Homework1

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1 Data Mining Homework 1

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1.1 Part I

1.1

The following program is used to sort the list according to the second element in the sublist. Please complete the missing lines.

1.2

The following program is used to clean up the string and remove all the punctuation marks. Please complete the missing lines.

```
In [7]: # define punctuation
    punctuations = '''!()-[]{};:'"\,<>./?@#$%^&*_~'''

my_str = "Hello!!!, he said ---and went."

# To take input from the user
    # my_str = input("Enter a string: ")

# remove punctuation from the string
no_punct = ""
for char in my_str:
    if(not char in punctuations):
```

```
# display the unpunctuated string
        print(no_punct)
Hello he said and went
  1.3
  Complete the following program to multiply two matrices using nested loops.
In [8]: sum([1,2])
Out[8]: 3
In [17]: # Program to multiply two matrices using nested loops
         # 3x3 matrix
         X = [[12,7,3],
             [4,5,6],
             [7,8,9]]
         # 3x4 matrix
         Y = [[5,8,1,2],
             [6,7,3,0],
             [4,5,9,1]]
         # result is 3x4
         result = [[0,0,0,0],
                  [0,0,0,0],
                  [0,0,0,0]]
         \# iterate through rows of X
         for i in range(len(X)):
            # iterate through columns of Y
            for j in range(len(Y[0])):
                 # iterate through rows of Y
                 #for all values in the x row, multiply by the correspoding value in the Y col
                 runningTotal = 0
                 for index in range(len(X)):
                     runningTotal = runningTotal + X[i][index]*Y[index][j]
                 result[i][j] = runningTotal
         for r in result:
             print(r)
[114, 160, 60, 27]
[74, 97, 73, 14]
[119, 157, 112, 23]
```

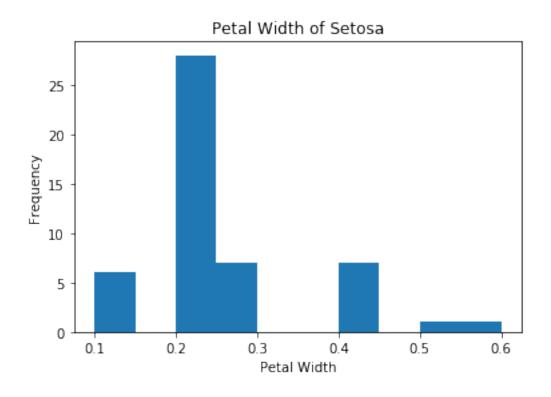
no_punct = no_punct + char

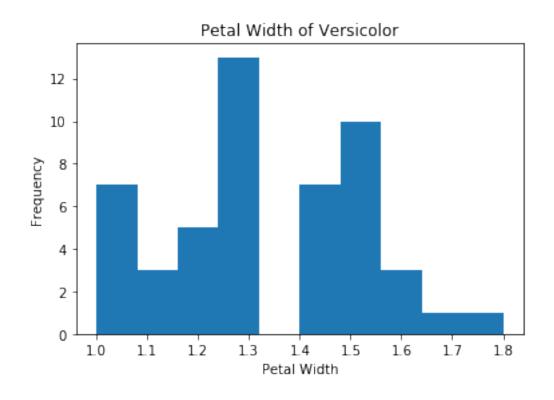
1.2 Part II

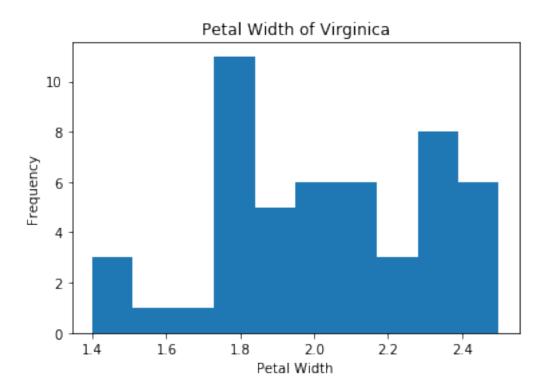
2.1

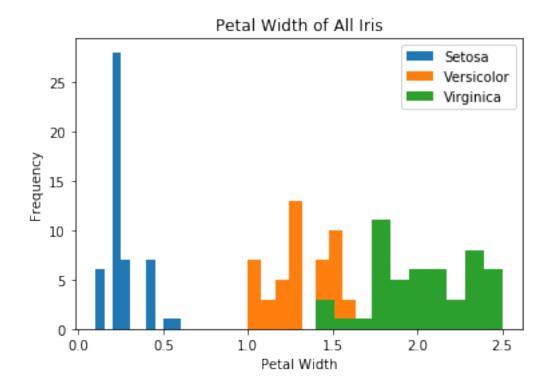
For each of the three flower categories, please quantize the petal width into 10 bins, and account the frequency of each bin, i.e., generating a 1-D histogram. Compare the three histograms side by side and explain your observations. You may use the function hist() or write your codes to implement the above objective.

```
In [22]: import pandas
         iris = pandas.read csv('iris.csv')
In [23]: iris.head()
Out [23]:
            sepal_length sepal_width petal_length petal_width species
         0
                     5.1
                                  3.5
                                                1.4
                                                             0.2 setosa
                     4.9
                                  3.0
         1
                                                1.4
                                                             0.2 setosa
         2
                     4.7
                                  3.2
                                                1.3
                                                             0.2 setosa
         3
                     4.6
                                  3.1
                                                1.5
                                                             0.2 setosa
                     5.0
                                  3.6
                                                1.4
                                                             0.2 setosa
In [34]: iris['species'].unique()
Out[34]: array(['setosa', 'versicolor', 'virginica'], dtype=object)
In [46]: import matplotlib.pyplot as plt
         plt.hist(iris.where(iris['species'] == 'setosa')['petal_width'],bins=10)
         plt.title("Petal Width of Setosa")
         plt.xlabel("Petal Width")
         plt.ylabel("Frequency")
         plt.show()
         plt.hist(iris.where(iris['species'] == 'versicolor')['petal_width'],bins=10)
         plt.title("Petal Width of Versicolor")
         plt.xlabel("Petal Width")
         plt.ylabel("Frequency")
         plt.show()
         plt.hist(iris.where(iris['species'] == 'virginica')['petal_width'],bins=10)
         plt.title("Petal Width of Virginica")
         plt.xlabel("Petal Width")
         plt.ylabel("Frequency")
         plt.show()
```









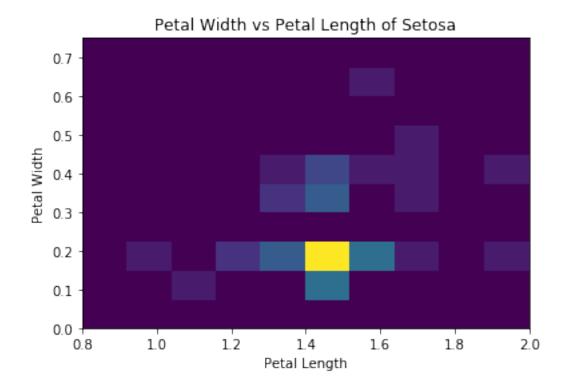
The setosa iris is the smallest of the three. Versicolor has a medium petal width, with some overlap with virginica. Virginica has the widest petal of the iris's being compared.

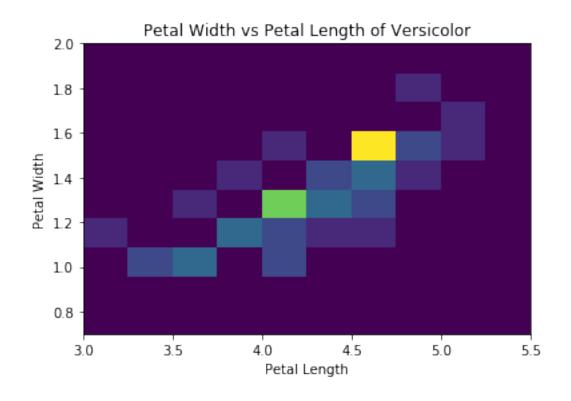
2.2

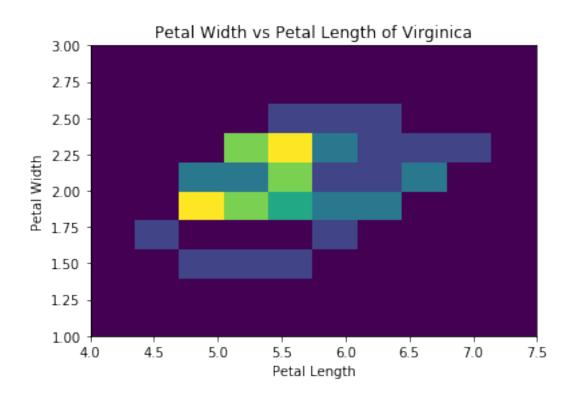
For each of the three categories, generate a 2D histogram

```
In [86]: plt.hist2d(x=iris.where(iris['species'] == 'setosa')['petal_length'].tolist(),
                    y=iris.where(iris['species'] == 'setosa')['petal_width'].tolist(),
                    range=[[.8,2],[0,.75]])
         plt.title("Petal Width vs Petal Length of Setosa")
         plt.xlabel("Petal Length")
         plt.ylabel("Petal Width")
         plt.show()
         plt.hist2d(x=iris.where(iris['species'] == 'versicolor')['petal_length'].tolist(),
                    y=iris.where(iris['species'] == 'versicolor')['petal_width'].tolist(),
                    range=[[3,5.5],[.7,2]])
         plt.title("Petal Width vs Petal Length of Versicolor")
         plt.xlabel("Petal Length")
         plt.ylabel("Petal Width")
         plt.show()
         plt.hist2d(x=iris.where(iris['species'] == 'virginica')['petal_length'].tolist(),
                    y=iris.where(iris['species'] == 'virginica')['petal_width'].tolist(),
                    range=[[4,7.5],[1,3]])
         plt.title("Petal Width vs Petal Length of Virginica")
         plt.xlabel("Petal Length")
```

plt.ylabel("Petal Width")
plt.show()





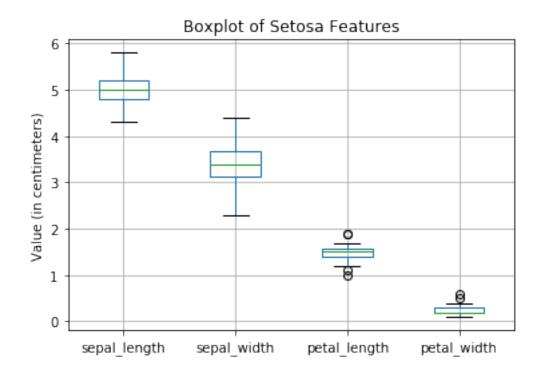


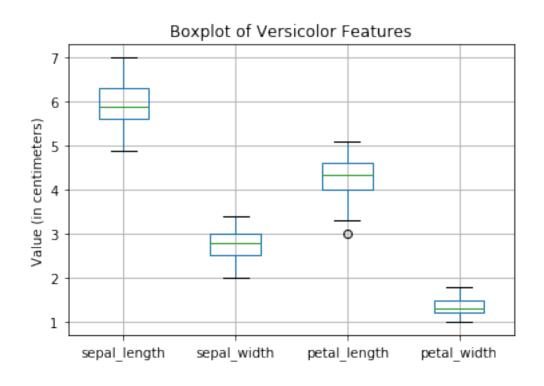
The range of size of the larger flowers tends to be wider. The axis labels are important for these, as we can see the Setosa's maximum petal length is shorter than the smallest Versicolor or Virginica.

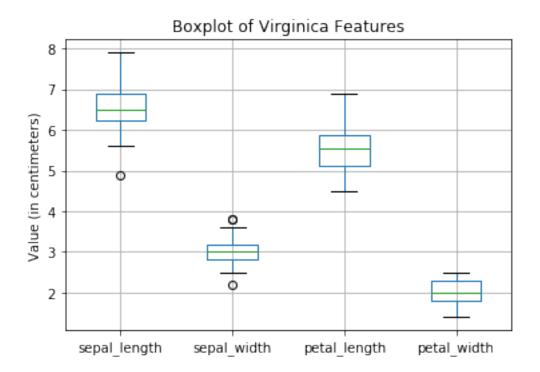
2.3

For each category, generate the BOX PLOTs for all four features

```
In [103]: iris.where(iris['species'] == 'setosa').boxplot(['sepal_length','sepal_width','petal_plt.title("Boxplot of Setosa Features")
    plt.ylabel("Value (in centimeters)")
    plt.show()
    iris.where(iris['species'] == 'versicolor').boxplot(['sepal_length','sepal_width','petal_plt.title("Boxplot of Versicolor Features")
    plt.ylabel("Value (in centimeters)")
    plt.show()
    iris.where(iris['species'] == 'virginica').boxplot(['sepal_length','sepal_width','petal_plt.title("Boxplot of Virginica Features")
    plt.ylabel("Value (in centimeters)")
    plt.show()
```







2.4 Visualization of similarity matrix.

plt.show()

plt.imshow(euclid, interpolation='none')
plt.title("Similarity Matrix of Iris's")

plt.ylabel("Virginica, Versicolor, and Setosa")

