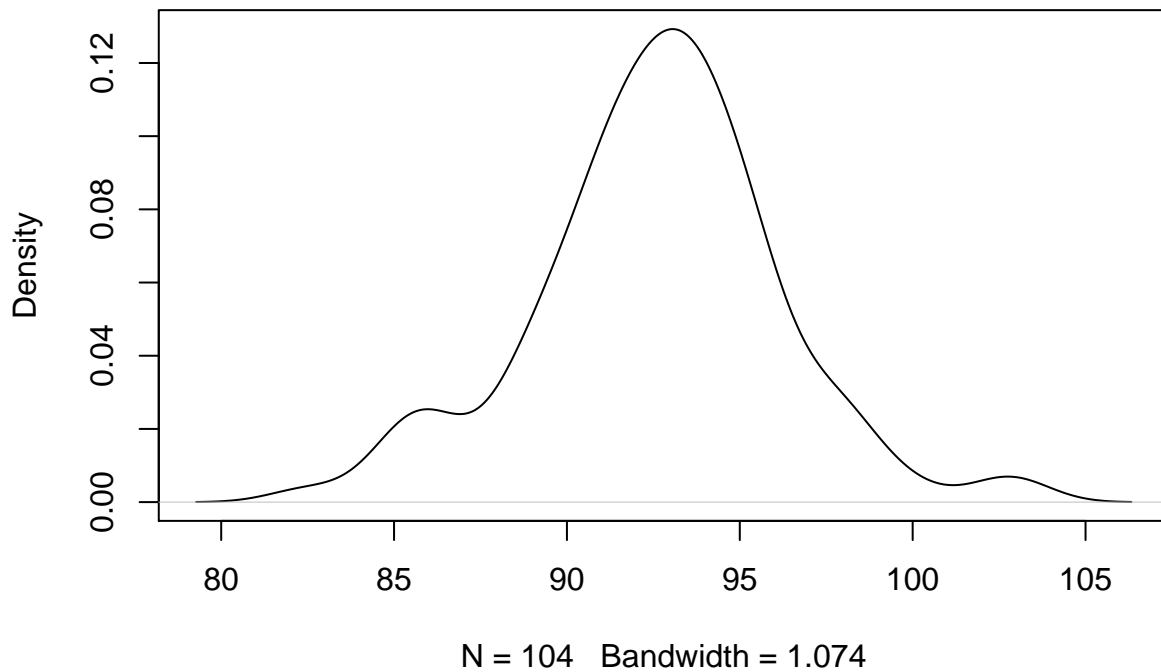
```
# 3.3(c)
qqnorm(possum$hdlength)
```


Normal Q-Q Plot



```
# 3.3(d)  
plot(density(possum$hdlength), main = "Estimated density of possum head length")
```

Estimated density of possum head length

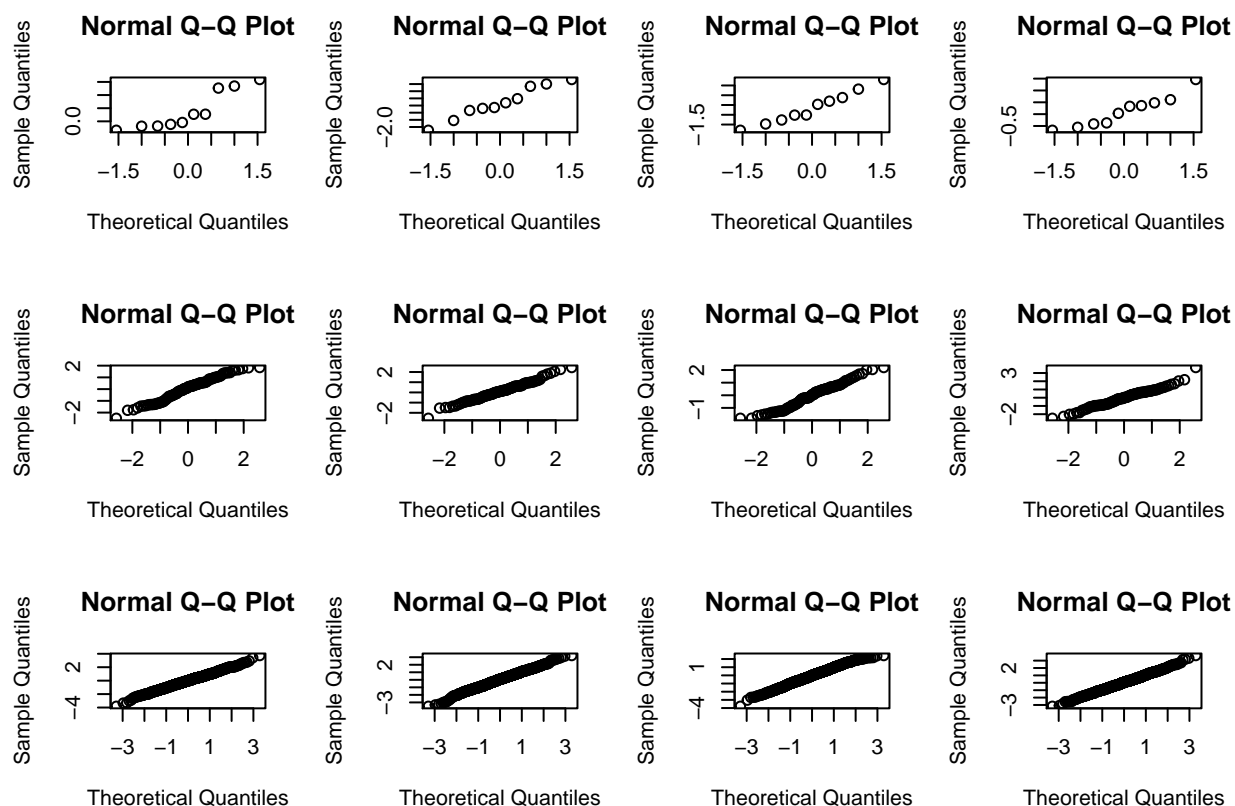


These displays provide slightly different types of information. The stem and leaf plot is essentially a sideways histogram, with the bin widths fixed at 1. It allows you to see all the individual data points. The histogram

provides more flexibility in that bin widths are adjustable, but it is sensitive to the bin width chosen. The density plot is a smoothed version of the histogram, which may give a better sense of the shape of the distribution, but it is sensitive to the parameters of kernel used to estimate the curve. The normal QQ plot makes it easy to see deviations from normality, but it is hard to see what the distribution is if it is not normal. The other displays make it much easier to see the shape of the distribution.

Exercise 5

```
par(mfrow = c(3, 4))
for (i in 1:4) {
  qqnorm(rnorm(10))
}
for (i in 1:4) {
  qqnorm(rnorm(100))
}
for (i in 1:4) {
  qqnorm(rnorm(1000))
}
```



```
# Typically you'll want to reset the grid after changing it.
par(mfrow = c(1, 1))
```

The larger the sample size, the more representative the sample (i.e. the closer to normal), and hence the straighter the line in the QQ plot.