Assignment_2_Iris_dataset

September 26, 2021

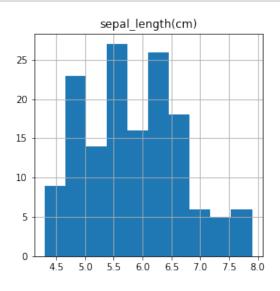
```
[2]: %pylab inline
    import warnings
    warnings.filterwarnings('ignore')
    import pandas as pd
    import numpy as np
   Populating the interactive namespace from numpy and matplotlib
[3]: column_names = ["sepal_length(cm)", "sepal_width(cm)", "petal_length(cm)",
     df = pd.read csv("http://archive.ics.uci.edu/ml/machine-learning-databases/iris/
     →iris.data", names=column_names, header=None)
    df.head()
[3]:
       sepal_length(cm)
                       sepal_width(cm) petal_length(cm) petal_width(cm) \
    0
                   5.1
                                  3.5
                                                   1.4
                                                                   0.2
                   4.9
                                  3.0
                                                   1.4
                                                                   0.2
    1
                   4.7
                                  3.2
                                                                   0.2
    2
                                                   1.3
                                  3.1
                                                   1.5
                                                                   0.2
    3
                   4.6
                   5.0
                                  3.6
                                                   1.4
                                                                   0.2
            class
    0 Iris-setosa
    1 Iris-setosa
    2 Iris-setosa
    3 Iris-setosa
    4 Iris-setosa
[4]: print("range ", df['sepal_length(cm)'].max() - df['sepal_length(cm)'].min())
    print("variance", df['sepal_length(cm)'].var())
    print(df['sepal_length(cm)'].describe())
    print("======="")
    print("range ", df['sepal_width(cm)'].max() - df['sepal_width(cm)'].min())
    print("variance", df['sepal_width(cm)'].var())
    print(df['sepal_width(cm)'].describe())
    print("======="")
                  ", df['petal_length(cm)'].max() - df['petal_length(cm)'].min())
```

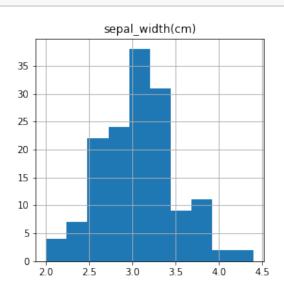
```
print("variance", df['petal_length(cm)'].var())
print(df['petal_length(cm)'].describe())
print("======"")
print("range
             ", df['petal_width(cm)'].max() - df['petal_width(cm)'].min())
print("variance", df['petal_width(cm)'].var())
print(df['petal_width(cm)'].describe())
        3.6000000000000005
range
variance 0.6856935123042507
       150.000000
count
mean
         5.843333
std
         0.828066
min
         4.300000
25%
         5.100000
50%
         5.800000
75%
         6.400000
         7.900000
max
Name: sepal_length(cm), dtype: float64
_____
       2.4000000000000004
range
variance 0.1880040268456376
count
       150.000000
         3.054000
mean
std
         0.433594
         2.000000
min
25%
         2.800000
50%
         3.000000
         3.300000
75%
         4.400000
max
Name: sepal_width(cm), dtype: float64
5.9
range
variance 3.113179418344519
count 150.000000
mean
         3.758667
         1.764420
std
min
         1.000000
25%
         1.600000
50%
         4.350000
75%
         5.100000
         6.900000
Name: petal_length(cm), dtype: float64
_____
        2.4
range
variance 0.582414317673378
       150.000000
count
         1.198667
mean
         0.763161
std
```

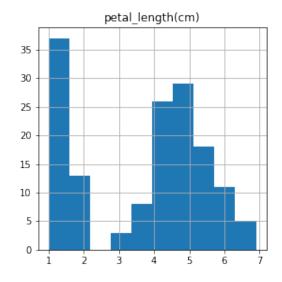
```
min 0.100000
25% 0.300000
50% 1.300000
75% 1.800000
max 2.500000
```

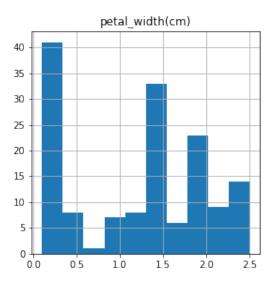
Name: petal_width(cm), dtype: float64

```
[9]: # Histogram
fig = plt.figure(figsize = (10,10))
ax = fig.gca()
histogram = df.hist(ax = ax)
```

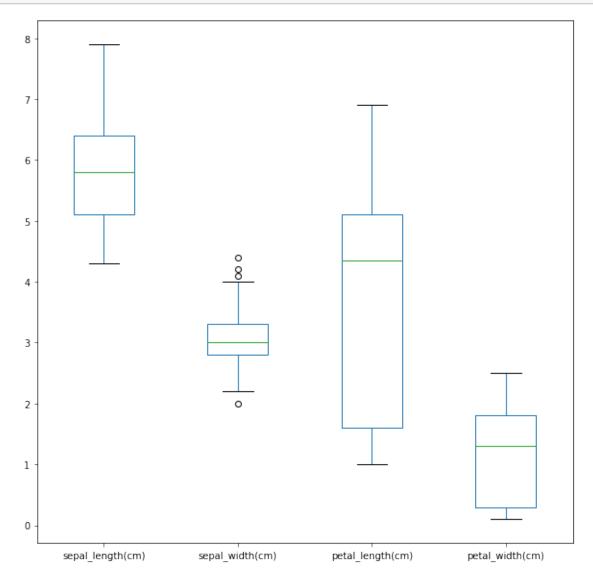








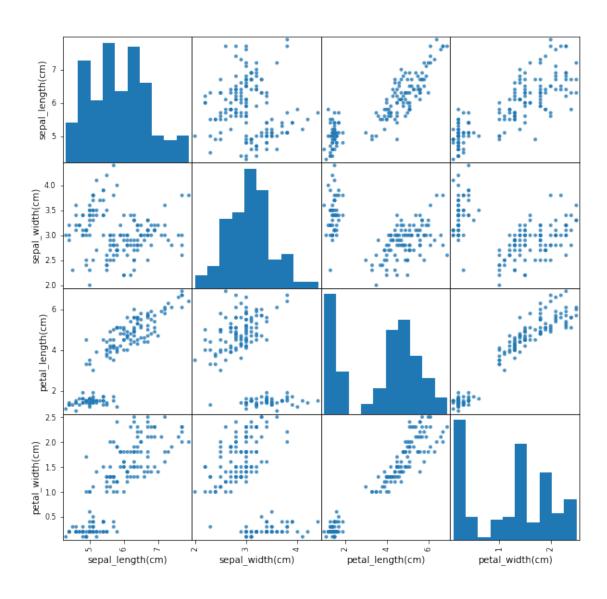
```
[39]: # Box plots
fig = plt.figure(figsize = (10,10))
ax = fig.gca()
box_plot = df.boxplot(ax = ax, grid=False, return_type='axes')
```



```
[7]: # Pairwise Plots

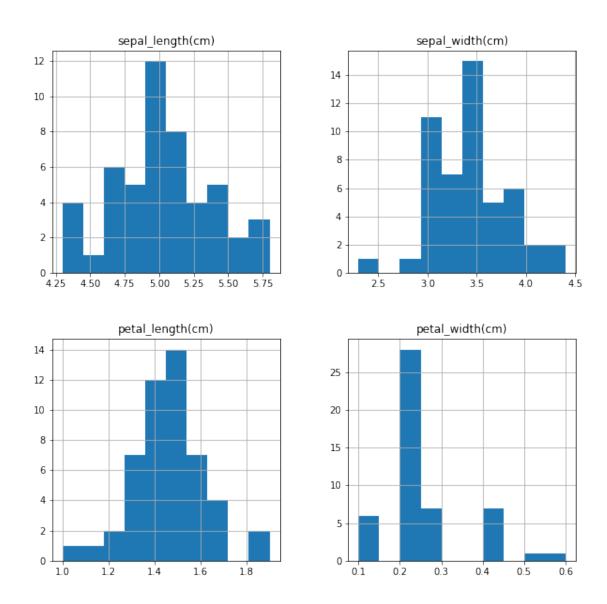
pairwise_plots = pd.plotting.scatter_matrix(df, figsize=(10,10), marker = '.', □

→s = 60, alpha = 0.8)
```



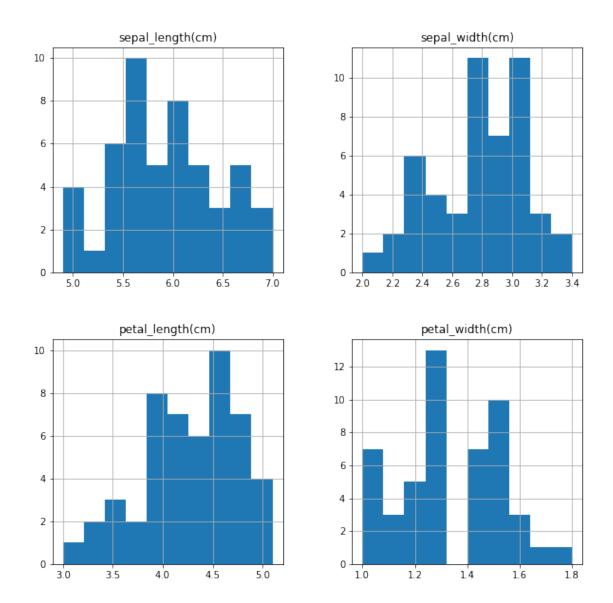
```
[12]: x = df[df['class'] == 'Iris-setosa']
print("SETOSA CLASS WISE VISUALIZATION")
setosa = x.hist(figsize=(10,10))
```

SETOSA CLASS WISE VISUALIZATION



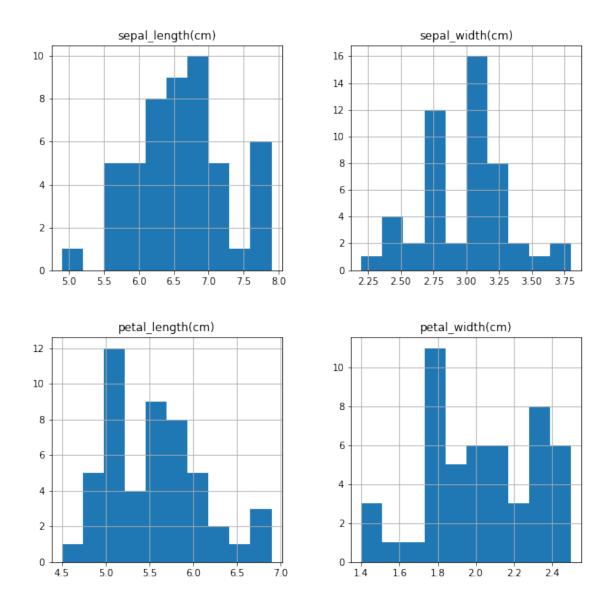
```
[68]: x = df[df['class'] == 'Iris-versicolor']
print("VERSICOLOR CLASS WISE VISUALIZATION")
setosa = x.hist(figsize=(10,10))
```

VERSICOLOR CLASS WISE VISUALIZATION



```
[69]: x = df[df['class'] == 'Iris-virginica']
print("VIRGINICA CLASS WISE VISUALIZATION")
setosa = x.hist(figsize=(10,10))
```

VIRGINICA CLASS WISE VISUALIZATION



[15]: #df.groupby(['class']).hist(figsize=(30,10))

1 1.3 - CONCEPTUAL QUESTIONS

1.0.1 1.

There are four features (Sepal Length (cm), Sepal Width (cm), Petal Length (cm), Petal Width (cm)). They are all numeric.

1.0.2 2.

From the initial glance we can see that there are considerable amount of flowers in every bin of sepal length and width. This means the data from this perspective is less grained. But when you

look at petal length and width there are gaps which suggest that flowers with that petal length and width are rare to find.

In the histograms for whole data, the petal length histogram is divided into two parts. The first part ends at pental_length = 2.1 cms and the second part starts at petal_length ~ 2.8 cms. So, any value in this range can segment the histogram into two separate parts.

1.0.3 3.

The pair of (sepal_length, petal_width) has the biggest parity in their medians. And, petal_length is the feature that explains the greatest amount of data.

1.0.4 4.

From the pairwise plot, the three pairs that have a strong correlation are (petal_length, petal_width), (petal_length, sepal_length), and (petal_width, sepal_length).

1.0.5 5.

IRIS SETOSA For Iris-Setosa class of flowers, the petal_width is the most differentiating factor, with all of them falling in the low ranges of the values when compared to other classes. This might be the reason for the low median score for petal_width which is represented in the box plot of the whole data.

IRIS VERSICOLOR For Iris-Versicolor class of flowers, while sepal_length and petal_width occupy the spots around the median, sepal_width and petal_length are mostly > median of the whole dataset.

IRIS VIRGINICA For Iris-Virginica class of flowers, the sepal_length is mostly >= median of the whole dataset, while the rest of the features are mostly populated around the median.

[3]: import nbconvert