

Question 1

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Files

- sample_data
- original_diabetes.xlsx

```
import pandas as pd
import matplotlib.pyplot as plt
import numpy as np

def calculate_mean(data):
    return np.nanmean(data)

def calculate_variance(data):
    return np.nanvar(data)

def calculate_std_dev(data):
    return np.nanstd(data)

def calculate_z_scores(data):
    mean = calculate_mean(data)
    std_dev = calculate_std_dev(data)
    z_scores = [(x - mean) / std_dev if not np.isnan(x) else np.nan for x in data]
    return z_scores

def calculate_quartiles(data):
    data_sorted = np.sort(data[~np.isnan(data)])
    n = len(data_sorted)
    q2 = np.nanmedian(data)
    q1 = data_sorted[n // 4]
    q3 = data_sorted[(3 * n) // 4]
    return q1, q2, q3

data = pd.read_excel("original_diabetes.xlsx")

# Extracting the "Glucose" and "BloodPressure" columns
glucose_data = data["Glucose"]
blood_pressure_data = data["BloodPressure"]

# Calculating statistics for "Glucose"
glucose_mean = calculate_mean(glucose_data)
glucose_variance = calculate_variance(glucose_data)
glucose_std_dev = calculate_std_dev(glucose_data)
glucose_z_scores = calculate_z_scores(glucose_data)
```

Table of contents

- Getting started
- Data science
- Machine learning
- More Resources
- Featured examples
- Section

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```
q1, median, q3 = calculate_quartiles(glucose_data)

# Calculating statistics for "BloodPressure"
bp_mean = calculate_mean(blood_pressure_data)
bp_variance = calculate_variance(blood_pressure_data)
bp_std_dev = calculate_std_dev(blood_pressure_data)
bp_z_scores = calculate_z_scores(blood_pressure_data)
bp_q1, bp_median, bp_q3 = calculate_quartiles(blood_pressure_data)

# Print the statistics for "Glucose"
print("Statistics for Glucose:")
print(f"Mean: {glucose_mean}")
print(f"Variance: {glucose_variance}")
print(f"Standard Deviation: {glucose_std_dev}")
print(f"Z-Scores: {glucose_z_scores}")
print(f"Q1: {q1}, Median (Q2): {median}, Q3: {q3}")
print()

# Print the statistics for "BloodPressure"
print("Statistics for BloodPressure:")
print(f"Mean: {bp_mean}")
print(f"Variance: {bp_variance}")
print(f"Standard Deviation: {bp_std_dev}")
print(f"Z-Scores: {bp_z_scores}")
print(f"Q1: {bp_q1}, Median (Q2): {bp_median}, Q3: {bp_q3}")
print()

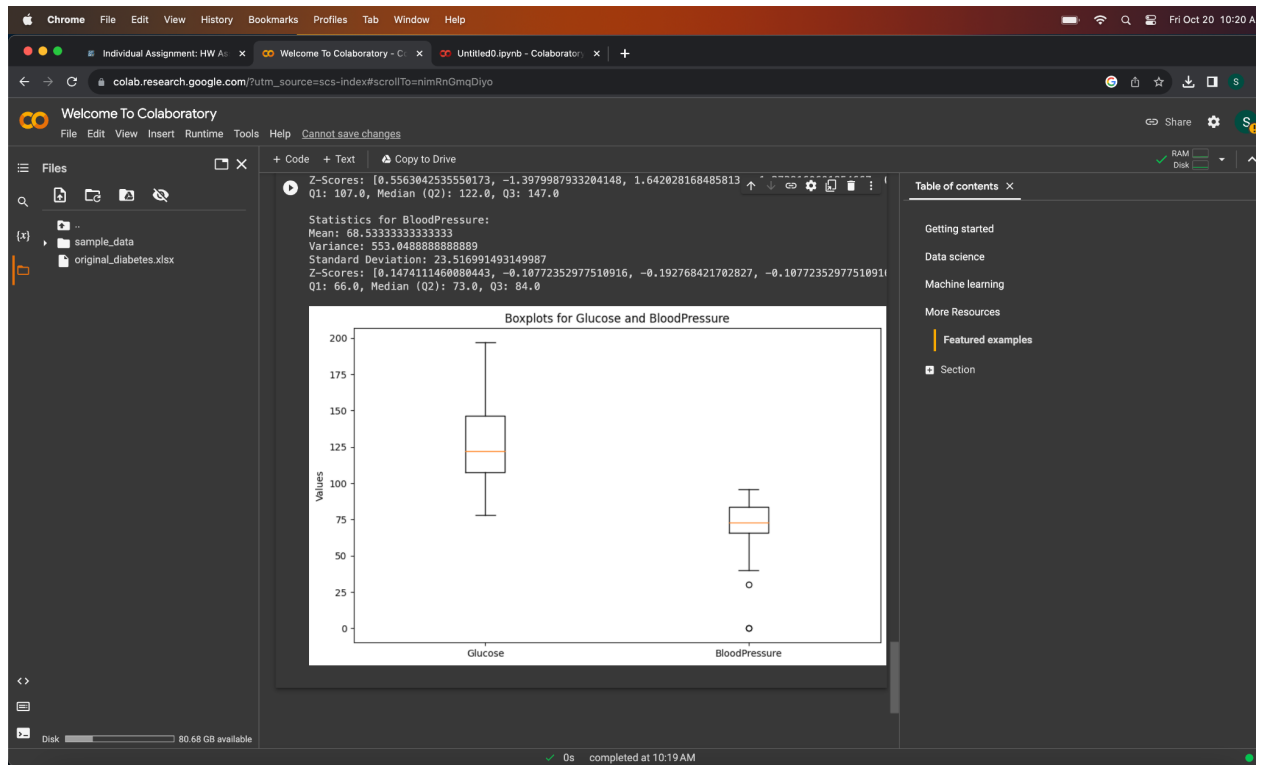
# Plotting boxplots for "Glucose" and "BloodPressure" in one frame
plt.figure(figsize=(10, 6))
plt.boxplot([glucose_data[~np.isnan(glucose_data)], blood_pressure_data[~np.isnan(blood_pressure_data)]], labels=["Glucose", "BloodPressure"])
plt.title("Boxplots for Glucose and BloodPressure")
plt.ylabel("Values")
plt.show()

Statistics for Glucose:
Mean: 130.06666666666666
Variance: 1039.1955555555555
Standard Deviation: 32.2365619875603
Z-Scores: [0.5563042535550173, -1.3979987933204148, 1.642028168485813, -1.2739160601854667, ...]
```

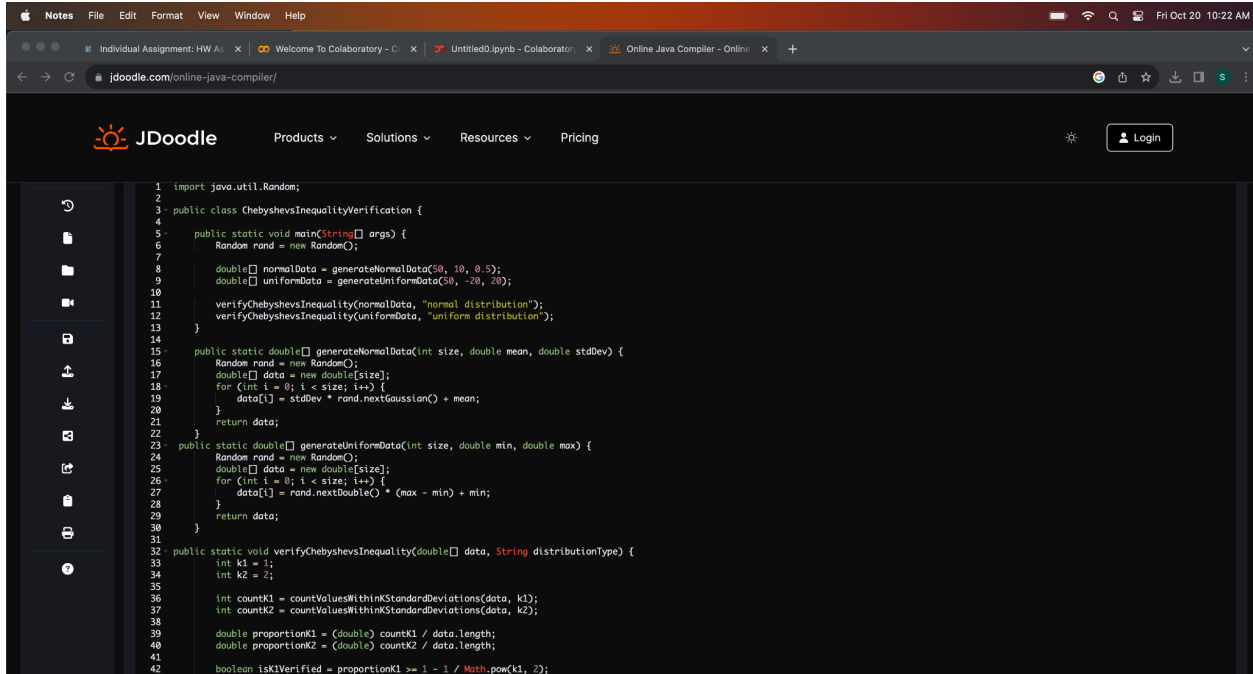
Table of contents

- Getting started
- Data science
- Machine learning
- More Resources
- Featured examples
- Section

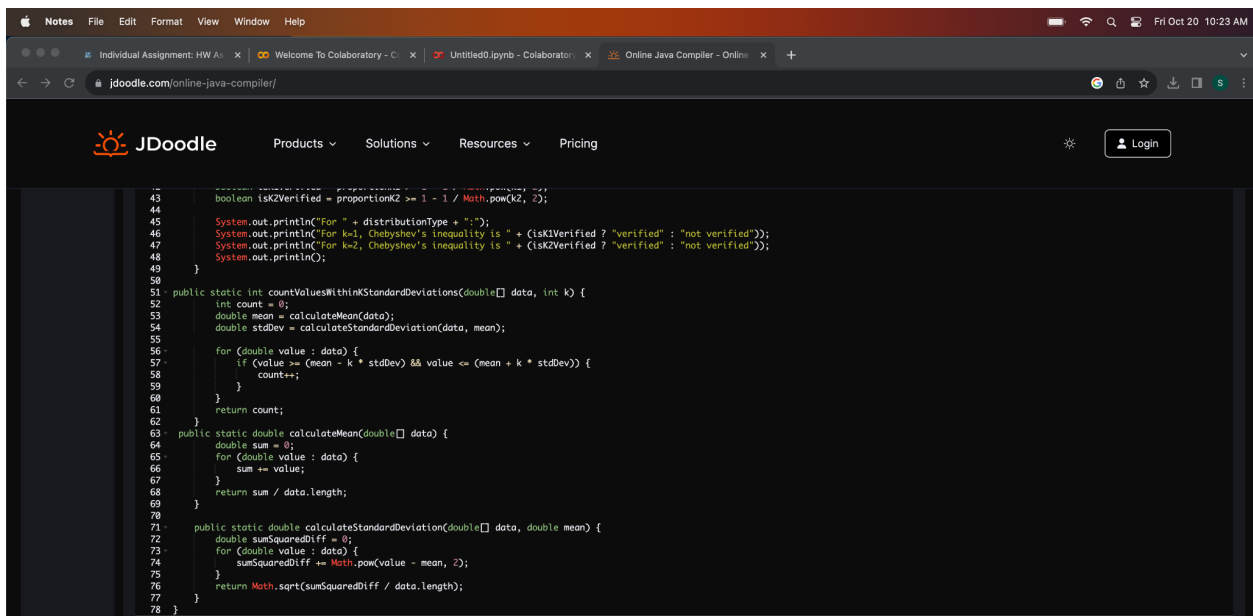
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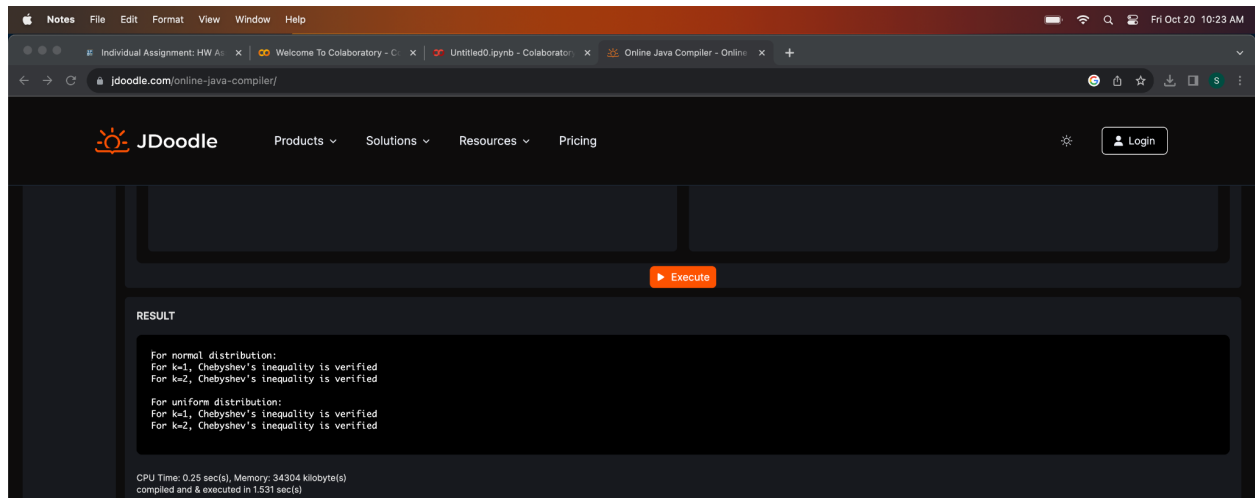
Question 2



```
1 import java.util.Random;
2
3 public class ChebyshevsInequalityVerification {
4
5     public static void main(String[] args) {
6         Random rand = new Random();
7
8         double[] normalData = generateNormalData(50, 10, 0.5);
9         double[] uniformData = generateUniformData(50, -20, 20);
10
11         verifyChebyshevsInequality(normalData, "normal distribution");
12         verifyChebyshevsInequality(uniformData, "uniform distribution");
13     }
14
15     public static double[] generateNormalData(int size, double mean, double stdDev) {
16         Random rand = new Random();
17         double[] data = new double[size];
18         for (int i = 0; i < size; i++) {
19             data[i] = stdDev * rand.nextGaussian() + mean;
20         }
21         return data;
22     }
23
24     public static double[] generateUniformData(int size, double min, double max) {
25         Random rand = new Random();
26         double[] data = new double[size];
27         for (int i = 0; i < size; i++) {
28             data[i] = rand.nextDouble() * (max - min) + min;
29         }
30         return data;
31     }
32
33     public static void verifyChebyshevsInequality(double[] data, String distributionType) {
34         int k1 = 1;
35         int k2 = 2;
36
37         int countK1 = countValuesWithinKStandardDeviations(data, k1);
38         int countK2 = countValuesWithinKStandardDeviations(data, k2);
39
40         double proportionK1 = (double) countK1 / data.length;
41         double proportionK2 = (double) countK2 / data.length;
42
43         boolean isK1Verified = proportionK1 >= 1 - 1 / Math.pow(k1, 2);
```



```
43         boolean isK2Verified = proportionK2 >= 1 - 1 / Math.pow(k2, 2);
44
45         System.out.println("For " + distributionType + ":");
46         System.out.println("For k=1, Chebyshev's inequality is " + (isK1Verified ? "verified" : "not verified"));
47         System.out.println("For k=2, Chebyshev's inequality is " + (isK2Verified ? "verified" : "not verified"));
48         System.out.println();
49     }
50
51     public static int countValuesWithinKStandardDeviations(double[] data, int k) {
52         int count = 0;
53         double mean = calculateMean(data);
54         double stdDev = calculateStandardDeviation(data, mean);
55
56         for (double value : data) {
57             if (value >= (mean - k * stdDev) && value <= (mean + k * stdDev)) {
58                 count++;
59             }
60         }
61         return count;
62     }
63
64     public static double calculateMean(double[] data) {
65         double sum = 0;
66         for (double value : data) {
67             sum += value;
68         }
69         return sum / data.length;
70     }
71
72     public static double calculateStandardDeviation(double[] data, double mean) {
73         double sumSquaredDiff = 0;
74         for (double value : data) {
75             sumSquaredDiff += Math.pow(value - mean, 2);
76         }
77         return Math.sqrt(sumSquaredDiff / data.length);
78     }
79 }
```



Question 3

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```
[5] #Question 3
import numpy as np
from scipy.stats import linregress
import matplotlib.pyplot as plt

X = np.array([2, 3, 4, 5, 6, 7, 8, 9, 10])
Y = np.array([30, 25, 95, 115, 265, 325, 570, 700, 1085, 1300])

slope, intercept, r_value, p_value, std_err = linregress(X, Y)
correlation_coefficient = r_value

predicted_Y = slope * X + intercept

plt.scatter(X, Y, label='Data Points')
plt.plot(X, predicted_Y, color='red', label='Linear Fit')
plt.xlabel('X')
plt.ylabel('Y')
plt.legend()

plt.show()

print(f'Linear Regression: Y = {slope:.2f}X + {intercept:.2f}')
print(f'Coefficient of Correlation (r): {correlation_coefficient:.4f}')

if correlation_coefficient >= 0.95:
    print('The linear model is a good fit for the dataset.')
else:
    print('The linear model may not be the best fit for the dataset. Consider exploring oth
```

Table of contents

- Getting started
- Data science
- Machine learning
- More Resources
- Featured examples
- Section

Chrome File Edit View History Bookmarks Profiles Tab Window Help

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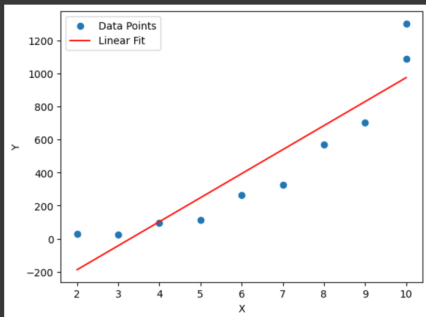
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```
print('The linear model may not be the best fit for the dataset. Consider exploring oth
```



Linear Regression: Y = 145.17X + -478.12
Coefficient of Correlation (r): 0.9212
The linear model may not be the best fit for the dataset. Consider exploring other models.

Table of contents

- Getting started
- Data science
- Machine learning
- More Resources
- Featured examples
- Section

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