Homework 7

1. Textbook 6.10

Here is my code for this section:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
url =
column names = ['Sex', 'Length', 'Diameter', 'Height', 'Whole
abalone data = pd.read csv(url, names=column names)
length data = abalone data['Length']
population median = np.median(length data)
print(f"Population Median: {population median}")
def bootstrap confidence interval(data, n samples, sample size,
confidence level=0.90):
  boot medians = []
   for _ in range(n samples):
       sample = np.random.choice(data, size=sample size,
replace=True)
      boot medians.append(np.median(sample))
   lower percentile = (1 - confidence level) / 2 * 100
   upper percentile = (1 + confidence level) / 2 * 100
   lower bound = np.percentile(boot medians, lower percentile)
   upper bound = np.percentile(boot medians, upper percentile)
   return boot_medians, (lower_bound, upper_bound)
n \text{ samples a} = 2000
sample size a = 100
```

```
boot medians a, ci a = bootstrap confidence interval(length data,
n samples a, sample size a)
print(f"90% CI for part (a): {ci a}")
inside ci a = np.sum((ci a[0] <= population median) &</pre>
(population_median <= ci_a[1])) / n_samples_a</pre>
print(f"Fraction for part (a): {inside ci a}")
plt.hist(boot medians a, bins=30, edgecolor='k')
plt.title('Bootstrap Sample Medians - Part (a) (100 Records)')
plt.xlabel('Median')
plt.ylabel('Frequency')
plt.show()
n samples c = 2000
sample size c = 10
boot medians c, ci c = bootstrap confidence interval(length data,
n samples c, sample size c)
print(f"90% CI for part (c): {ci c}")
inside ci c = np.sum((ci c[0] <= population median) &
(population_median <= ci_c[1])) / n_samples_c</pre>
print(f"Fraction for part (c): {inside ci c}")
plt.hist(boot medians c, bins=30, edgecolor='k')
plt.title('Bootstrap Sample Medians - Part (c) (10 Records)')
plt.xlabel('Median')
plt.ylabel('Frequency')
plt.show()
```

The output of that code is:

Population Median: 0.545

90% CI for part (a): (0.515, 0.56749999999999)

Fraction for part (a): 0.0005

90% CI for part (c): (0.457375, 0.61)

Fraction for part (c): 0.0005c

First, let's calculate the standard error of the Mean.

$$SEM = \frac{10}{\sqrt{50}} \approx 1.41$$

Then lets calculate the Z score:

$$Z = \frac{200-169}{1.41} \approx 21.99$$

With a Z score so high, the chances of the average being greater than 200cm is basically 0.

First, let's calculate the standard error of the Mean.

$$SEM = \frac{0.7}{\sqrt{30}} \approx 0.128$$

Then let's calculate the Z score:

$$Z = \frac{4-5}{0.128} \approx -7.81$$

The chances of the mean being less than 4k are very close to 0.

$$2.55 \times 10^{-15}$$

b. b

First, let's calculate the standard error of the Mean.

$$SEM = \frac{0.7}{\sqrt{300}} \approx 0.04$$

Then let's calculate the Z score:

$$Z = \frac{4-5}{0.04} \approx -25$$

$$1.82 \times 10^{-135}$$

c. c

The reason the chances are smaller for larger sample sizes is that in a smaller sample size, it is more realistic to find a couple of outliers from the usual average and make an unusually high mean. When the sample size is larger the chances of that happening diminish.

The p value in terms of the integral is equal to

$$2 \times \int_{30}^{\infty} \frac{1}{\sqrt{2\pi}} e^{\frac{-z^2}{2}} dz$$

Evaluating this using wolfram I get 2.06×10^{-198}

Since the p value is extremely close to 0 it is very unlikely that the mean of the Parktown Prawns is 10cm. The true mean is probably closer to 7cm.

The expected number of boys (E[X]) for this problem is:

$$E[X] = 2009 \times 0.5 = 1004.5$$

The standard error is:

$$SE = \sqrt{2009 \times 0.5 \times 0.5} \approx 22.42$$

The Z score is:

$$\frac{983-1004.3}{22.42} \approx -0.96$$

Now lets look at his hypothesis. The P value integral is:

$$\int_{0.96}^{\infty} \frac{1}{\sqrt{2\pi}} e^{\frac{-z^2}{2}} dz$$

Using Wolfram Alpha we find the p-value to be 0.3372. Since the p value is fairly large, his hypothesis could be true that boys are born with probability 0.5.