Lab 5: Sampling Distributions

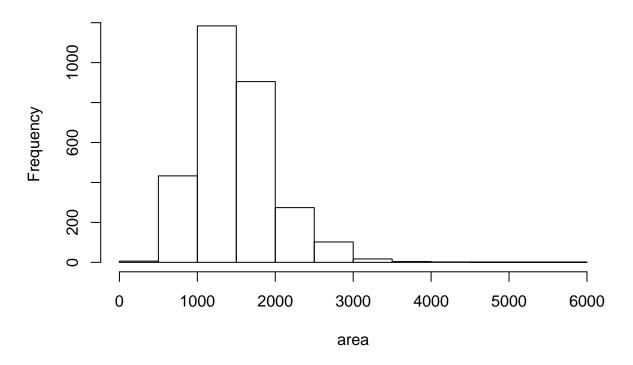
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```
download.file("http://www.openintro.org/stat/data/ames.RData", destfile = "ames.RData")
load("ames.RData")
area <- ames$Gr.Liv.Area
price <- ames$SalePrice</pre>
```

Exercise 1: Describe the shape, center (mean), and spread (standard deviation) of this population distribution

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 334 1126 1442 1500 1743 5642
hist(area, main = "Histogram of living area of the house")
```

Histogram of living area of the house



```
sd(area)
```

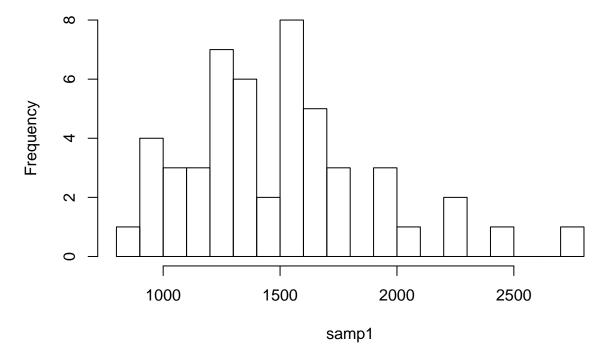
[1] 505.5089

-The Shape of this distribution is right skewed. The center of the distribution is 1500 and the standard deviation is 505.5

Exercise 2: Calculate summary statistics and plot a histogram of your sample. Describe the shape, center (mean), and spread (standard deviation) of this sample distribution. How does it compare to the population distribution you described in Exercise 1?

```
set.seed(001)
samp1 <- sample(area, 50)</pre>
summary(samp1)
##
      Min. 1st Qu.
                     Median
                                Mean 3rd Qu.
                                                 Max.
##
               1218
                                1491
                                                 2730
       864
                       1463
                                         1679
hist(samp1, main = "Histogram of living area of houses bases on a sample", breaks = 20)
```

Histogram of living area of houses bases on a sample



```
sd(samp1)
```

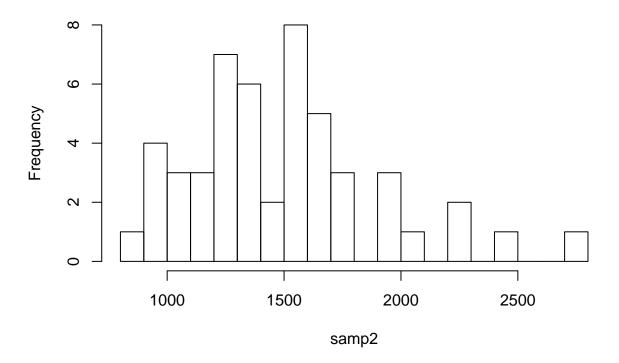
[1] 401.2159

⁻The histogram of the sample is also right-skewed, The center is 1491 and the standard deviation is 401.2. In comparison the sample mean is lower than the mean in the population distribution in exercise 1.the standard deviation is also lower, however, they are both right-skewed.

Exercise 3: Take a second sample, also of size 50, and name it samp2. How does the mean of samp2 compare with the mean of samp1? Suppose we took two more samples, one of size 100 and one of size 1000. Which would you think would provide a more accurate estimate of the population mean? Why?

```
set.seed(001)
samp2 <- sample(area,50)</pre>
samp100 <- sample(area, 100)</pre>
samp1000 <- sample (area, 1000)</pre>
set.seed(001)
summary(samp2)
##
      Min. 1st Qu.
                     Median
                                 Mean 3rd Qu.
                                                  Max.
##
               1218
                        1463
                                 1491
                                         1679
                                                  2730
hist(samp2, main = "Histogram of living area of houses bases on second sample", breaks = 20)
```

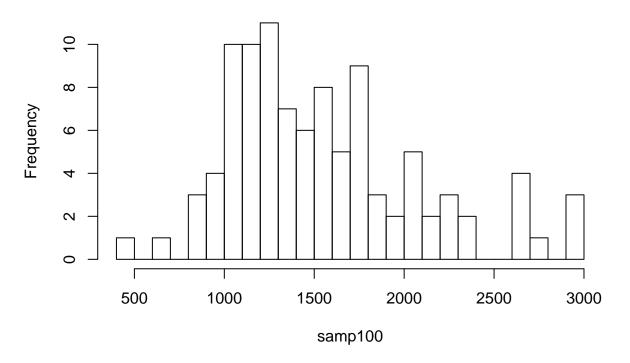
Histogram of living area of houses bases on second sample



```
sd(samp2)
## [1] 401.2159
summary(samp100)
## Min. 1st Qu. Median Mean 3rd Qu. Max.
```

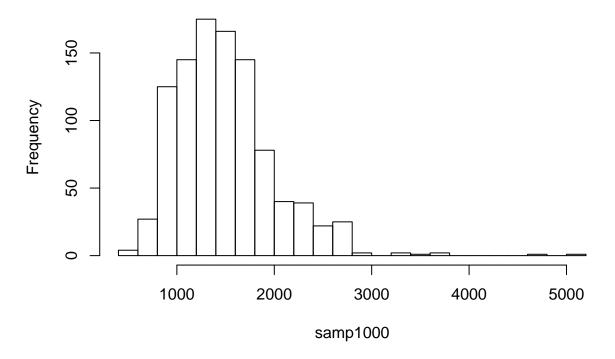
```
## 492 1178 1442 1557 1806 2956
hist(samp100, main = "Histogram of living area of houses bases on a sample of 100", breaks = 20)
```

Histogram of living area of houses bases on a sample of 100



```
sd(samp100)
## [1] 533.3047
summary(samp1000)
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 407 1128 1430 1492 1728 5095
hist(samp1000, main = "Histogram of living area of houses bases on a sample of 1000", breaks = 20)
```

Histogram of living area of houses bases on a sample of 1000



```
sd(samp1000)
```

[1] 505.5156

The second sample of size 50 is similar to the first sample, however the means are slightly different due to them being different samples and not the same seed. I think that the size of 1000 would provide a more accurate estimate for the mean because there are more samples to be summarized.

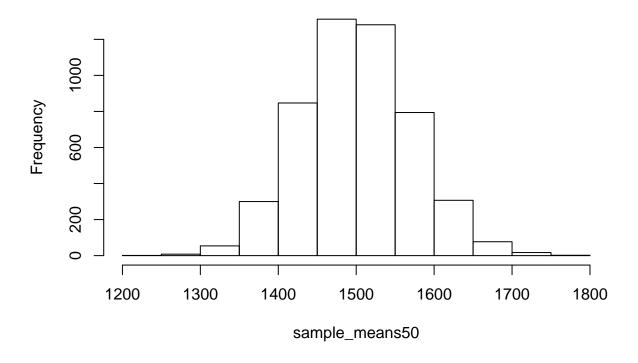
Exercise 4: How many elements are there in sample_means50? Describe the shape, center (mean), and spread (standard deviation) of the sampling distribution. How would you expect the sampling distribution to change if we instead collected 50,000 sample means?

```
set.seed(001)
sample_means50 <- rep(0, 5000)

for (i in 1:5000) {
    samp <- sample(area, 50)
    sample_means50[i] <- mean(samp)
}

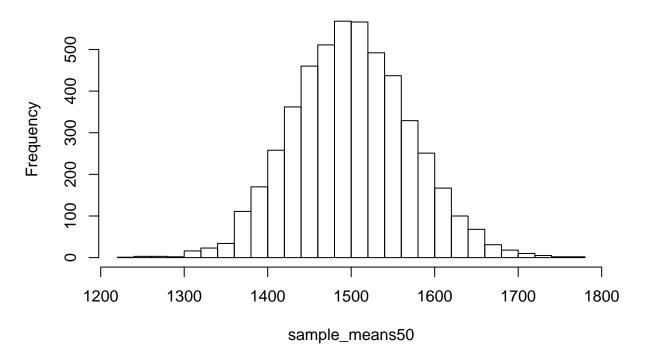
hist(sample_means50, main = "Histogram of Sample mean with size 50")</pre>
```

Histogram of Sample mean with size 50



hist(sample_means50, breaks = 25, main = "Histogram of Sample mean with size 50 and breaks of 25")

Histogram of Sample mean with size 50 and breaks of 25



```
summary(sample_means50)

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 1235 1452 1499 1501 1547 1766

sd(sample_means50)
```

[1] 70.51946

There are 5000 elements in the sample_means 50. The shape for this sampling distribution is uniform and symmetric. The center is 1501 which is similar to the previous sample means. The standard deviation however is much lower than the previous sample distributions at 70.519. If 50,000 sample means were collected instead, the standard deviation would be even smaller.

Exercise 5: When the sample size is larger, what happens to the center (mean) of the sampling distribution? What about the spread (standard deviation)?

```
sample_means10 <- rep(0, 5000)

sample_means100 <- rep(0, 5000)

for (i in 1:5000) {
    samp <- sample(area, 10)
    sample_means10[i] <- mean(samp)
    samp <- sample(area, 100)
    sample_means100[i] <- mean(samp)</pre>
```

```
par(mfrow = c(3, 1))

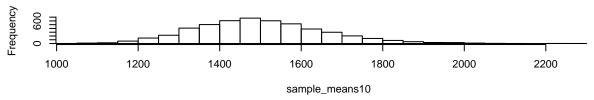
xlimits = range(sample_means10)

hist(sample_means10, breaks = 20, xlim = xlimits)

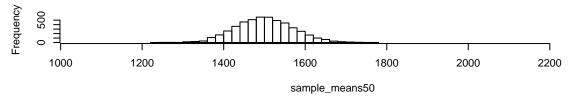
hist(sample_means50, breaks = 20, xlim = xlimits)

hist(sample_means100, breaks = 20, xlim = xlimits)
```

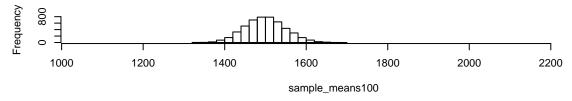
Histogram of sample_means10



Histogram of sample_means50



Histogram of sample means100

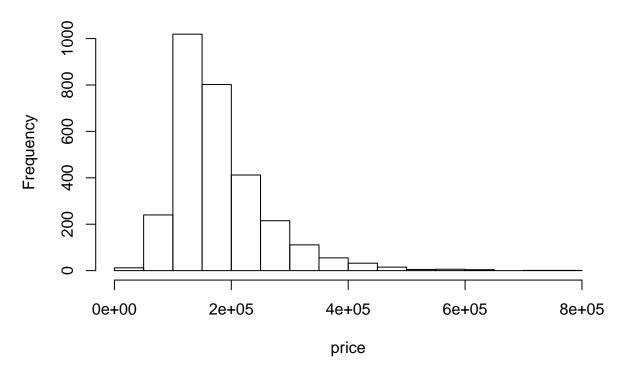


-When the sample size is larger, the mean stays the same dispite how large the sample size is. However, the spread of the sampling distribution gets smaller as the sample size increases.

1. Take a random sample of size 50 from price. Using this sample, what is your best point estimate of the population mean home price?

```
pricesamp <- sample(price, 50)
hist(price, breaks = 20, main = "Histogram of price sample")</pre>
```

Histogram of price sample



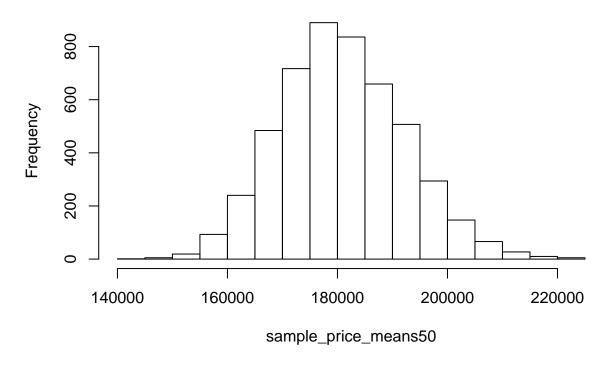
```
summary(price)
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 12789 129500 160000 180796 213500 755000
sd(price)
```

- ## [1] 79886.69
 - This population distribution is right-skewed and the ebst point estimate of the population is 180796
- 2. Since you have access to the population, simulate the sampling distribution for the sample mean of home price by taking 5000 samples from the population of size 50 and computing 5000 price sample means. Store these means in a vector called sample_price_means50. Plot the data, then describe the shape of this simulated sampling distribution. Based on this simulated sampling distribution, what would you guess the mean home price of the population to be?

```
set.seed(001)
sample_price_means50 <- rep(0,5000)

for(i in 1:5000){
samp<-sample(price,50)
sample_price_means50[i]<-mean(samp)
}</pre>
```

Histogram for sample price with n = 50



```
summary(sample_price_means50)

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 142663 173051 180294 180867 188314 224643

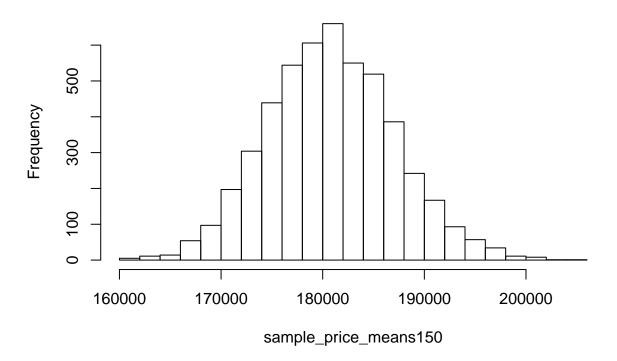
sd(sample_price_means50)
```

- ## [1] 11239.13
- -The shape of this population distribution is uniform and symmetrical. The mean home price of the population would then be 180867.
- 3. Change your sample size from 50 to 150, and then generate a simulated sampling distribution using the same method as above. Store these means in a new vector called sample_price_means150. Compare and contrast the shape, center (mean), and spread (standard deviation) of your simulated sampling distributions for n = 50 and n = 150. Based on your simulated sampling distribution for samples of size n = 150, what would you guess to be the mean sale price of homes in Ames? Finally, calculate and report the actual population mean.

```
set.seed(001)
sample_price_means150 <- rep(0,5000)</pre>
```

```
for(i in 1:5000){
samp<-sample(price,150)
sample_price_means150[i]<-mean(samp)
}
hist(sample_price_means150, breaks = 20, main = "Histogram for sample price with n = 150")</pre>
```

Histogram for sample price with n = 150



```
summary(sample_price_means150)
##
      Min. 1st Qu.
                    Median
                              Mean 3rd Qu.
                                               Max.
   160798 176505 180728
                           180823
                                   184994
                                            205015
summary(price)
##
      Min. 1st Qu.
                    Median
                              Mean 3rd Qu.
                                               Max.
     12789 129500
                    160000
                            180796
                                    213500
                                            755000
sd(sample_price_means150)
```

[1] 6272.882

• Again the shape of this distribution is uniform and symmetrical. The center is 180823, which is very similar to the previous sample. The spread is 6273 which is much lower than the sampling distribution for n = 50. The mean sale price for homes in Ames would be 180823. The actual population mean is very similar at 180796.

4. Of the sampling distributions from #2 and #3, which has a smaller spread (standard deviation)? If we're concerned with making estimates that are more often close to the true value, would we prefer a sampling distribution with a large or small spread? Explain your reasoning.

⁻The sampling distribution with n=150 has a smaller spread. If we are concerned with making estimates we would perfer to have the sampling distribution of n=150 due to the fact that a smaller spread means that it is closer to the true value.