Capstone-1

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Import all Libraries

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from scipy.stats import zscore, pearsonr, spearmanr
```

Download and read dataset

```
male_url = "https://raw.githubusercontent.com/gagolews/teaching-data/master/marek/nhanes_adu
female_url = "https://raw.githubusercontent.com/gagolews/teaching-data/master/marek/nhanes_a
male = np.genfromtxt(male_url, delimiter=",", skip_header=1, dtype=float)
female = np.genfromtxt(female_url, delimiter=",", skip_header=1, dtype=float)

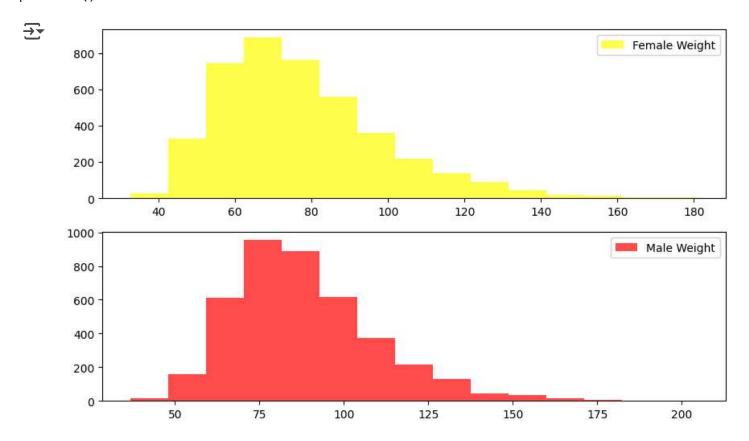
print(f"Male data shape: {male.shape}")
print(f"Female data shape: {female.shape}")

Ale data shape: (4082, 7)
Female data shape: (4222, 7)
```

Compare the weight distributions of male and female participants

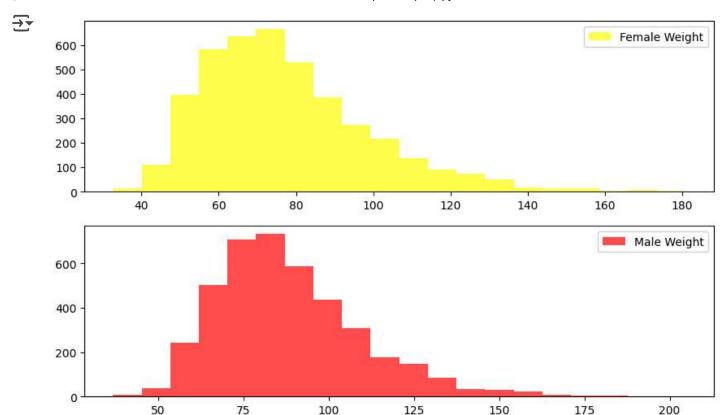
```
plt.figure(figsize=(10, 6))
plt.subplot(2, 1, 1)
plt.hist(female[:, 0], bins=15, alpha=0.7, color='yellow', label='Female Weight')
plt.legend()
plt.subplot(2, 1, 2)
```

```
plt.hist(male[:, 0], bins=15, alpha=0.7, color='red', label='Male Weight')
plt.legend()
plt.show()
```



Plot Histograms of Weights

```
plt.figure(figsize=(10, 6))
plt.subplot(2, 1, 1)
plt.hist(female[:, 0], bins=20, alpha=0.7, color='yellow', label='Female Weight')
plt.legend()
plt.subplot(2, 1, 2)
plt.hist(male[:, 0], bins=20, alpha=0.7, color='red', label='Male Weight')
plt.legend()
plt.show()
```



Check for NaN or invalid values

```
print("Male dataset preview:\n", male[:5])
print("Female dataset preview:\n", female[:5])
print("Any NaNs in Male Data:", np.isnan(male).any())
print("Any NaNs in Female Data:", np.isnan(female).any())
    Male dataset preview:
      [[ nan
                nan
                      nan
                            nan
                                         nan
                                               nan]
                                  nan
      [ 98.8 182.3
                    42.
                          40.1
                                38.2 108.2 120.4]
                                 30.2 94.5 86.8]
      74.3 184.2
                          41.
                    41.1
      [103.7 185.3
                    47.
                          44.
                                 32.
                                     107.8 109.6]
      [ 86. 167.8
                    39.5
                          38.4
                                29.
                                     106.4 108.3]]
     Female dataset preview:
      [[ nan
                nan
                      nan
                            nan
                                  nan
                                         nan
                                               nan]
      [ 97.1 160.2
                    34.7
                          40.8 35.8 126.1 117.9]
      [ 91.1 152.7
                    33.5
                          33.
                                 38.5 125.5 103.1]
      [ 73. 161.2
                    37.4
                          38.
                                 31.8 106.2 92. ]
      [ 61.7 157.4
                    38.
                                29.
                                             90.5]]
                          34.7
                                      101.
     Any NaNs in Male Data: True
     Any NaNs in Female Data: True
```

Remove NaN data

```
male = male[~np.isnan(male).any(axis=1)]
female = female[~np.isnan(female).any(axis=1)]
```

Checking Data post clearing

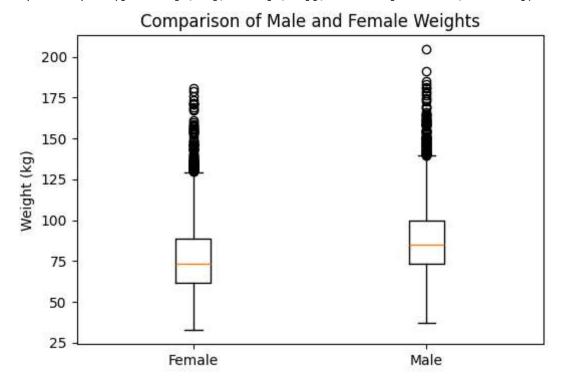
```
print("Male dataset preview:\n", male[:5])
print("Female dataset preview:\n", female[:5])
print("Any NaNs in Male Data:", np.isnan(male).any())
print("Any NaNs in Female Data:", np.isnan(female).any())
→ Male dataset preview:
     [[ 98.8 182.3 42.
                        40.1 38.2 108.2 120.4]
     74.3 184.2 41.1 41.
                              30.2 94.5 86.8]
     [103.7 185.3 47. 44.
                              32. 107.8 109.6]
     [ 86. 167.8 39.5 38.4 29.
                                   106.4 108.3
     [ 99.4 181.6 40.4 39.9 36.
                                   120.2 107. ]]
    Female dataset preview:
     [[ 97.1 160.2 34.7 40.8 35.8 126.1 117.9]
     [ 91.1 152.7 33.5 33.
                              38.5 125.5 103.1]
     [ 73. 161.2 37.4 38.
                              31.8 106.2 92. ]
     [ 61.7 157.4 38.
                        34.7 29. 101.
                                         90.5]
     [ 55.4 154.6 34.6 34.
                              28.3 92.5 73.2]]
    Any NaNs in Male Data: False
    Any NaNs in Female Data: False
```

Boxplot for Weight Comparison

```
plt.figure(figsize=(6, 4))
plt.boxplot([female[:, 0], male[:, 0]], labels=['Female', 'Male'])
plt.ylabel("Weight (kg)")
plt.title("Comparison of Male and Female Weights")
plt.show()
```

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<ipython-input-8-5fd59ffc8dc8>:2: MatplotlibDeprecationWarning: The 'labels' parameter c
 plt.boxplot([female[:, 0], male[:, 0]], labels=['Female', 'Male'])



Compute Basic Numerical Aggregates

```
def describe_data(data, label):
   print(f"Statistics for {label}:")
   print(f"Mean: {np.mean(data):.2f}, Median: {np.median(data):.2f}")
   print(f"Standard Deviation: {np.std(data):.2f}")
   print(f"Min: {np.min(data)}, Max: {np.max(data)}")
   print("----")
describe_data(female[:, 0], "Female Weights")
describe_data(male[:, 0], "Male Weights")
→ Statistics for Female Weights:
    Mean: 77.40, Median: 73.60
    Standard Deviation: 21.54
    Min: 32.6, Max: 180.9
    -----
    Statistics for Male Weights:
    Mean: 88.36, Median: 85.00
    Standard Deviation: 21.42
    Min: 36.8, Max: 204.6
     _____
```

Compute and Append BMI

```
def compute_bmi(data):
    weight = data[:, 0]
    height_m = data[:, 1] / 100
    bmi = weight / (height_m ** 2)
    return np.column_stack((data, bmi))

female = compute_bmi(female)
male = compute_bmi(male)
```

Standardize Female Data

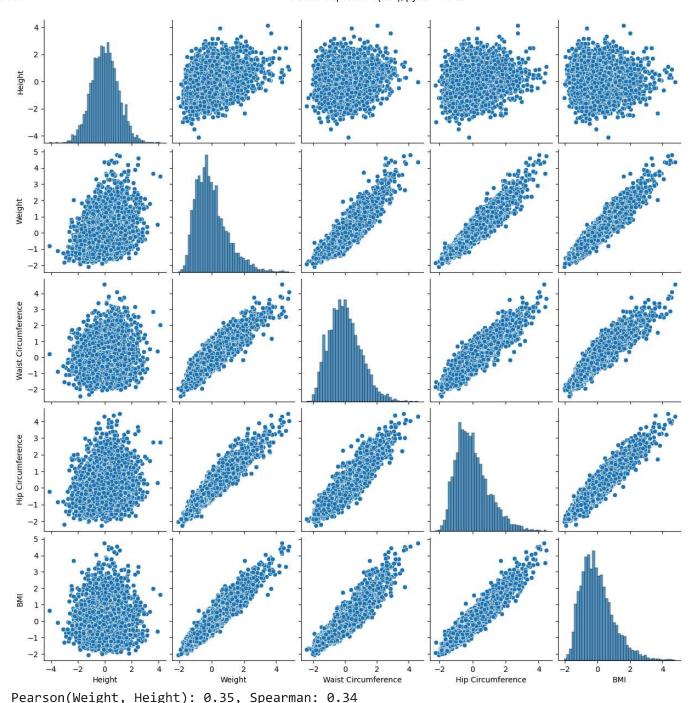
```
zfemale = np.apply_along_axis(zscore, 0, female)
```

Scatterplot Matrix and Correlations

```
df_zfemale = pd.DataFrame(zfemale, columns=["Weight", "Height", "Arm Length", "Leg Length",
sns.pairplot(df_zfemale[["Height", "Weight", "Waist Circumference", "Hip Circumference", "BN
plt.show()

for col1 in df_zfemale.columns:
    for col2 in df_zfemale.columns:
        if col1 != col2:
            pearson_corr, _ = pearsonr(df_zfemale[col1], df_zfemale[col2])
            spearman_corr, _ = spearmanr(df_zfemale[col1], df_zfemale[col2])
            print(f"Pearson({col1}, {col2}): {pearson_corr:.2f}, Spearman: {spearman_corr:.2}
```





Pearson(Weight, Arm Length): 0.55, Spearman: 0.54 Pearson(Weight, Leg Length): 0.19, Spearman: 0.20 Pearson(Weight, Arm Circumference): 0.91, Spearman: 0.91 Pearson(Weight, Hip Circumference): 0.95, Spearman: 0.95 Pearson(Weight, Waist Circumference): 0.90, Spearman: 0.90 Pearson(Weight, BMI): 0.95, Spearman: 0.94 Pearson(Height, Weight): 0.35, Spearman: 0.34 Pearson(Height, Arm Length): 0.67, Spearman: 0.67 Pearson(Height, Leg Length): 0.66, Spearman: 0.65 Pearson(Height, Arm Circumference): 0.15, Spearman: 0.14 Pearson(Height, Hip Circumference): 0.20, Spearman: 0.21 Pearson(Height, Waist Circumference): 0.13, Spearman: 0.11 Pearson(Height, BMI): 0.03, Spearman: 0.02 Pearson(Arm Length, Weight): 0.55, Spearman: 0.54 Pearson(Arm Length, Height): 0.67, Spearman: 0.67 Pearson(Arm Length, Leg Length): 0.48, Spearman: 0.46

```
Pearson(Arm Length, Arm Circumterence): 0.45, Spearman: 0.42
Pearson(Arm Length, Hip Circumference): 0.46, Spearman: 0.45
Pearson(Arm Length, Waist Circumference): 0.43, Spearman: 0.40
Pearson(Arm Length, BMI): 0.36, Spearman: 0.34
Pearson(Leg Length, Weight): 0.19, Spearman: 0.20
Pearson(Leg Length, Height): 0.66, Spearman: 0.65
Pearson(Leg Length, Arm Length): 0.48, Spearman: 0.46
Pearson(Leg Length, Arm Circumference): 0.08, Spearman: 0.08
Pearson(Leg Length, Hip Circumference): 0.10, Spearman: 0.12
Pearson(Leg Length, Waist Circumference): -0.03, Spearman: -0.04
Pearson(Leg Length, BMI): -0.01, Spearman: -0.02
Pearson(Arm Circumference, Weight): 0.91, Spearman: 0.91
Pearson(Arm Circumference, Height): 0.15, Spearman: 0.14
Pearson(Arm Circumference, Arm Length): 0.45, Spearman: 0.42
Pearson(Arm Circumference, Leg Length): 0.08, Spearman: 0.08
Pearson(Arm Circumference, Hip Circumference): 0.87, Spearman: 0.87
Pearson(Arm Circumference, Waist Circumference): 0.85, Spearman: 0.84
Pearson(Arm Circumference, BMI): 0.92, Spearman: 0.92
Pearson(Hip Circumference, Weight): 0.95, Spearman: 0.95
Pearson(Hip Circumference, Height): 0.20, Spearman: 0.21
Pearson(Hip Circumference, Arm Length): 0.46, Spearman: 0.45
Pearson(Hip Circumference, Leg Length): 0.10, Spearman: 0.12
Pearson(Hip Circumference, Arm Circumference): 0.87, Spearman: 0.87
Pearson(Hip Circumference, Waist Circumference): 0.90, Spearman: 0.89
Pearson(Hip Circumference, BMI): 0.94, Spearman: 0.93
Pearson(Waist Circumference, Weight): 0.90, Spearman: 0.90
Pearson(Waist Circumference, Height): 0.13, Spearman: 0.11
Pearson(Waist Circumference, Arm Length): 0.43, Spearman: 0.40
Pearson(Waist Circumference, Leg Length): -0.03, Spearman: -0.04
Pearson(Waist Circumference, Arm Circumference): 0.85, Spearman: 0.84
Pearson(Waist Circumference, Hip Circumference): 0.90, Spearman: 0.89
Pearson(Waist Circumference, BMI): 0.92, Spearman: 0.92
Pearson(BMI, Weight): 0.95, Spearman: 0.94
Pearson(BMI, Height): 0.03, Spearman: 0.02
Pearson(BMI, Arm Length): 0.36, Spearman: 0.34
Pearson(BMI, Leg Length): -0.01, Spearman: -0.02
Pearson(BMI, Arm Circumference): 0.92, Spearman: 0.92
Pearson(BMI, Hip Circumference): 0.94, Spearman: 0.93
Pearson(BMI, Waist Circumference): 0.92, Spearman: 0.92
```

Compute Waist-to-Height and Waist-to-Hip Ratios

```
# ## **9. Compute Waist-to-Height and Waist-to-Hip Ratios**

def compute_ratios(data):
    waist_to_height = data[:, 6] / data[:, 1]
    waist_to_hip = data[:, 6] / data[:, 5]
    return np.column_stack((data, waist_to_height, waist_to_hip))

female = compute_ratios(female)

male = compute ratios(male)
```

Boxplot Comparison of Ratios