## **Practical Question 6-KMeans**

Q6. Use Simple Kmeans, DBScan, Hierachical clustering algorithms for clustering. Compare the performance of clusters by changing the parameters involved in the algorithms.

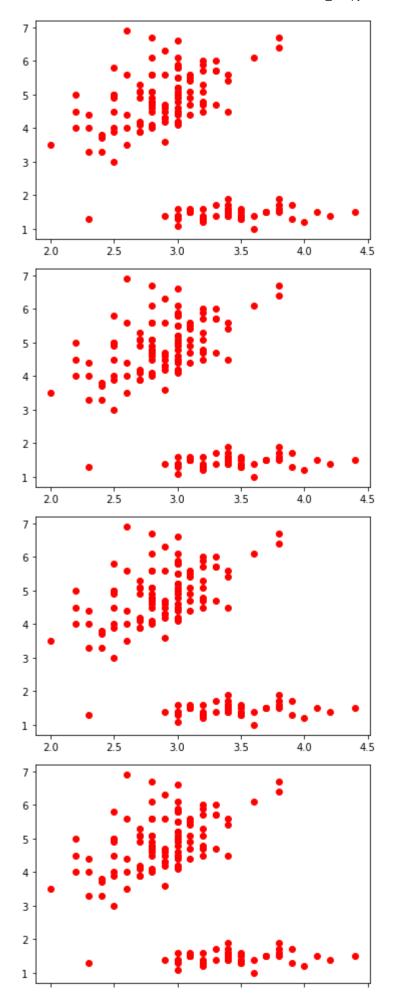
```
from google.colab import drive
drive.mount('/content/gdrive')
     Mounted at /content/gdrive
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
from sklearn.cluster import hierarchical
from sklearn.cluster import DBSCAN
from sklearn import preprocessing
from sklearn.decomposition import PCA
from sklearn.cluster import KMeans
iris=pd.read csv('/content/gdrive/MyDrive/q6 iris.csv')
print(iris)
#print(iris.keys())
          sepal length sepal width petal length
                                                    petal width
                                                                    species
     0
                   5.1
                                 3.5
                                               1.4
                                                             0.2
                                                                     setosa
     1
                   4.9
                                 3.0
                                               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
     4
                   5.0
                                                             0.2
                                 3.6
                                               1.4
                                                                     setosa
     145
                   6.7
                                 3.0
                                               5.2
                                                             2.3 virginica
     146
                   6.3
                                                             1.9 virginica
                                 2.5
                                               5.0
     147
                   6.5
                                 3.0
                                               5.2
                                                             2.0 virginica
                   6.2
                                                             2.3 virginica
     148
                                 3.4
                                               5.4
     149
                   5.9
                                 3.0
                                               5.1
                                                             1.8 virginica
     [150 rows x 5 columns]
sep l=iris.values[:,0]
print(sep_1)
#print(len(sep 1))
sep_w=iris.values[:,1]
print(sep_w)
#print(len(sep w))
pet_l=iris.values[:,2]
print(pet 1)
```

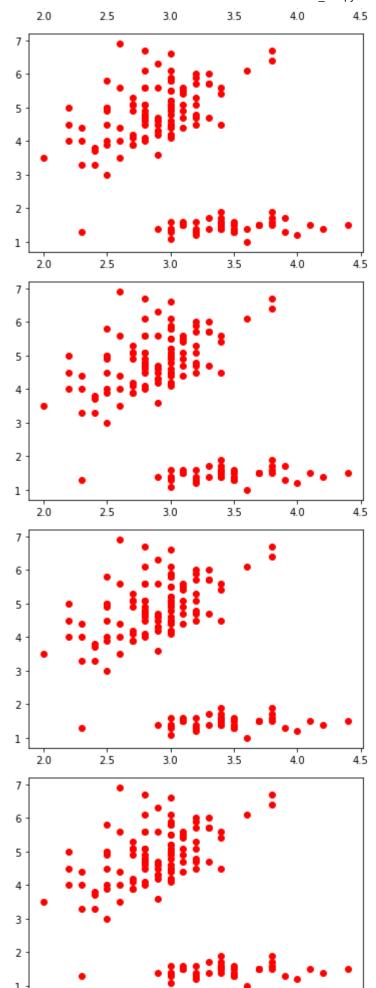
```
#print(len(pet_1))
pet_w=iris.values[:,3]
print(pet_w)
#print(len(pet_w))
```

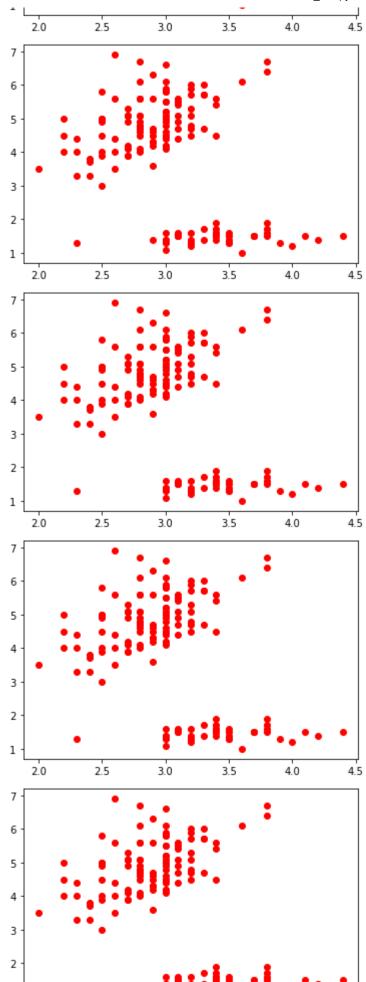
```
[5.1 4.9 4.7 4.6 5.0 5.4 4.6 5.0 4.4 4.9 5.4 4.8 4.8 4.3 5.8 5.7 5.4 5.1
 5.7 5.1 5.4 5.1 4.6 5.1 4.8 5.0 5.0 5.2 5.2 4.7 4.8 5.4 5.2 5.5 4.9 5.0
 5.5 4.9 4.4 5.1 5.0 4.5 4.4 5.0 5.1 4.8 5.1 4.6 5.3 5.0 7.0 6.4 6.9 5.5
 6.5 5.7 6.3 4.9 6.6 5.2 5.0 5.9 6.0 6.1 5.6 6.7 5.6 5.8 6.2 5.6 5.9 6.1
 6.3 6.1 6.4 6.6 6.8 6.7 6.0 5.7 5.5 5.5 5.8 6.0 5.4 6.0 6.7 6.3 5.6 5.5
 5.5 6.1 5.8 5.0 5.6 5.7 5.7 6.2 5.1 5.7 6.3 5.8 7.1 6.3 6.5 7.6 4.9 7.3
 6.7 7.2 6.5 6.4 6.8 5.7 5.8 6.4 6.5 7.7 7.7 6.0 6.9 5.6 7.7 6.3 6.7 7.2
 6.2 6.1 6.4 7.2 7.4 7.9 6.4 6.3 6.1 7.7 6.3 6.4 6.0 6.9 6.7 6.9 5.8 6.8
6.7 6.7 6.3 6.5 6.2 5.9]
[3.5 3.0 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 3.7 3.4 3.0 3.0 4.0 4.4 3.9 3.5
 3.8 3.8 3.4 3.7 3.6 3.3 3.4 3.0 3.4 3.5 3.4 3.2 3.1 3.4 4.1 4.2 3.1 3.2
 3.5 3.1 3.0 3.4 3.5 2.3 3.2 3.5 3.8 3.0 3.8 3.2 3.7 3.3 3.2 3.2 3.1 2.3
 2.8 2.8 3.3 2.4 2.9 2.7 2.0 3.0 2.2 2.9 2.9 3.1 3.0 2.7 2.2 2.5 3.2 2.8
 2.5 2.8 2.9 3.0 2.8 3.0 2.9 2.6 2.4 2.4 2.7 2.7 3.0 3.4 3.1 2.3 3.0 2.5
 2.6 3.0 2.6 2.3 2.7 3.0 2.9 2.9 2.5 2.8 3.3 2.7 3.0 2.9 3.0 3.0 2.5 2.9
 2.5 3.6 3.2 2.7 3.0 2.5 2.8 3.2 3.0 3.8 2.6 2.2 3.2 2.8 2.8 2.7 3.3 3.2
 2.8 3.0 2.8 3.0 2.8 3.8 2.8 2.8 2.6 3.0 3.4 3.1 3.0 3.1 3.1 3.1 2.7 3.2
 3.3 3.0 2.5 3.0 3.4 3.0]
[1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 1.6 1.4 1.1 1.2 1.5 1.3 1.4
 1.7 1.5 1.7 1.5 1.0 1.7 1.9 1.6 1.6 1.5 1.4 1.6 1.6 1.5 1.5 1.4 1.5 1.2
 1.3 1.5 1.3 1.5 1.3 1.3 1.3 1.6 1.9 1.4 1.6 1.4 1.5 1.4 4.7 4.5 4.9 4.0
 4.6 4.5 4.7 3.3 4.6 3.9 3.5 4.2 4.0 4.7 3.6 4.4 4.5 4.1 4.5 3.9 4.8 4.0
 4.9 4.7 4.3 4.4 4.8 5.0 4.5 3.5 3.8 3.7 3.9 5.1 4.5 4.5 4.7 4.4 4.1 4.0
 4.4 4.6 4.0 3.3 4.2 4.2 4.2 4.3 3.0 4.1 6.0 5.1 5.9 5.6 5.8 6.6 4.5 6.3
 5.8 6.1 5.1 5.3 5.5 5.0 5.1 5.3 5.5 6.7 6.9 5.0 5.7 4.9 6.7 4.9 5.7 6.0
 4.8 4.9 5.6 5.8 6.1 6.4 5.6 5.1 5.6 6.1 5.6 5.5 4.8 5.4 5.6 5.1 5.1 5.9
 5.7 5.2 5.0 5.2 5.4 5.1]
[0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 0.2 0.2 0.1 0.1 0.2 0.4 0.4 0.3
0.3 0.3 0.2 0.4 0.2 0.5 0.2 0.2 0.4 0.2 0.2 0.2 0.2 0.4 0.1 0.2 0.1 0.2
 0.2 0.1 0.2 0.2 0.3 0.3 0.2 0.6 0.4 0.3 0.2 0.2 0.2 0.2 1.4 1.5 1.5 1.3
 1.5 1.3 1.6 1.0 1.3 1.4 1.0 1.5 1.0 1.4 1.3 1.4 1.5 1.0 1.5 1.1 1.8 1.3
 1.5 1.2 1.3 1.4 1.4 1.7 1.5 1.0 1.1 1.0 1.2 1.6 1.5 1.6 1.5 1.3 1.3 1.3
 1.2 1.4 1.2 1.0 1.3 1.2 1.3 1.3 1.1 1.3 2.5 1.9 2.1 1.8 2.2 2.1 1.7 1.8
 1.8 2.5 2.0 1.9 2.1 2.0 2.4 2.3 1.8 2.2 2.3 1.5 2.3 2.0 2.0 1.8 2.1 1.8
 1.8 1.8 2.1 1.6 1.9 2.0 2.2 1.5 1.4 2.3 2.4 1.8 1.8 2.1 2.4 2.3 1.9 2.3
 2.5 2.3 1.9 2.0 2.3 1.8]
```

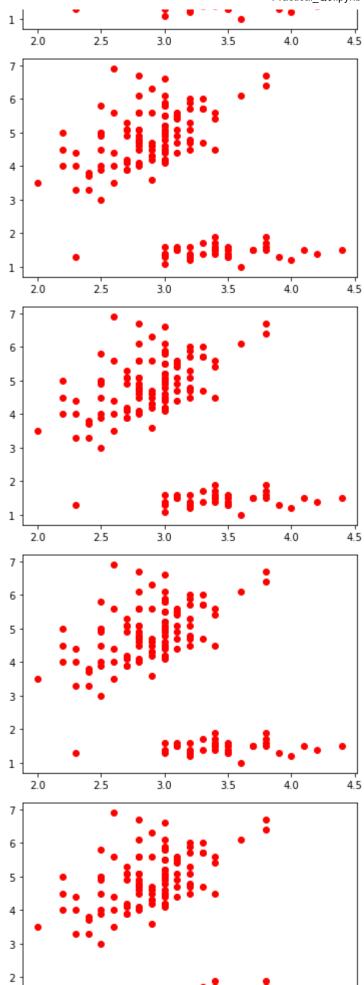
Taking Sepal Width and Petal Length as our two features for clustering

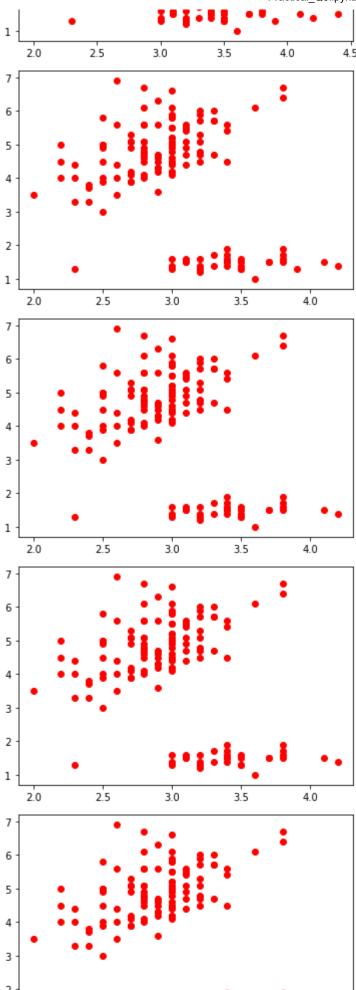
```
for i in range(150):
   if i<=49:
    plt.plot(iris.values[i:,1],iris.values[i:,2],'ro')
   if i>49 and i<=99:
     plt.plot(iris.values[i:,1],iris.values[i:,2],'bo')
   if i>99:
    plt.plot(iris.values[i:,1],iris.values[i:,2],'go')
   plt.show()
```

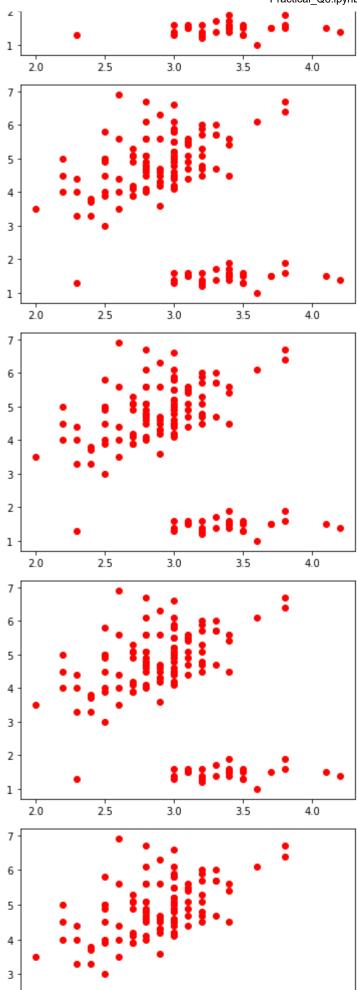


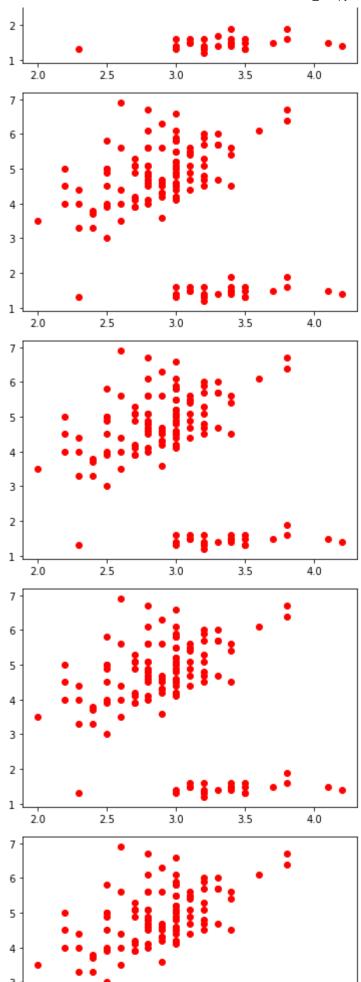


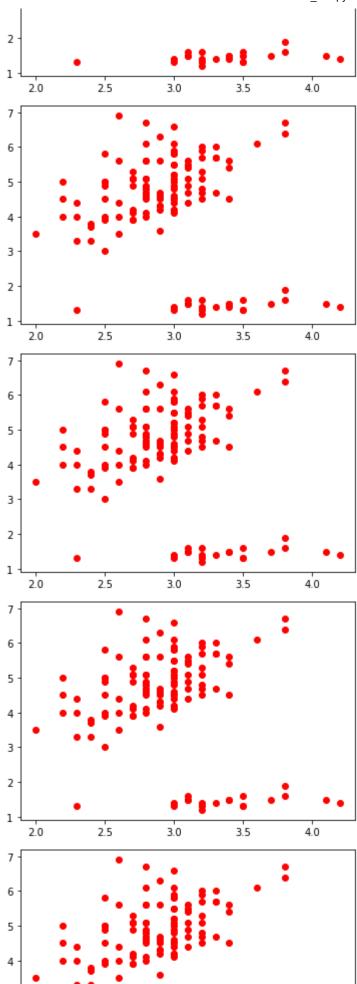


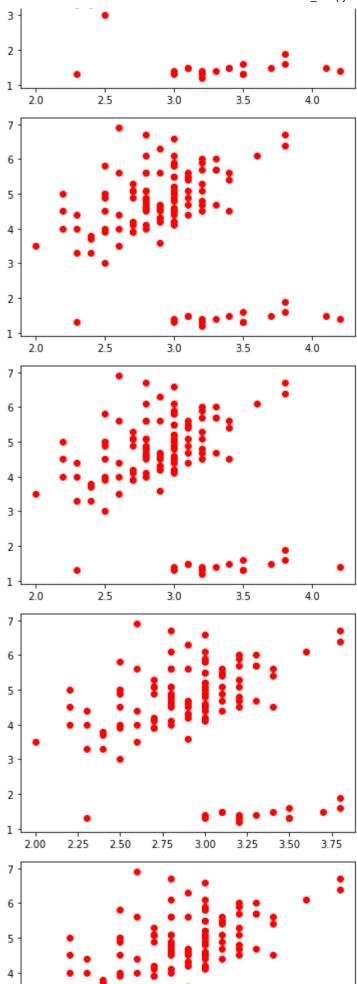


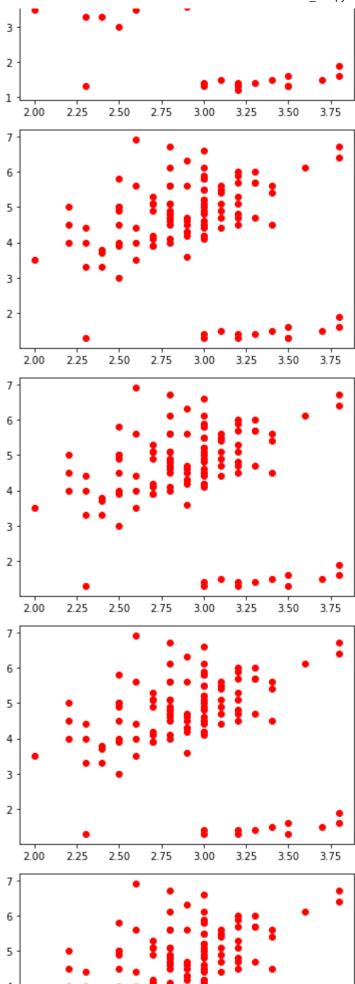


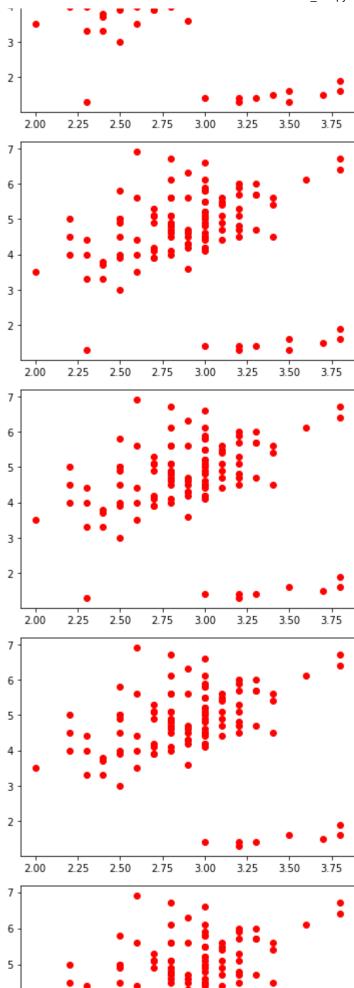


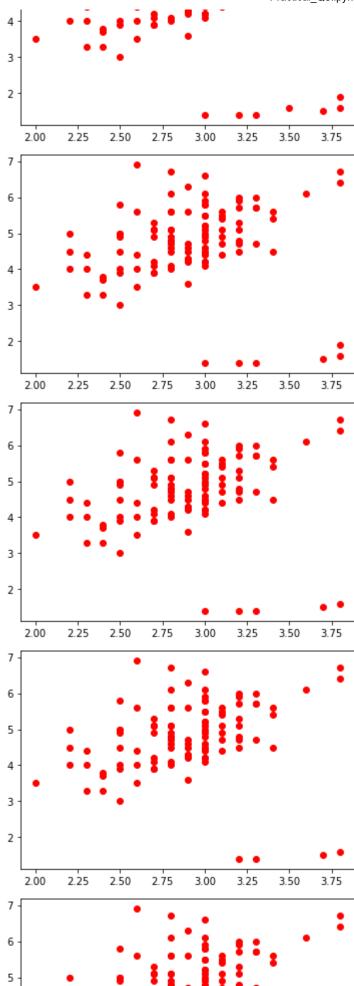


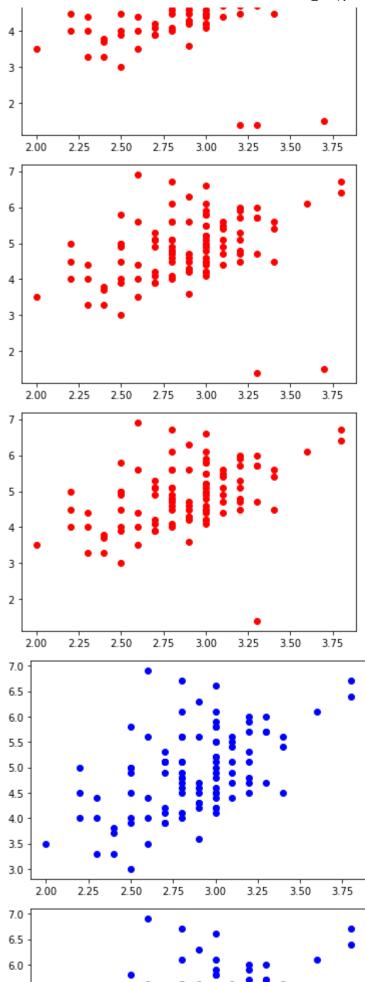


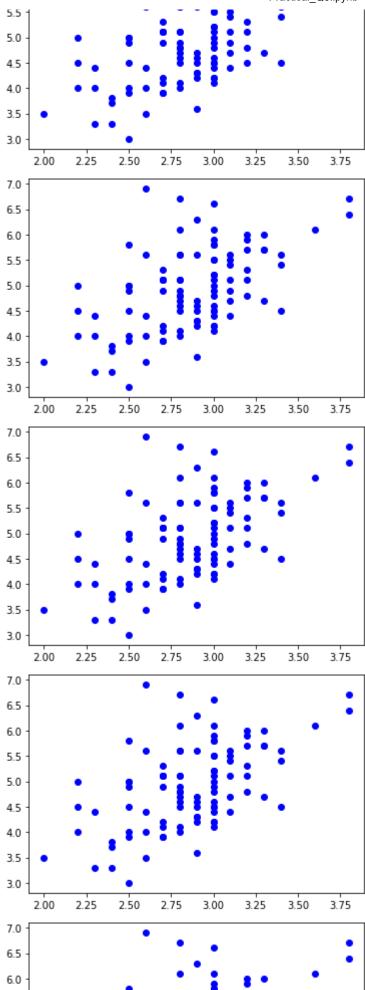


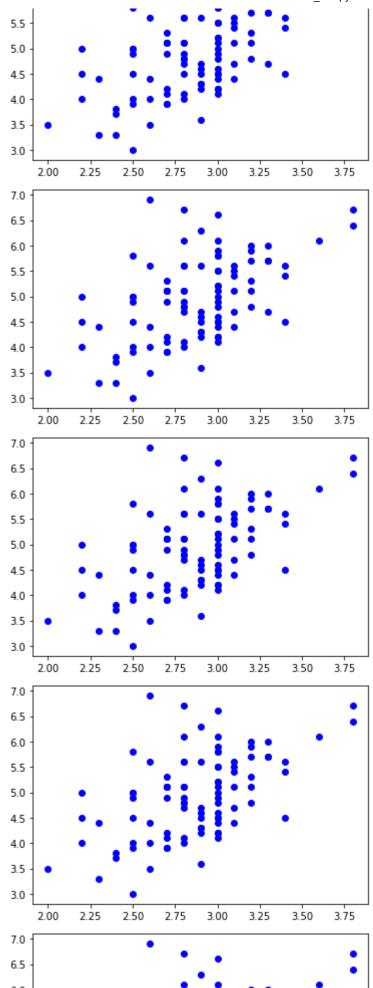


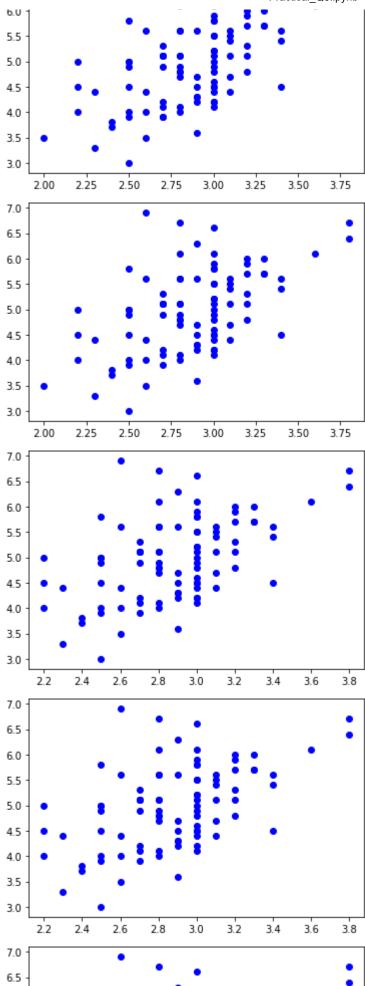


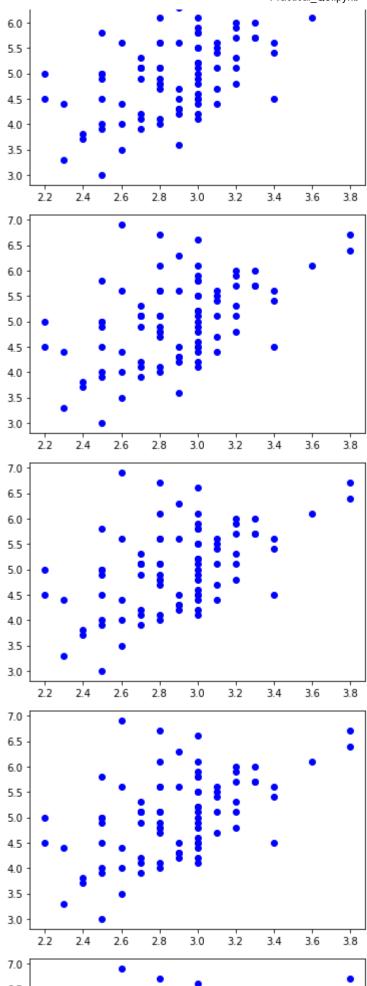


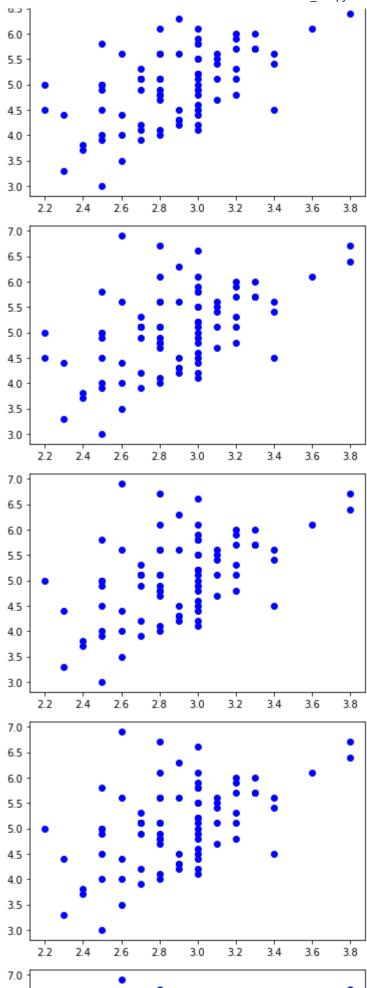


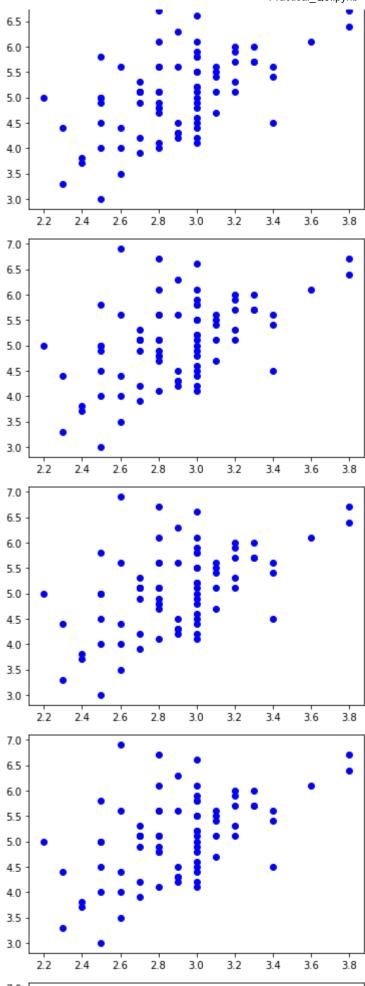


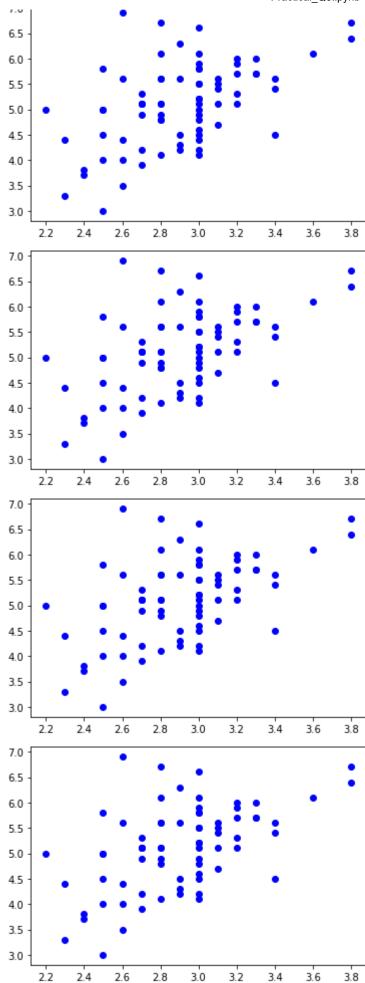


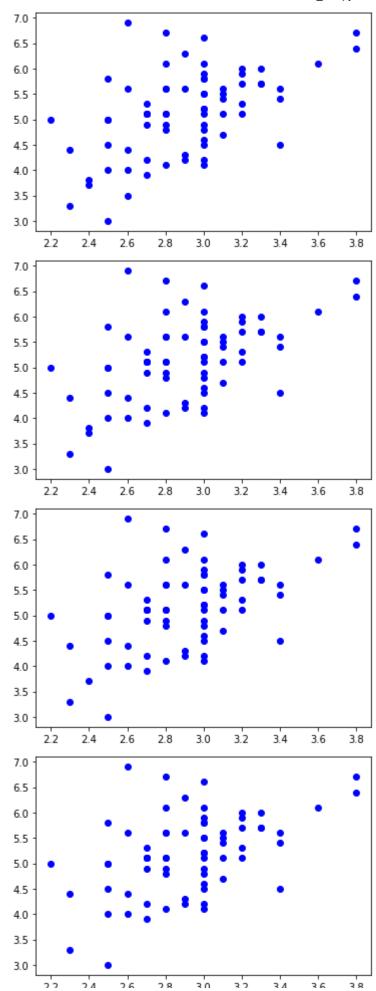


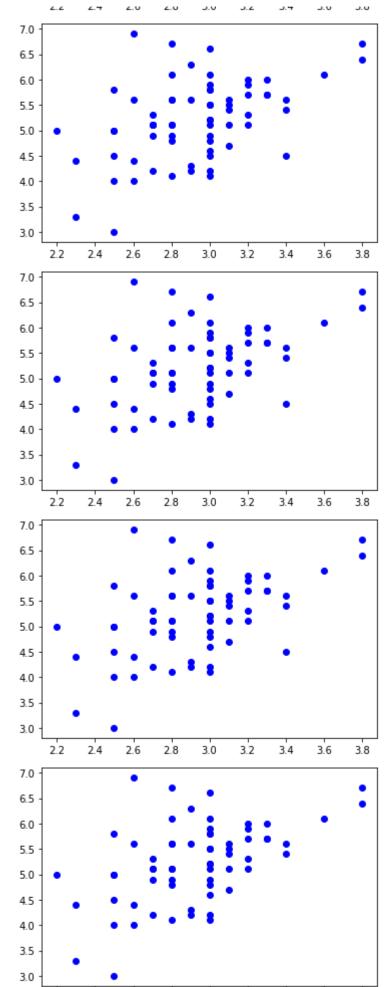


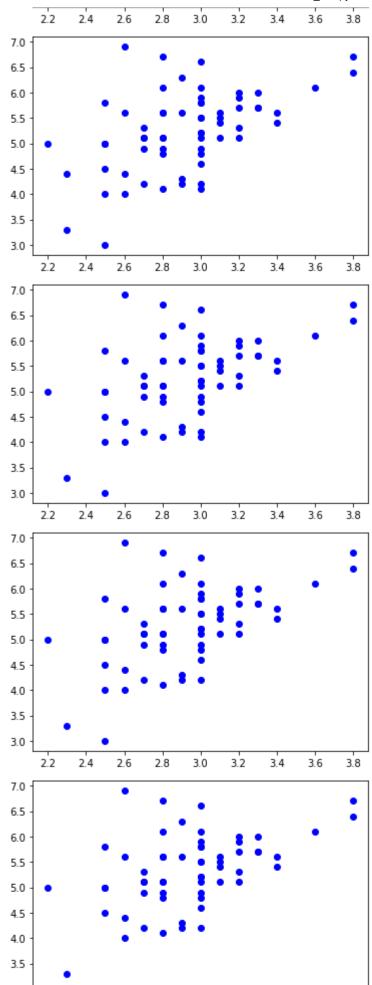


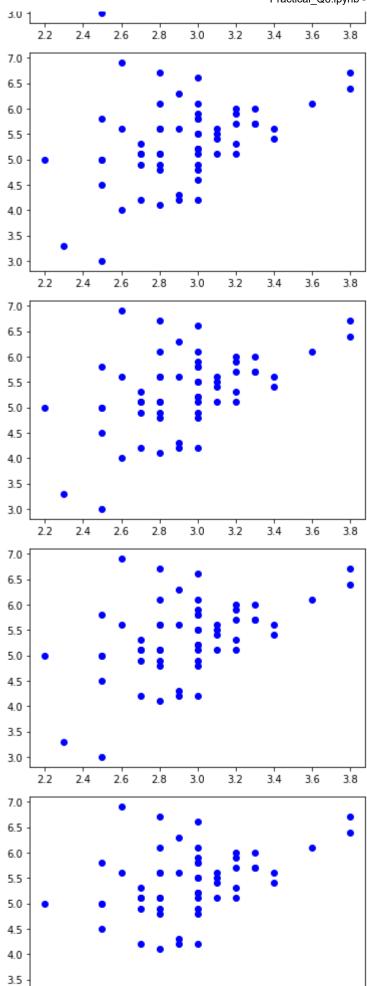


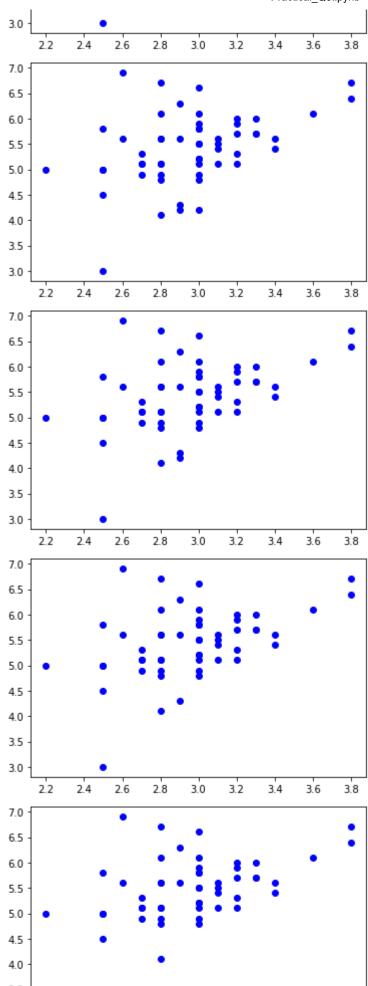


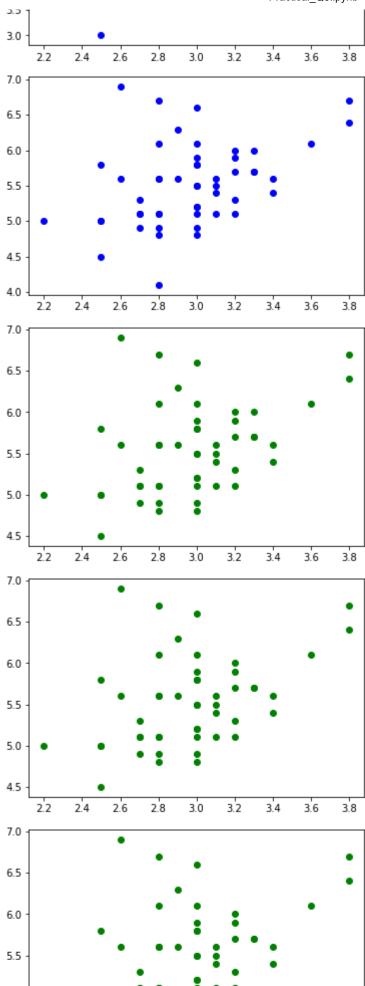


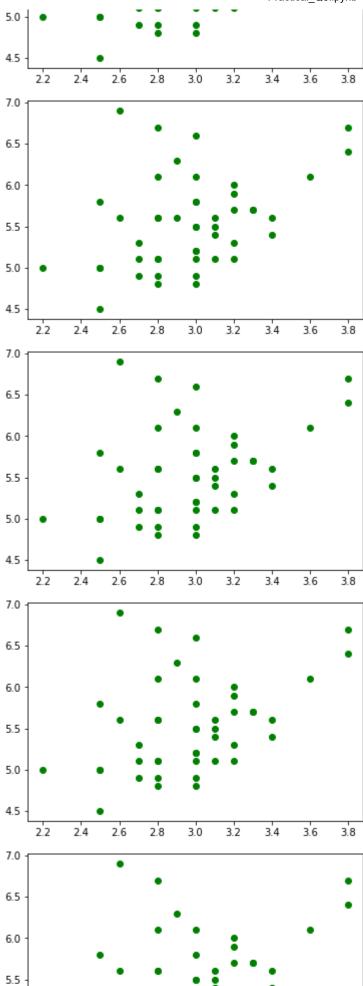


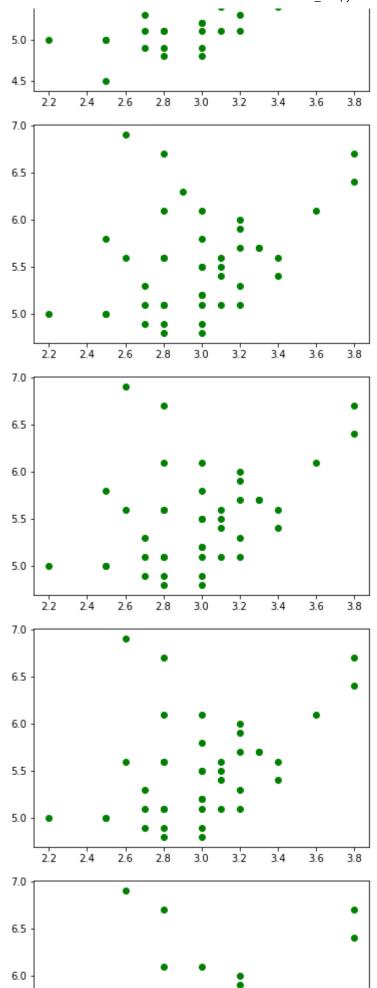


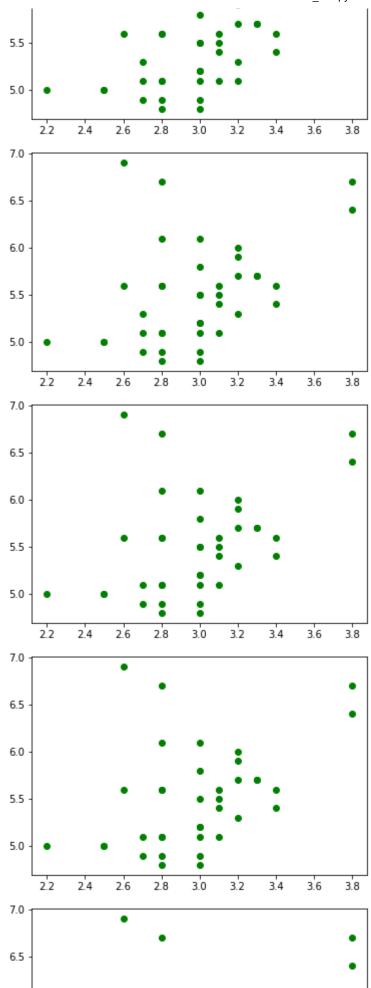


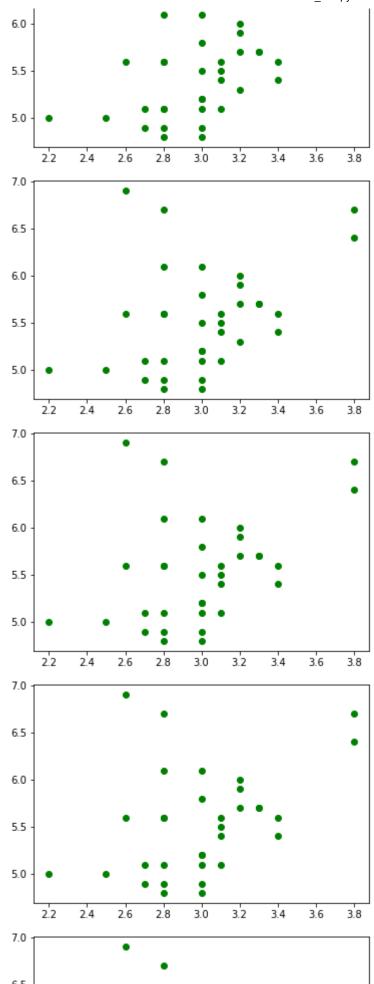


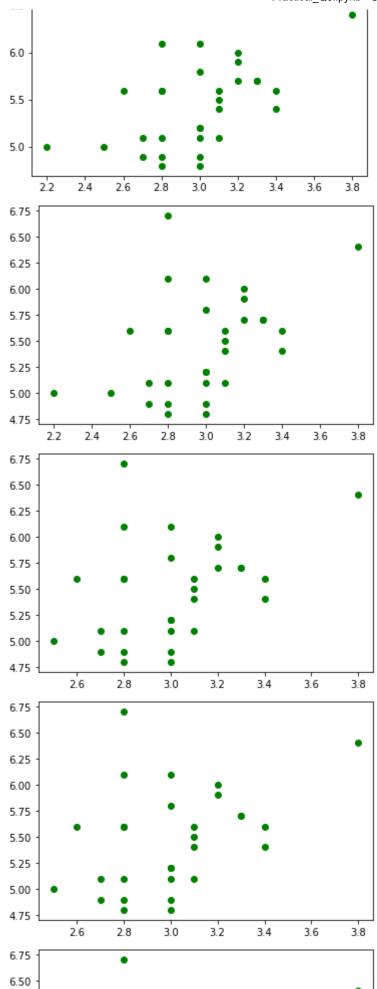


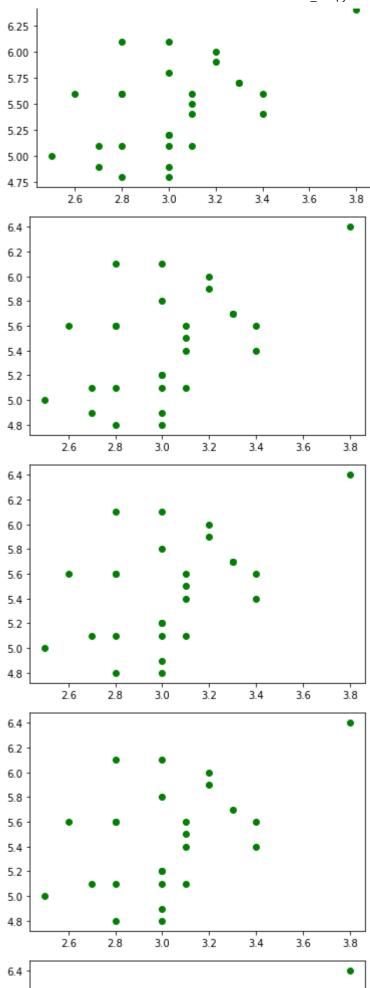


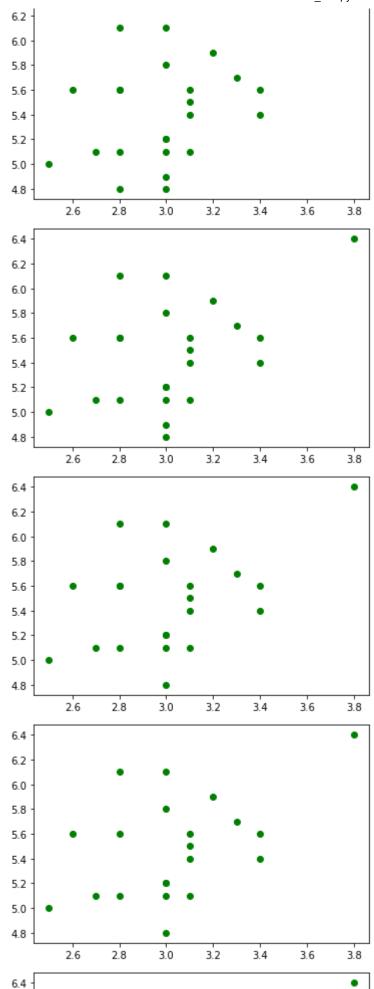


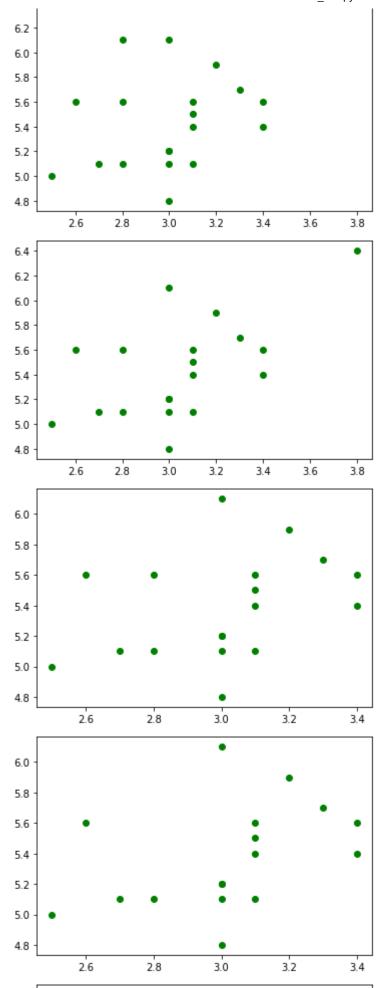


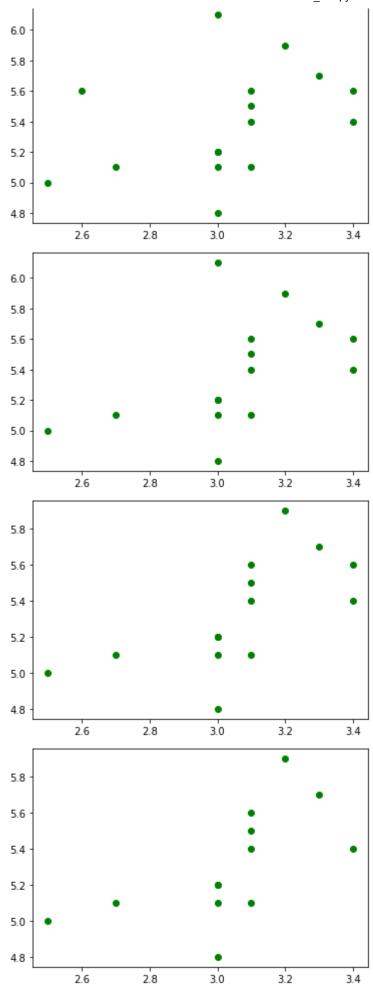


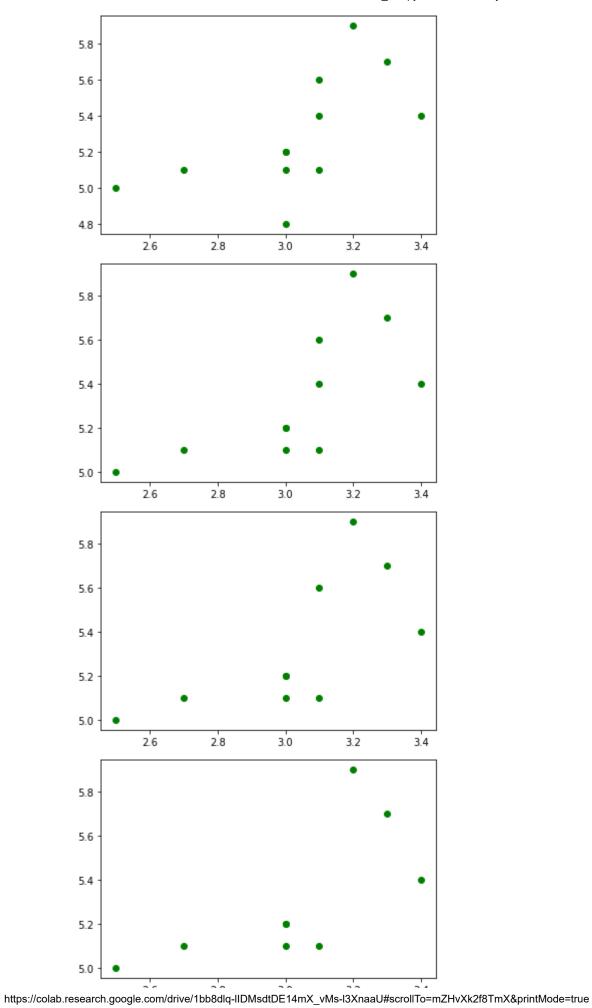


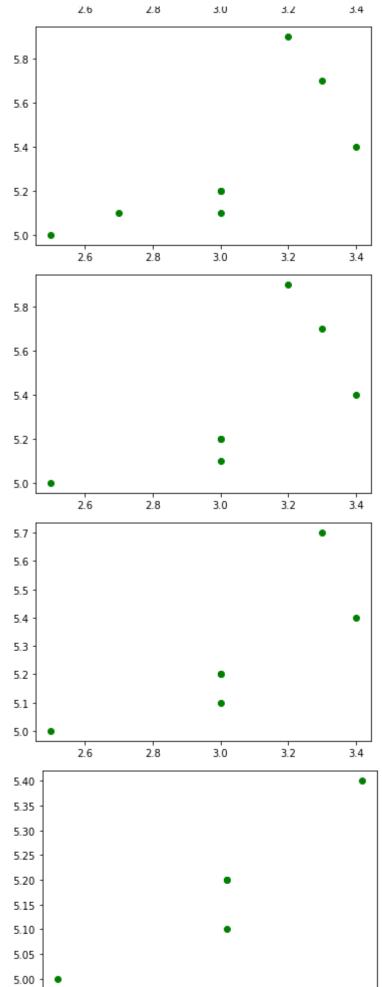


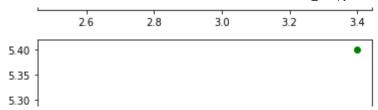






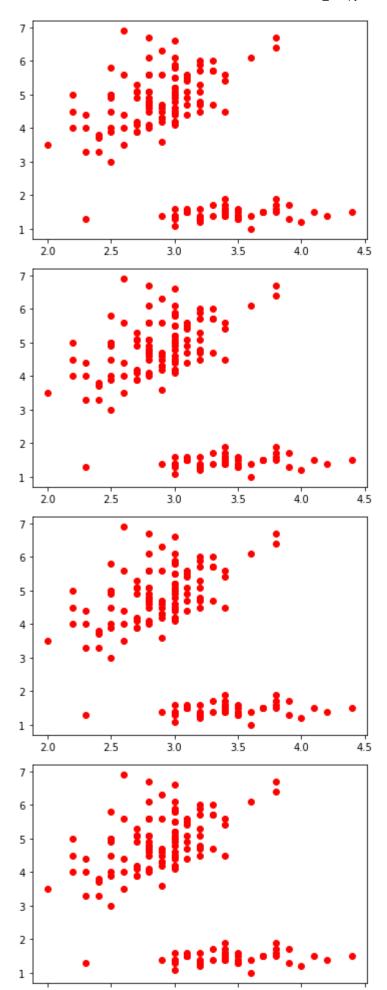


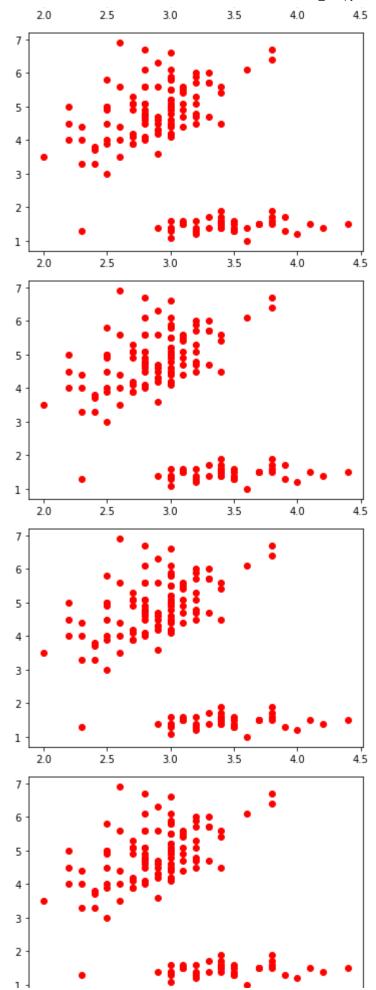


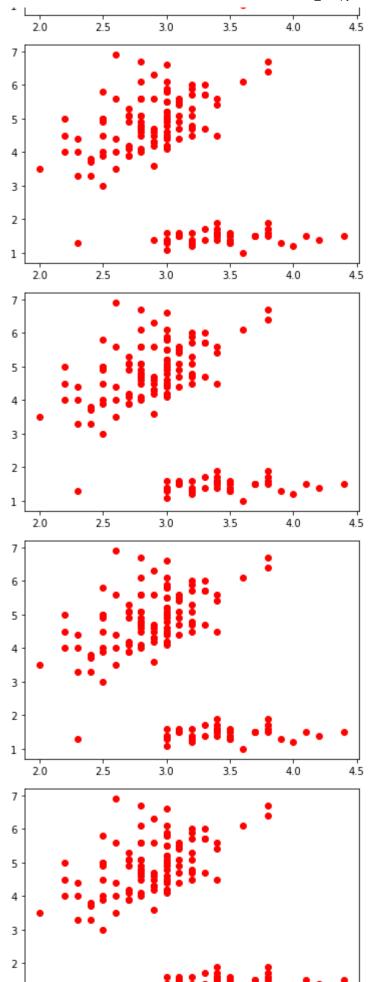


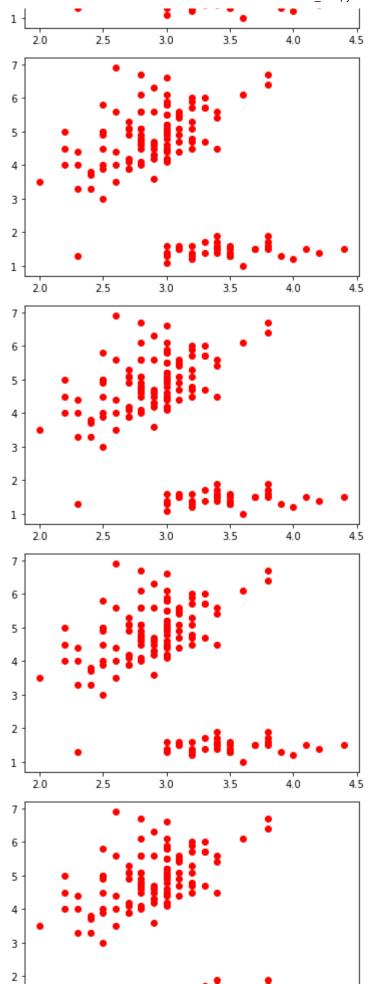
Taking Sepal Width and Petal Length as our two features for clustering

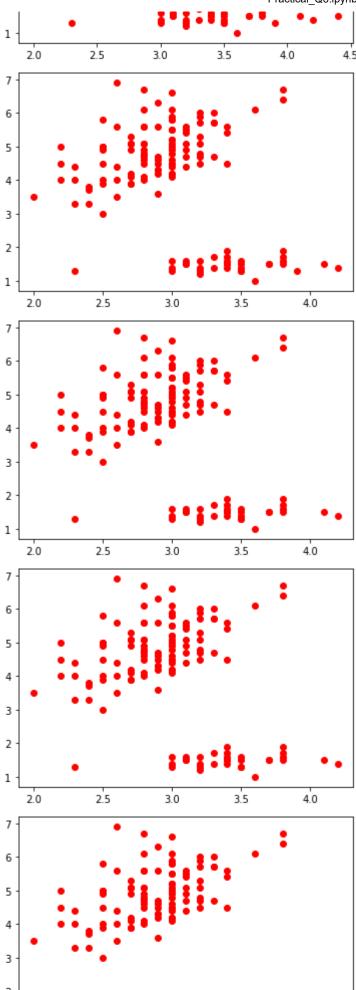
```
for i in range(150):
    if i<=49:
        plt.plot(iris.values[i:,1],iris.values[i:,2],'ro')
    if i>49 and i<=99:
        plt.plot(iris.values[i:,1],iris.values[i:,2],'bo')
    if i>99:
        plt.plot(iris.values[i:,1],iris.values[i:,2],'go')
    plt.show()
```

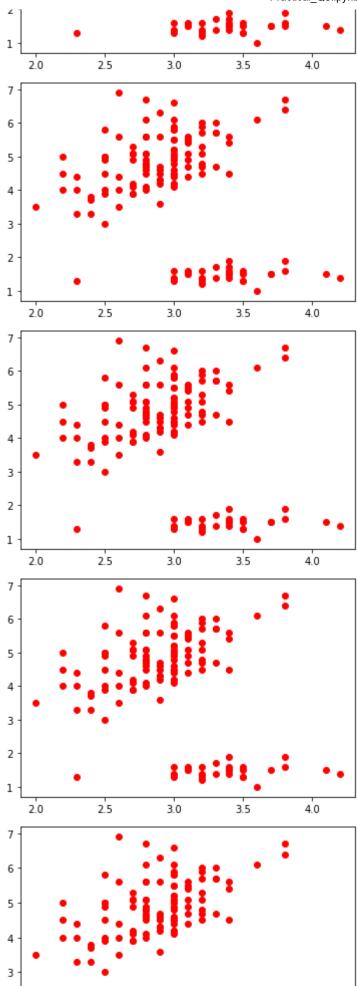


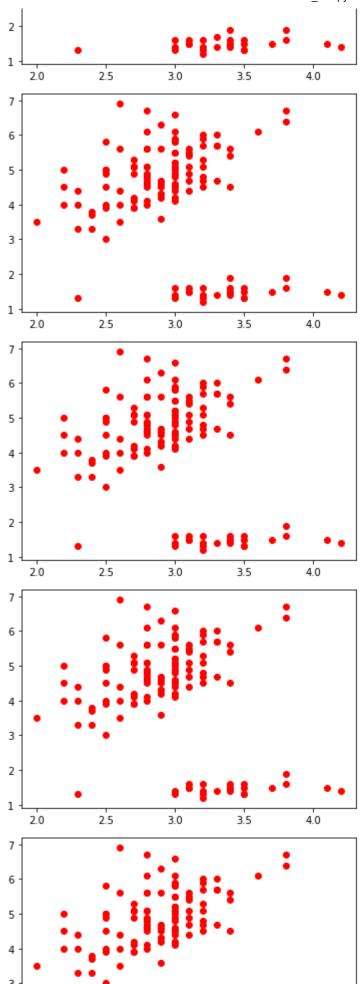


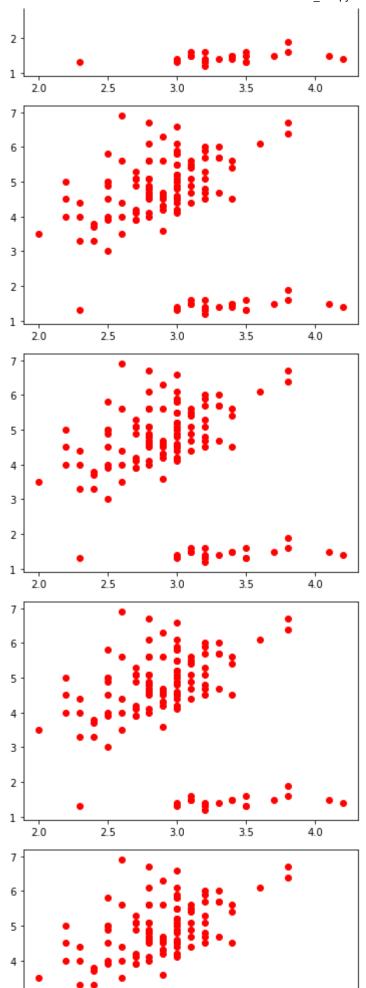


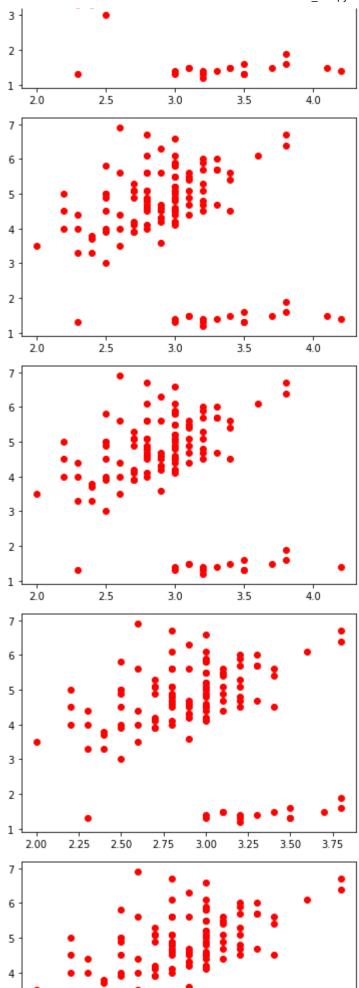


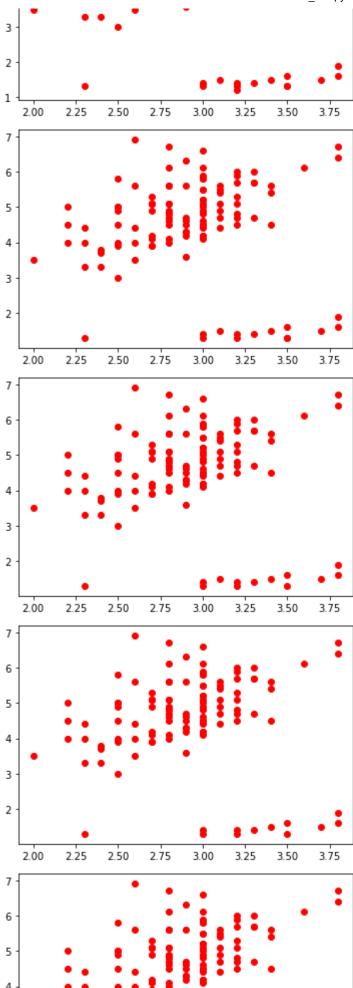


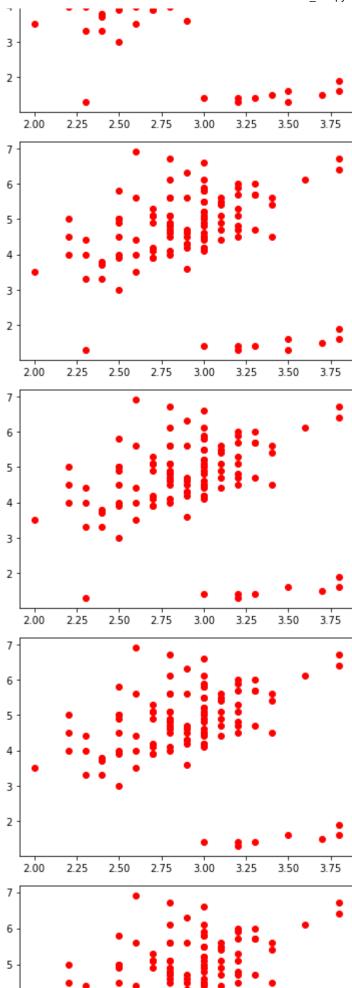


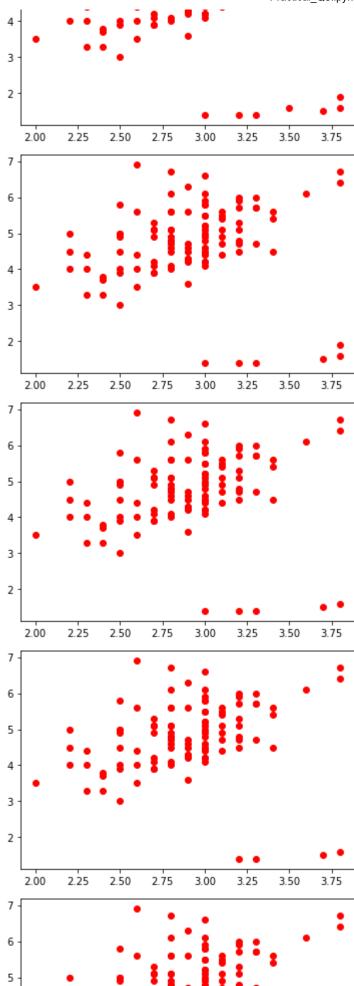


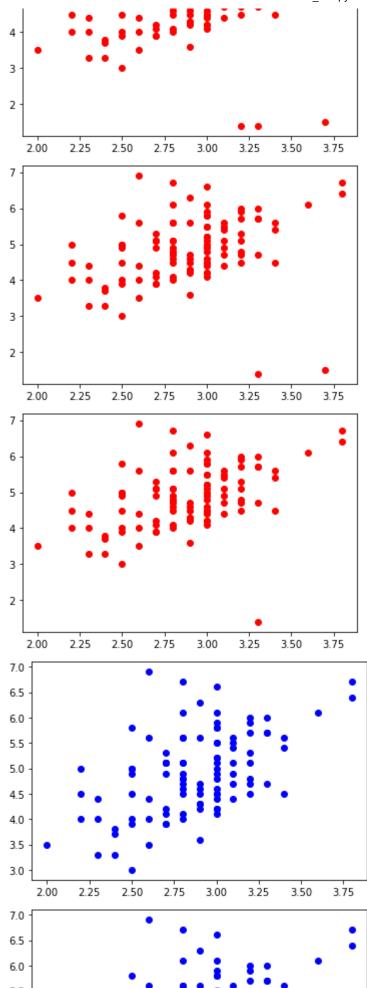


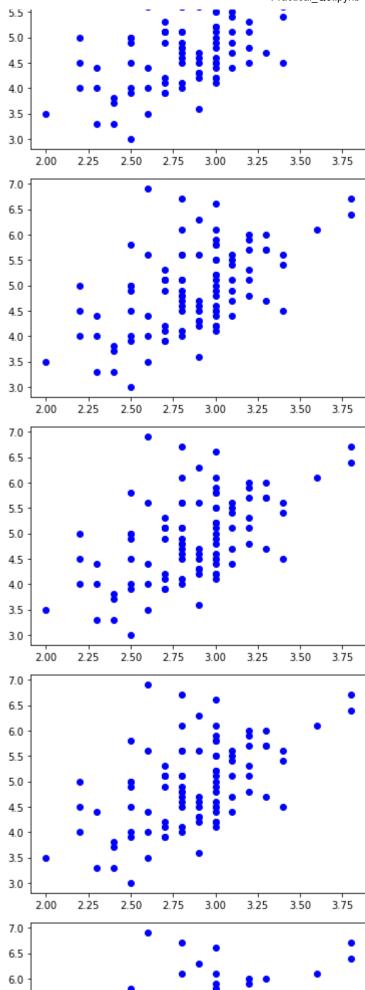


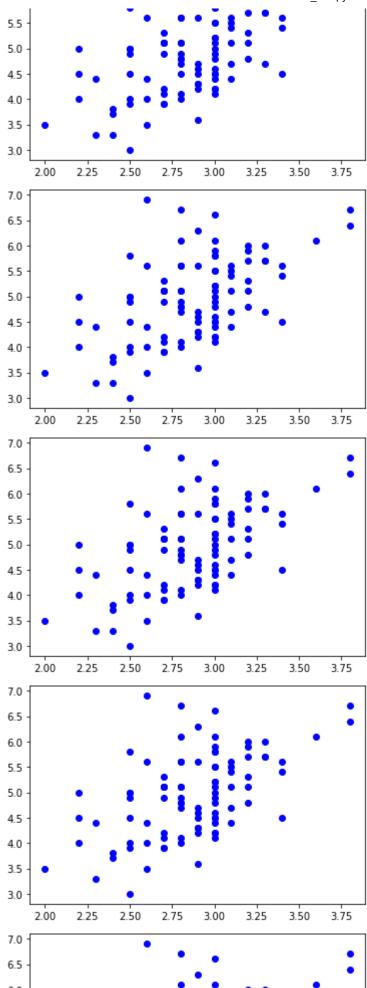


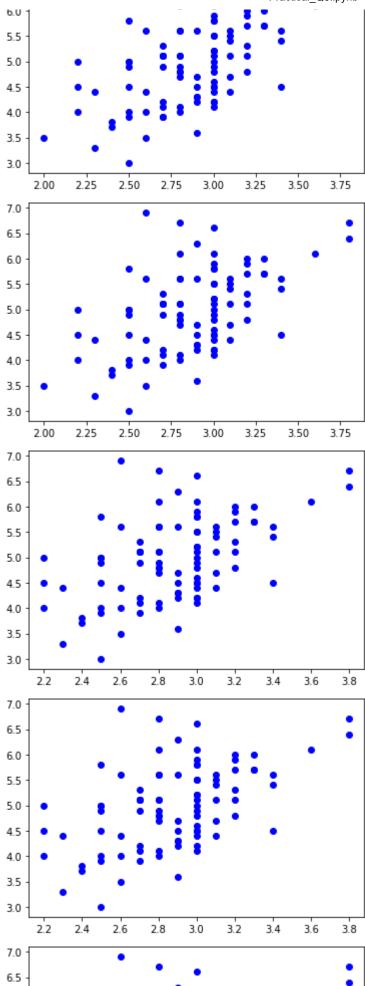


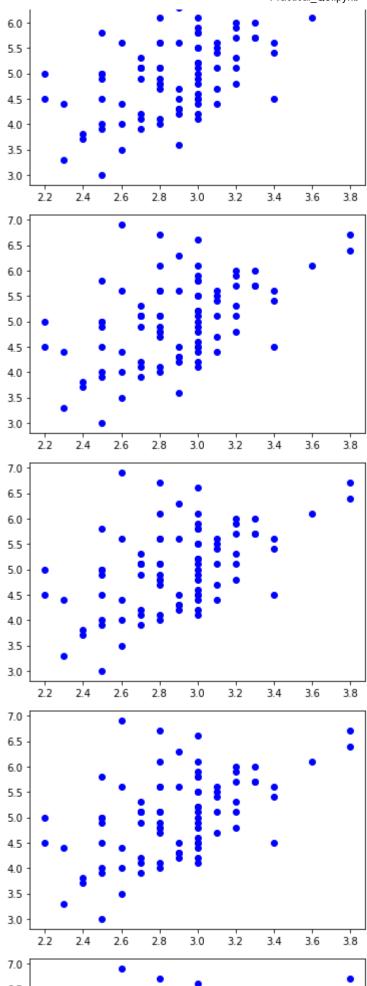


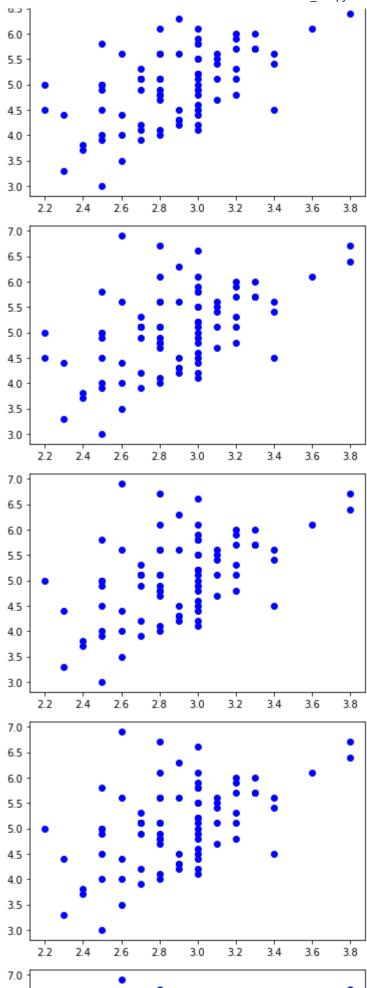


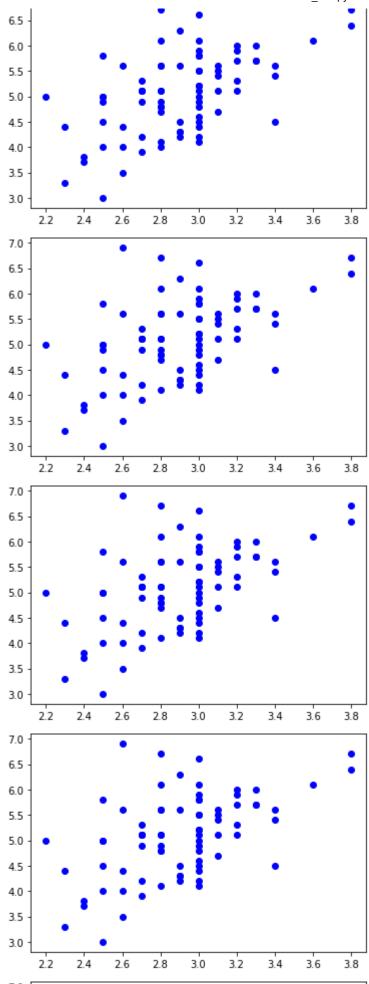


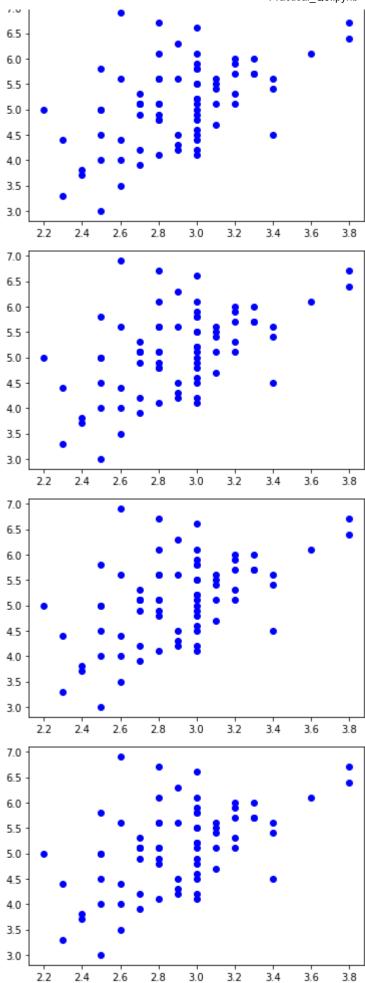


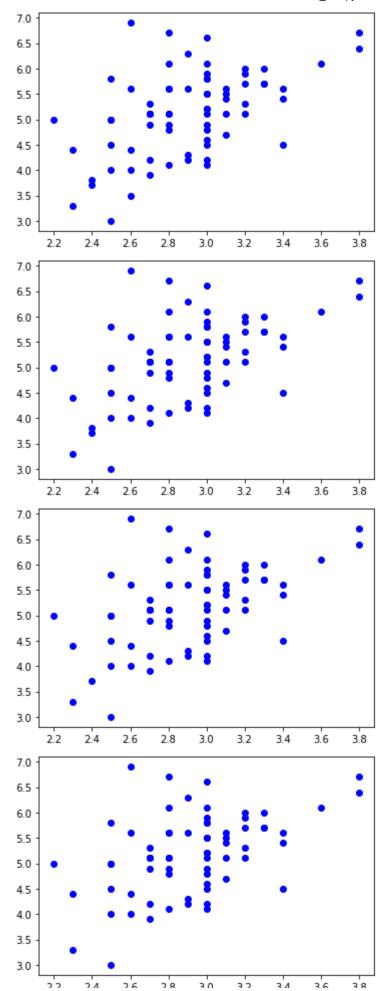


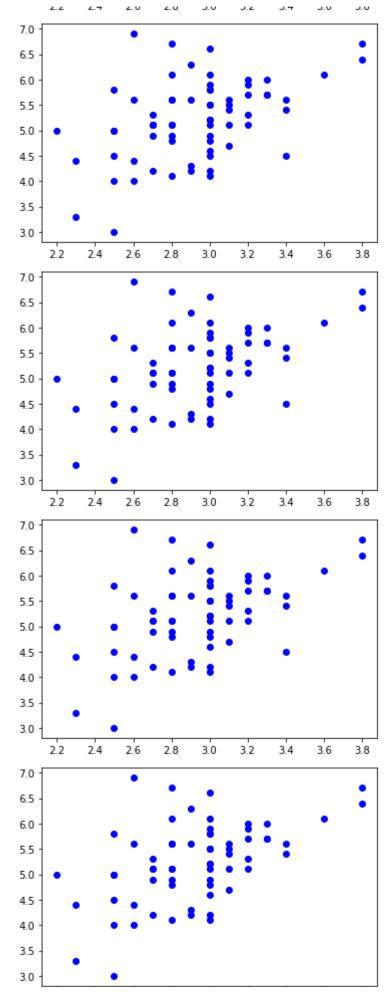


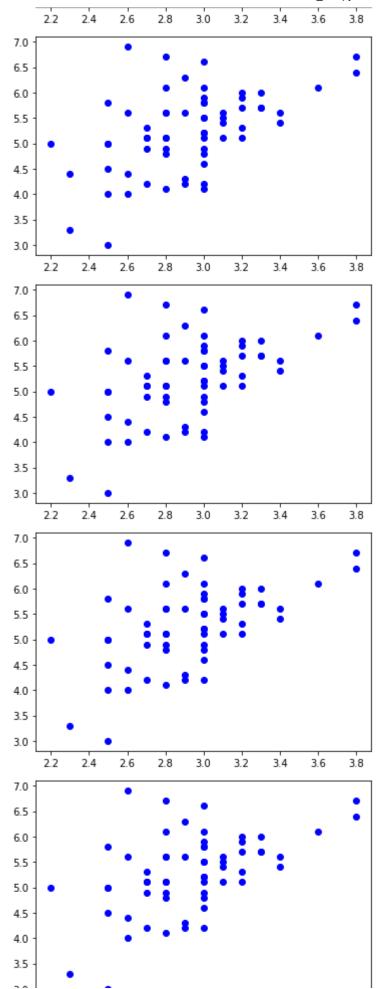


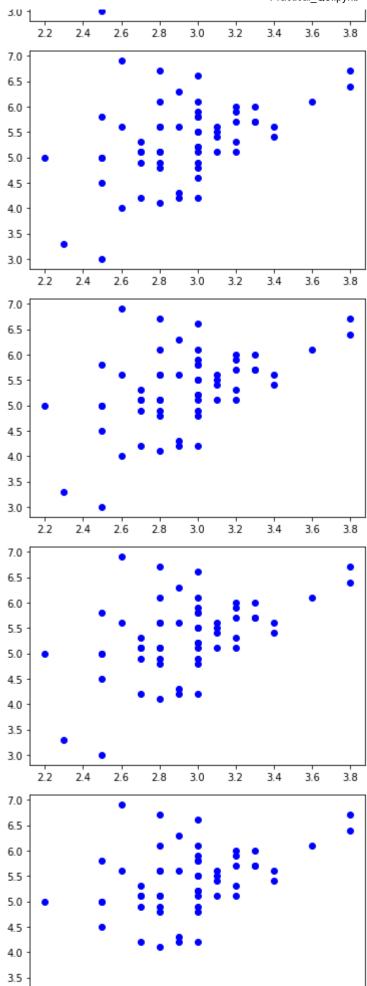


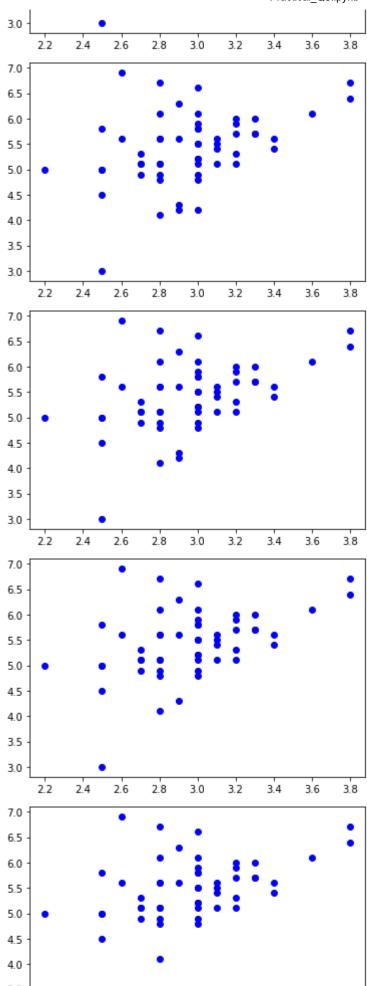


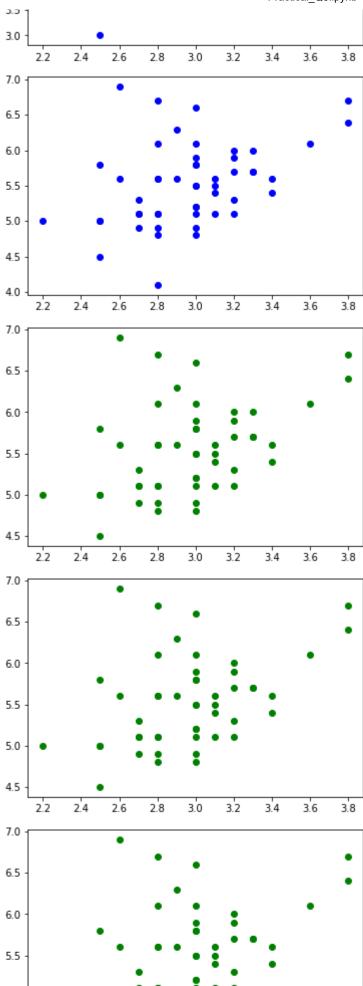


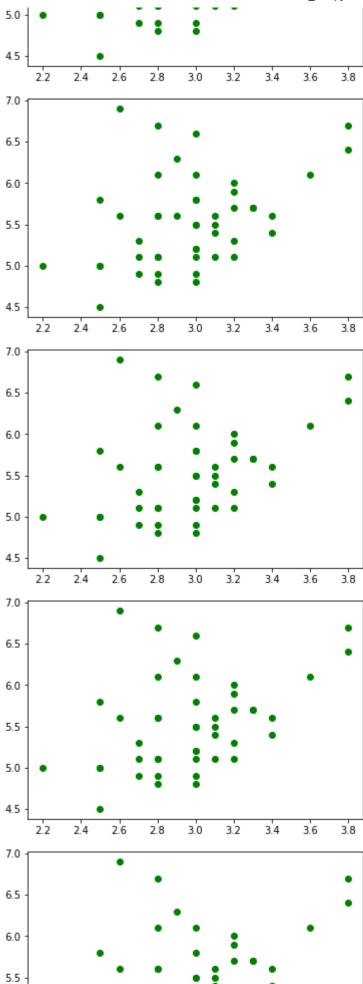


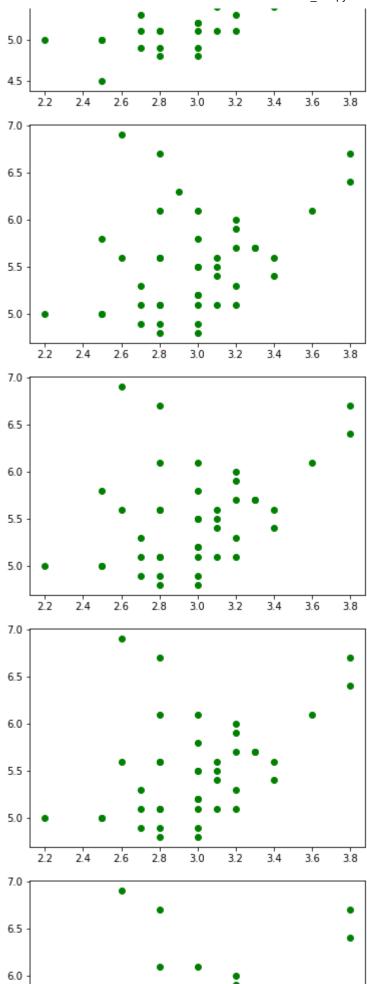


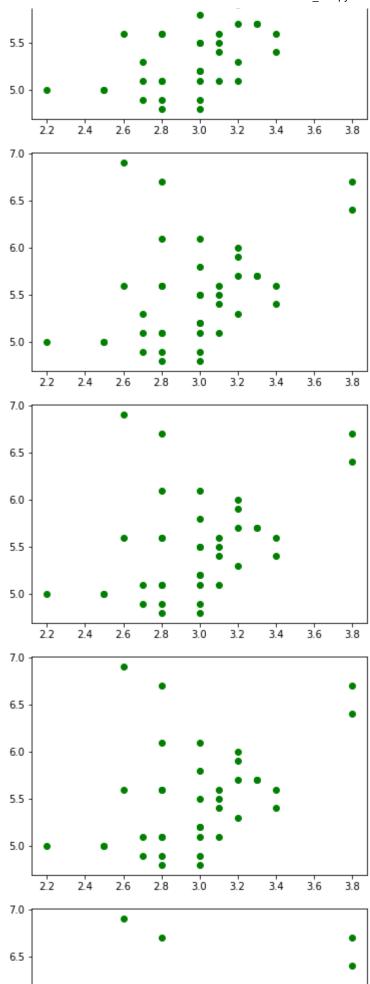


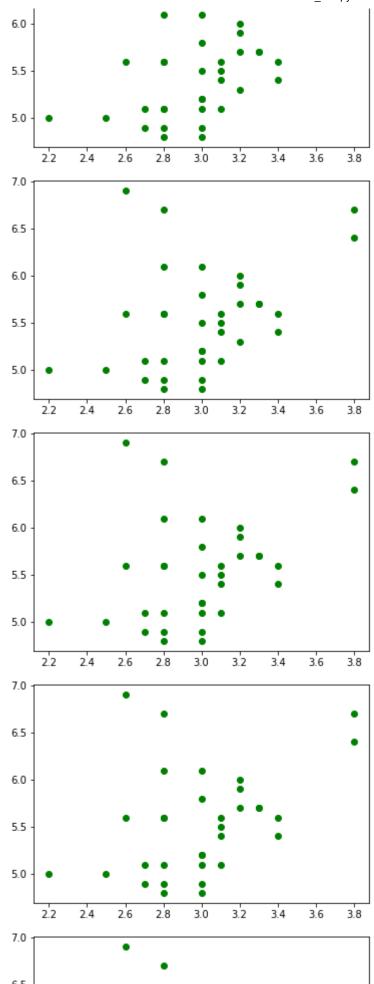


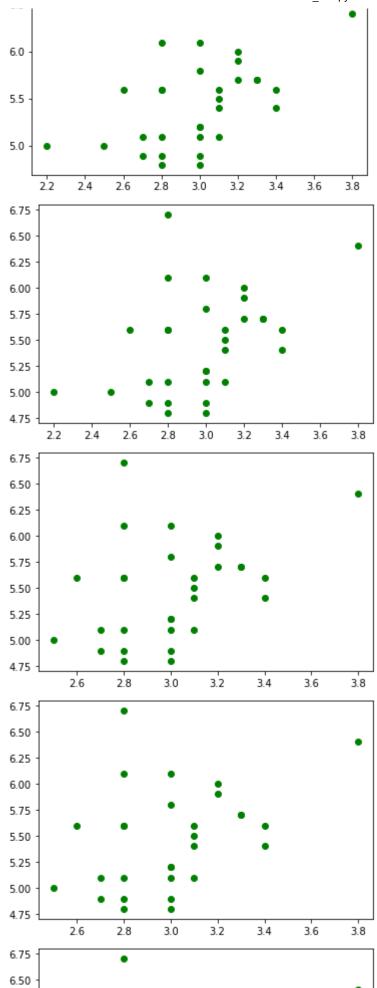


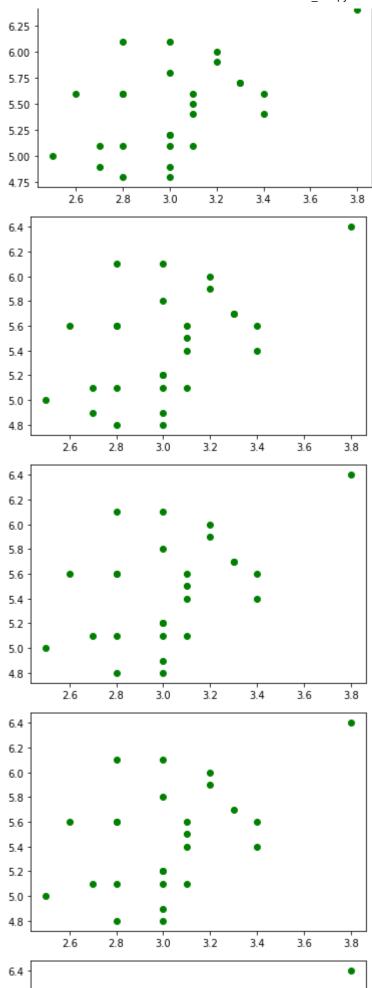


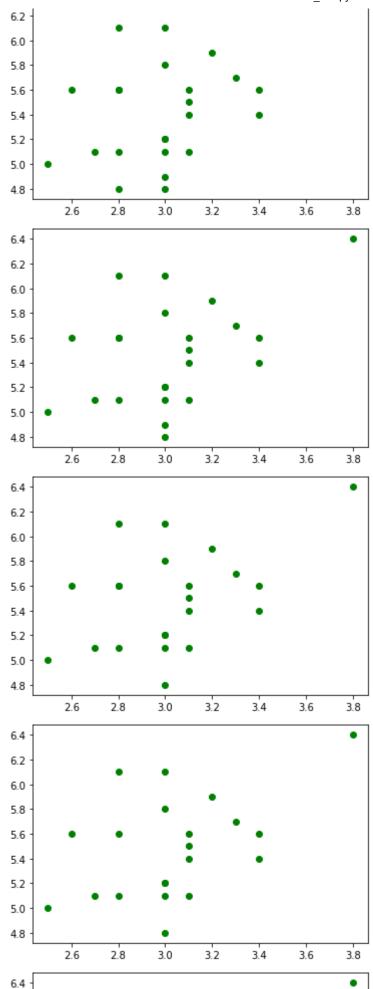


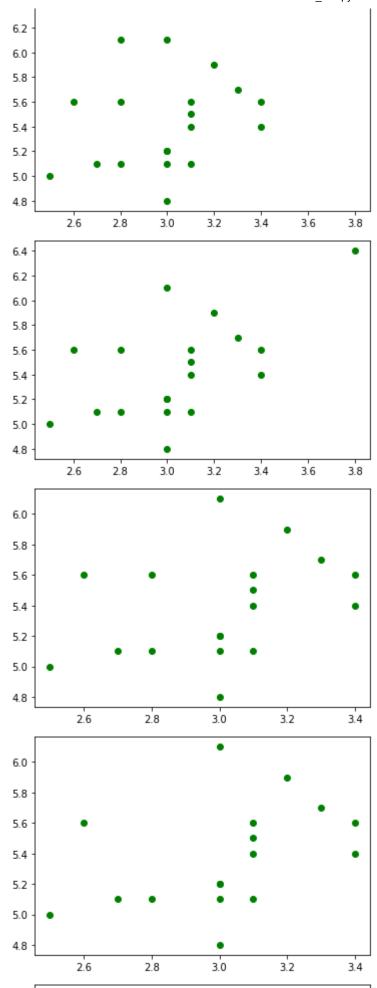


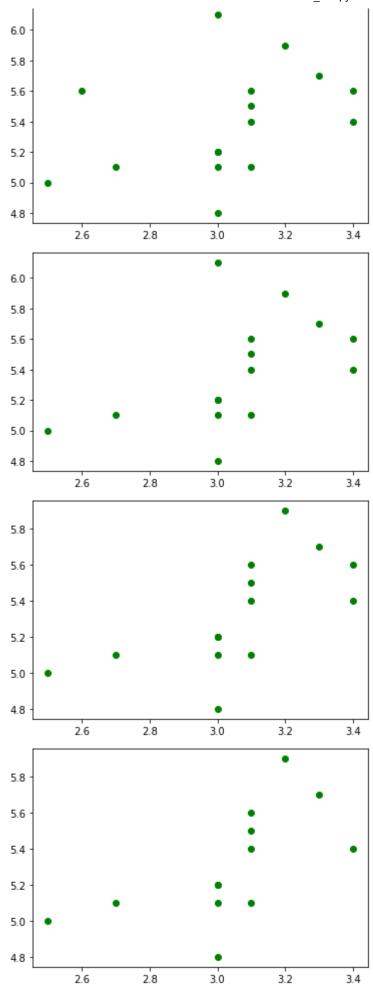


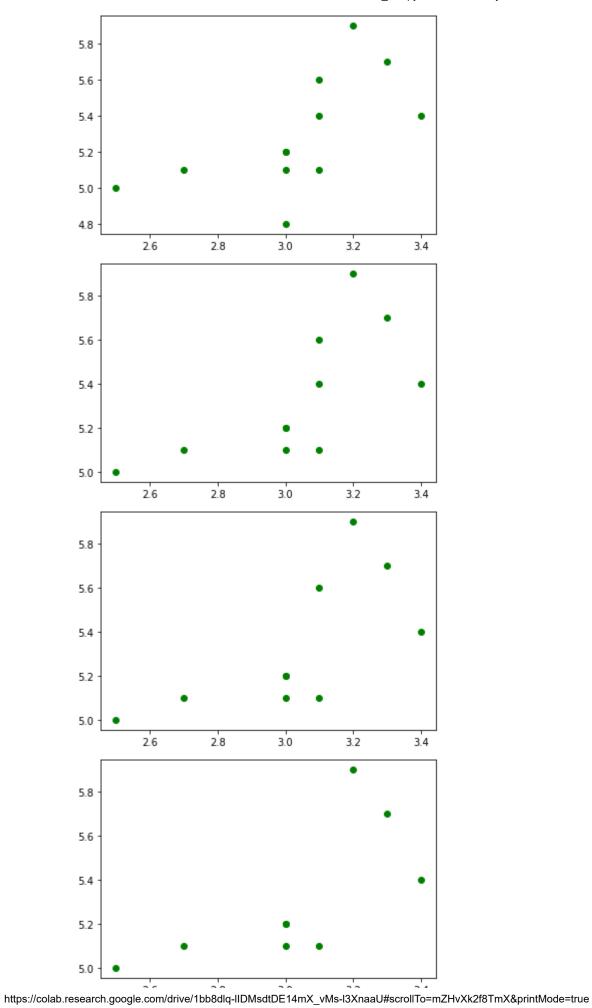


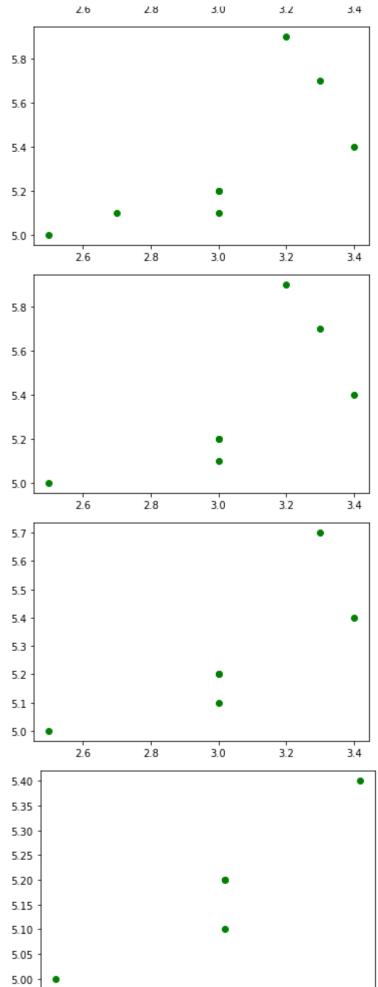


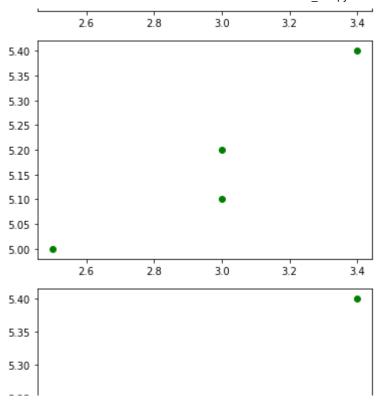






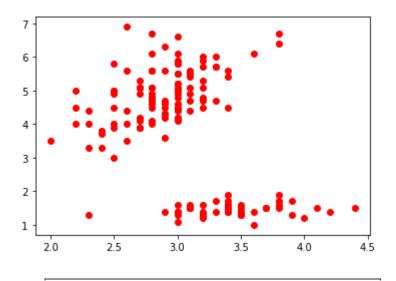






## Plotting unlabelized iris data set

plt.plot(iris.values[:,1],iris.values[:,2],'ro')
plt.show()

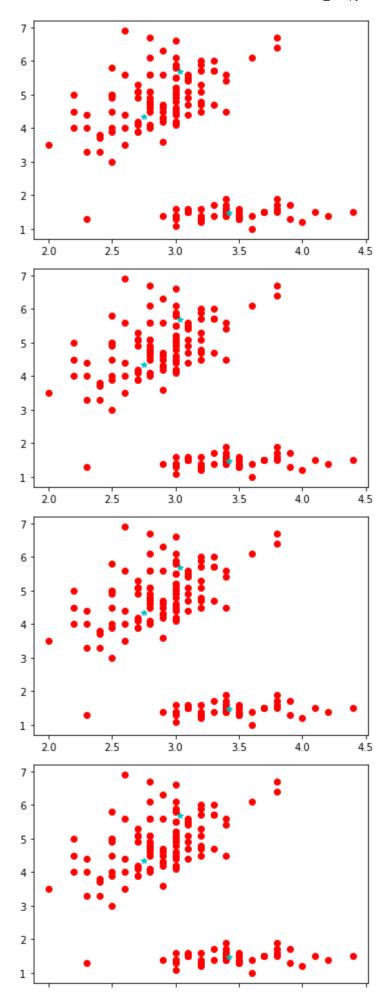


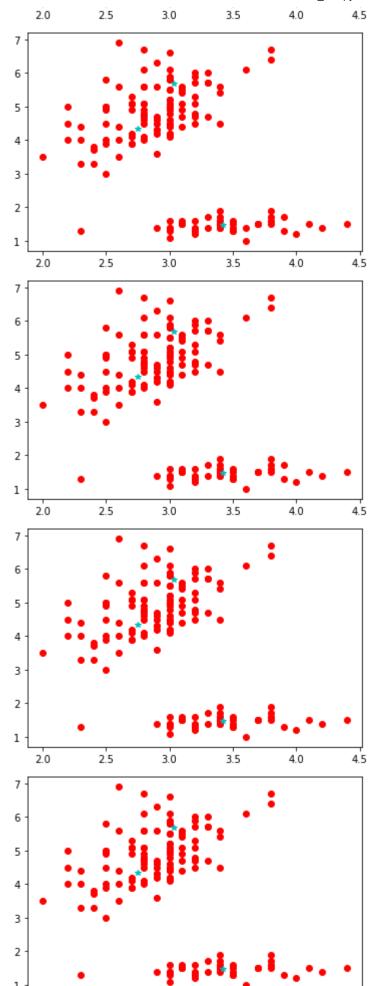
## Clustering using KMeans Clustering Algorithm

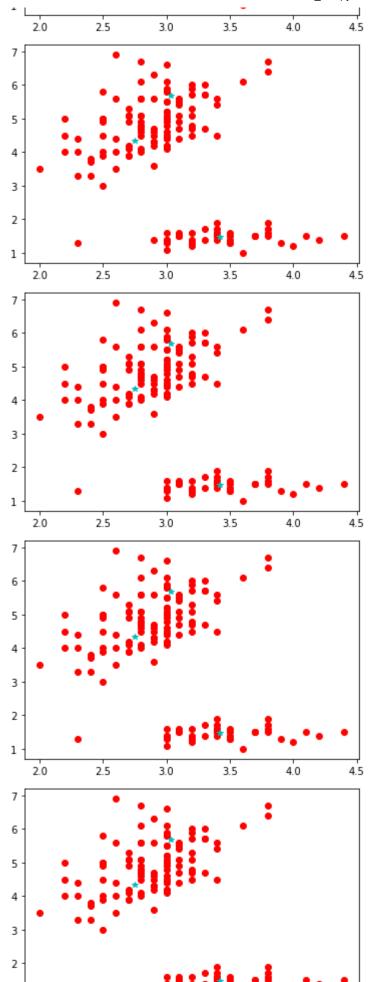
estimate1 = KMeans(n\_clusters=3)
estimate1.fit(iris.values[:,1:3])

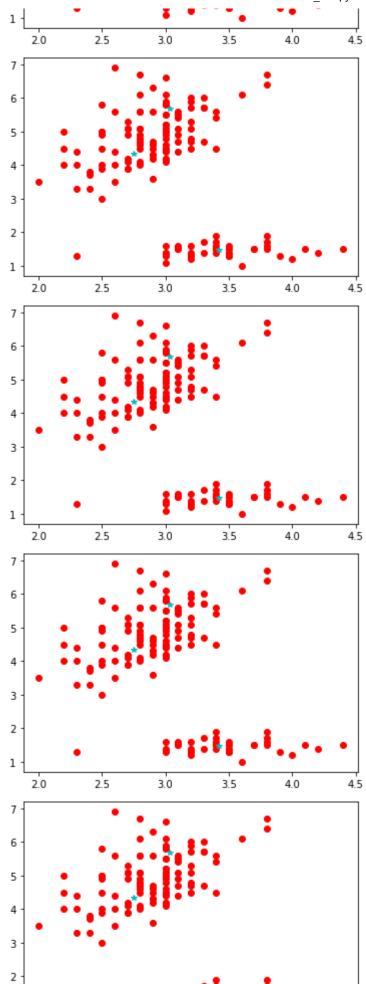
Plotting Clustered data points using K\_Means with 3 clusters

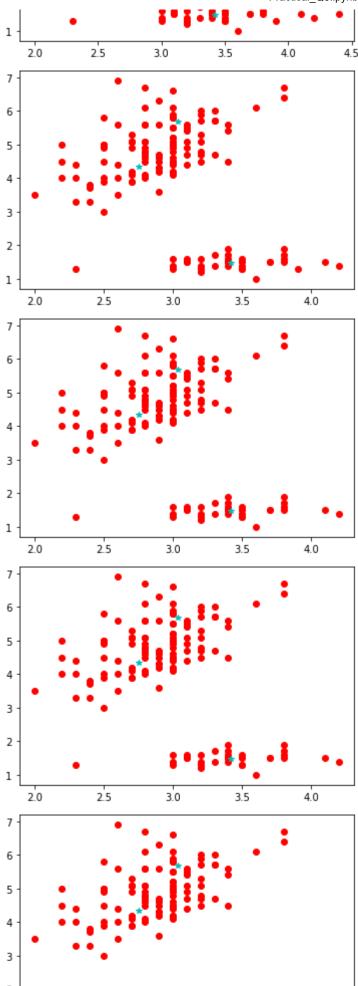
```
for i in range(150):
    if estimate1.labels_[i]==0:
        plt.plot(iris.values[i:,1],iris.values[i:,2],'go')
        plt.plot(estimate1.cluster_centers_[:,0],estimate1.cluster_centers_[:,1],'c*')
    elif estimate1.labels_[i]==1:
        plt.plot(iris.values[i:,1],iris.values[i:,2],'ro')
        plt.plot(estimate1.cluster_centers_[:,0],estimate1.cluster_centers_[:,1],'c*')
    elif estimate1.labels_[i]==2:
        plt.plot(iris.values[i:,1],iris.values[i:,2],'bo')
        plt.plot(estimate1.cluster_centers_[:,0],estimate1.cluster_centers_[:,1],'c*')
    plt.show()
```

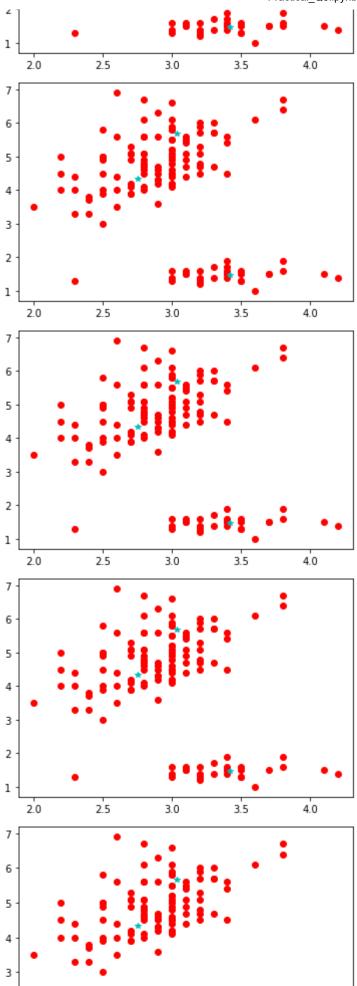


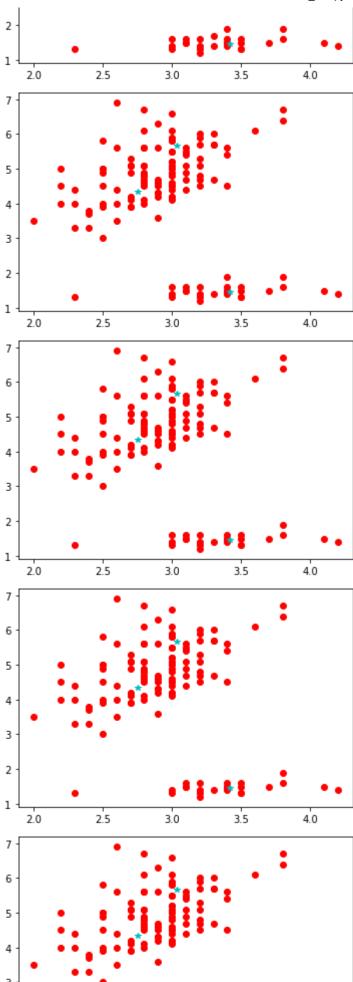


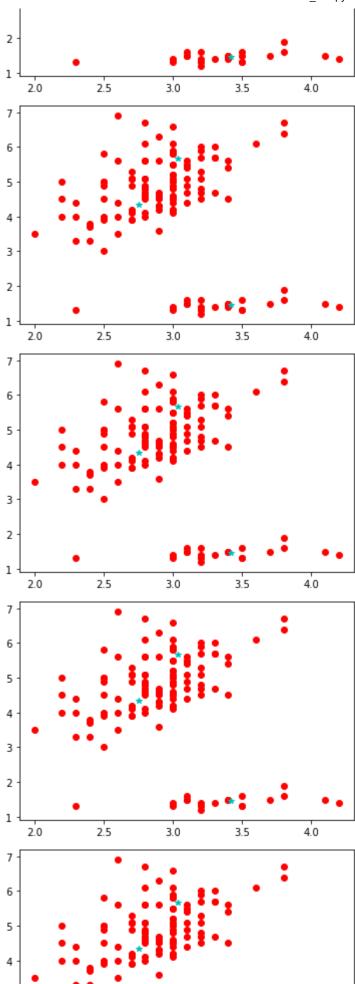


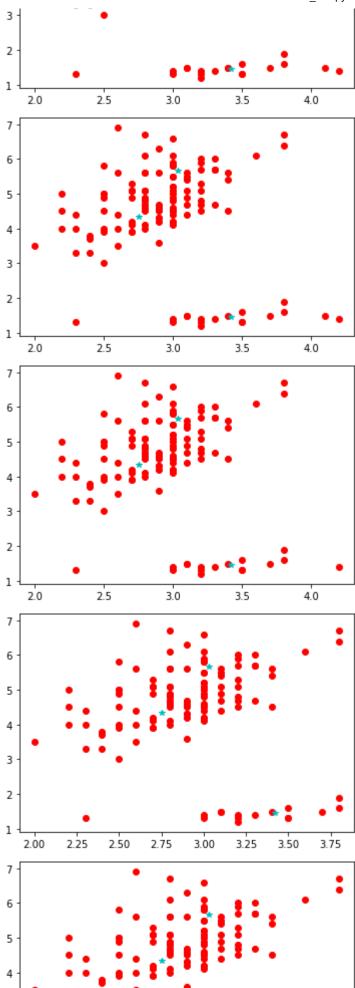


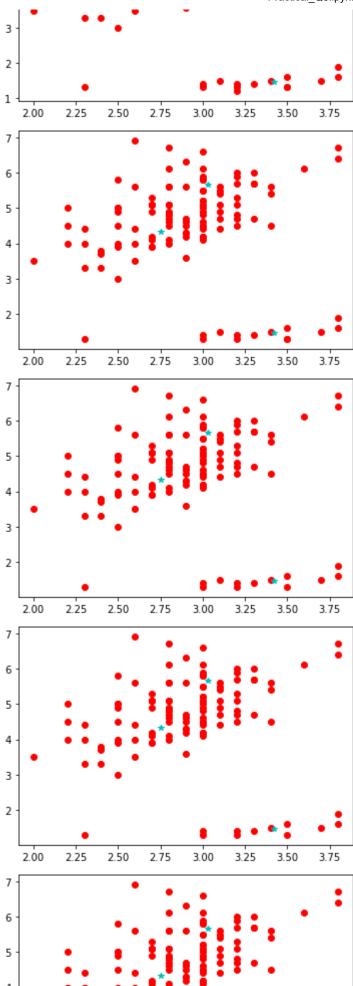


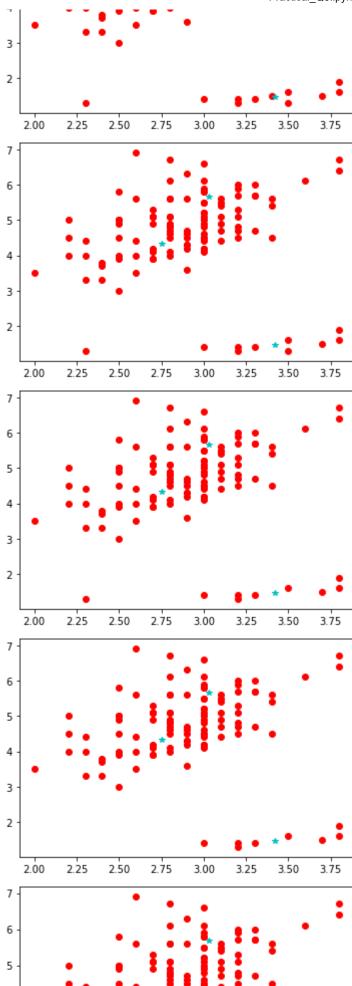


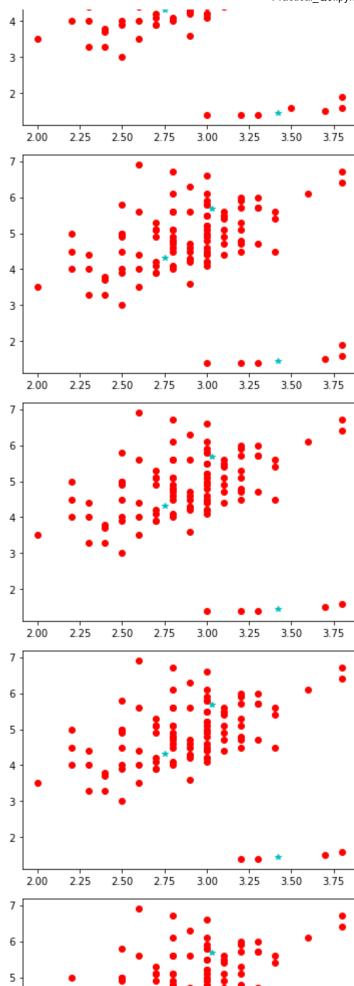


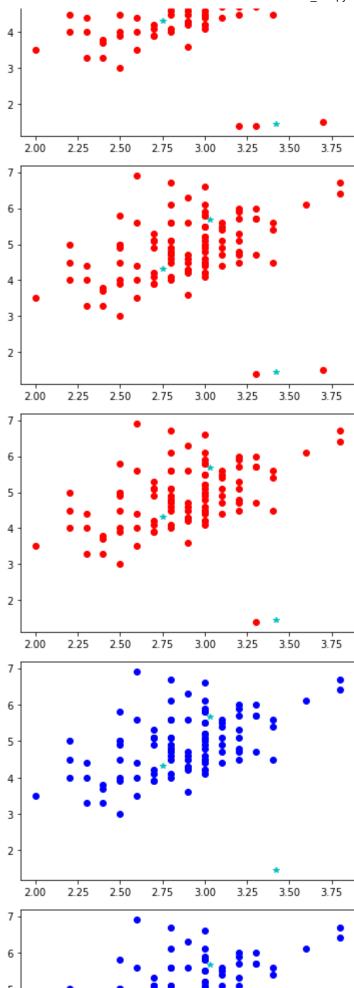


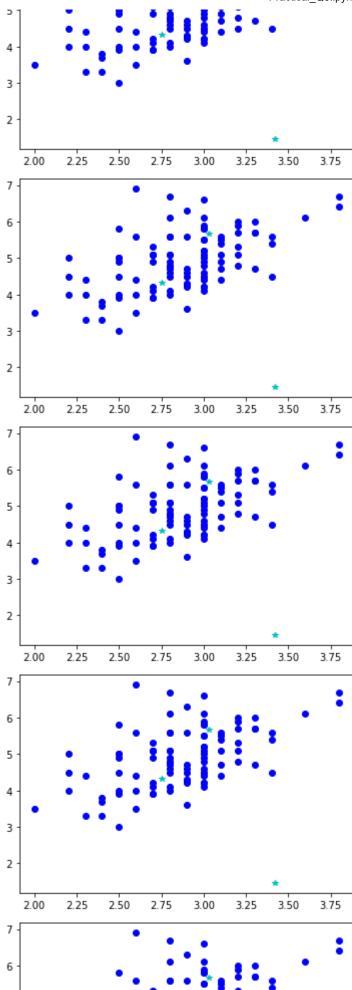


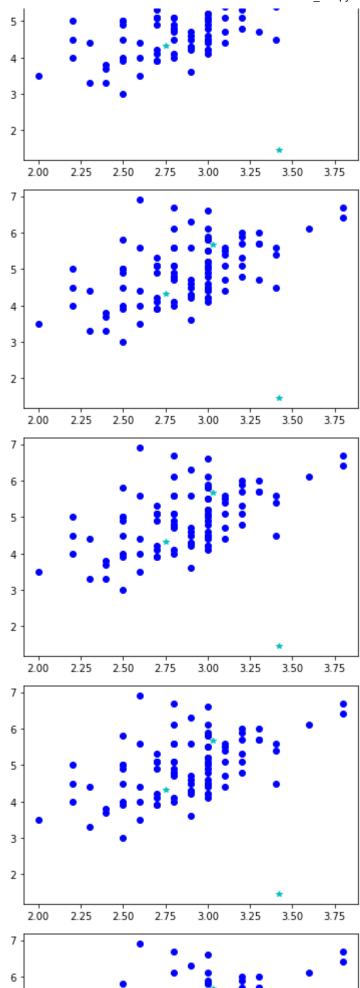


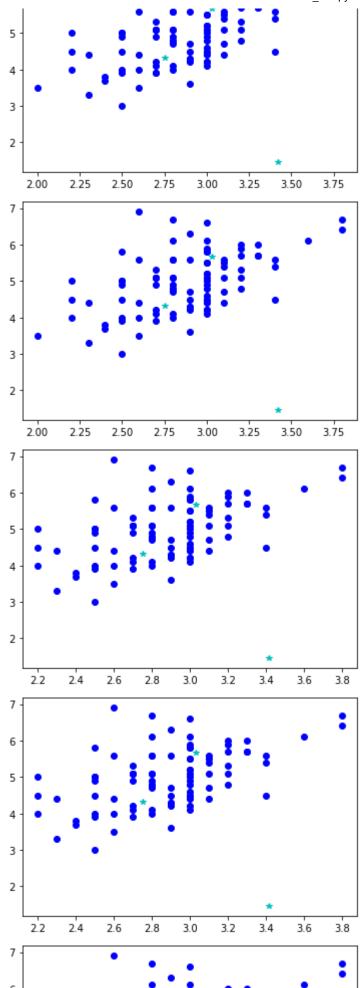


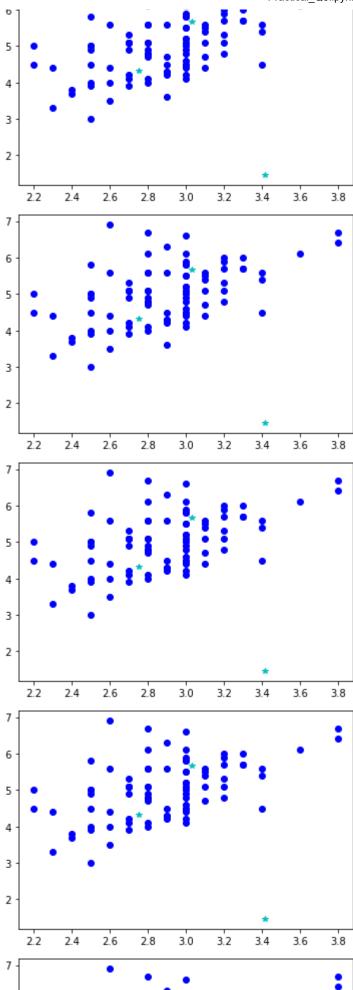


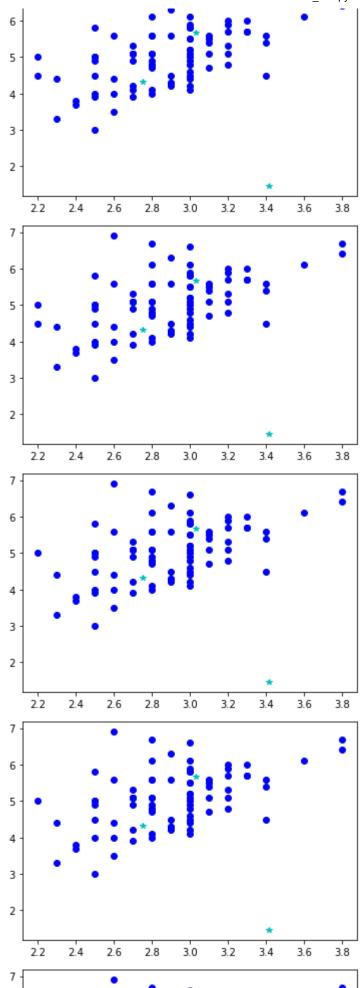


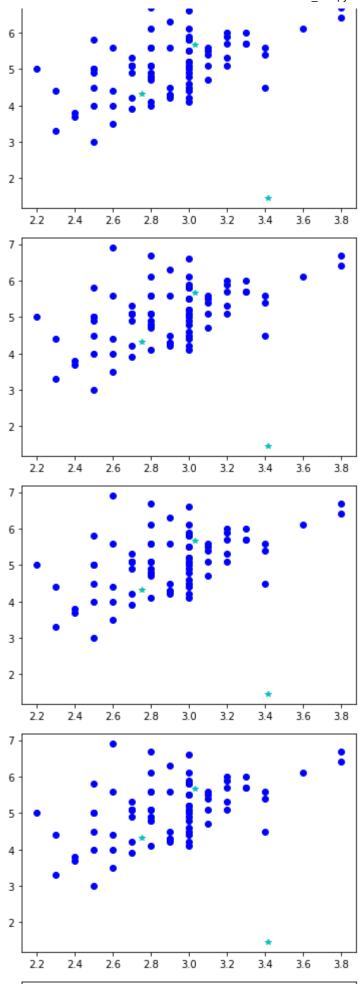


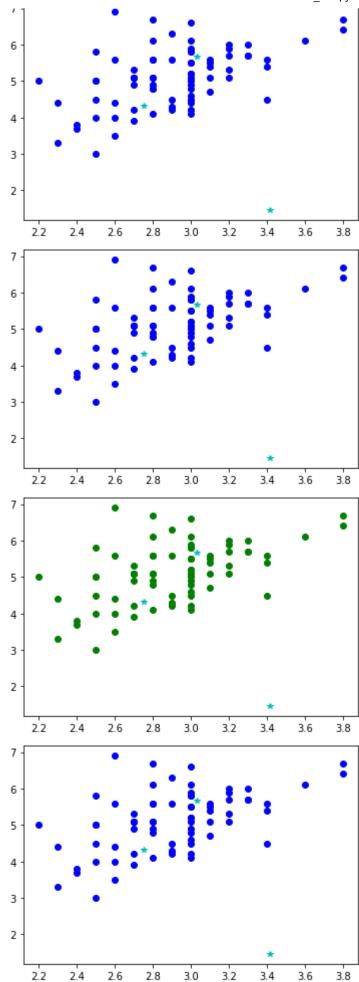


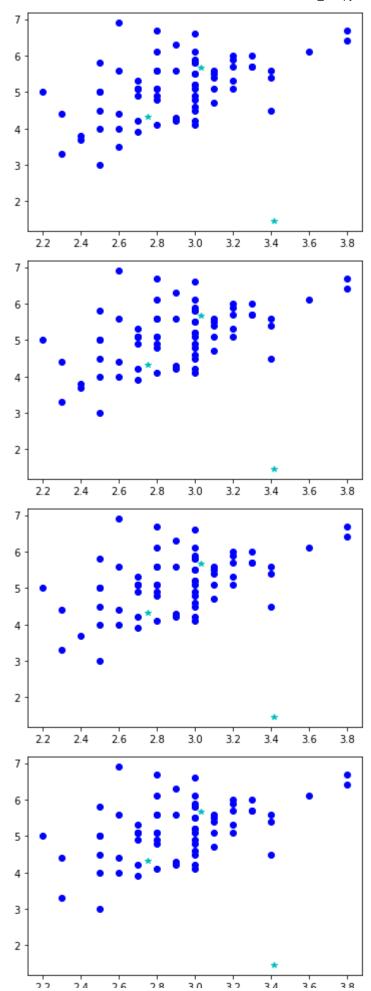


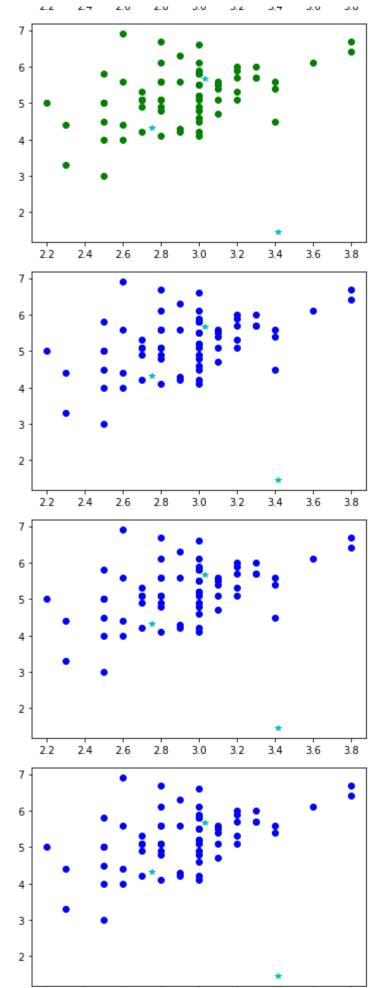


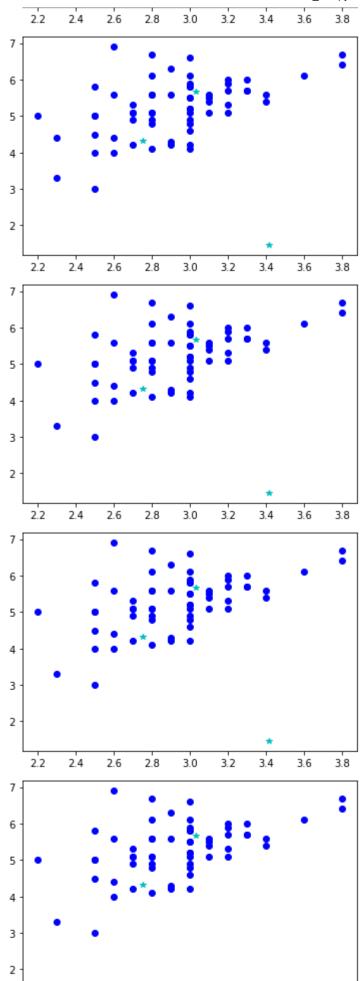


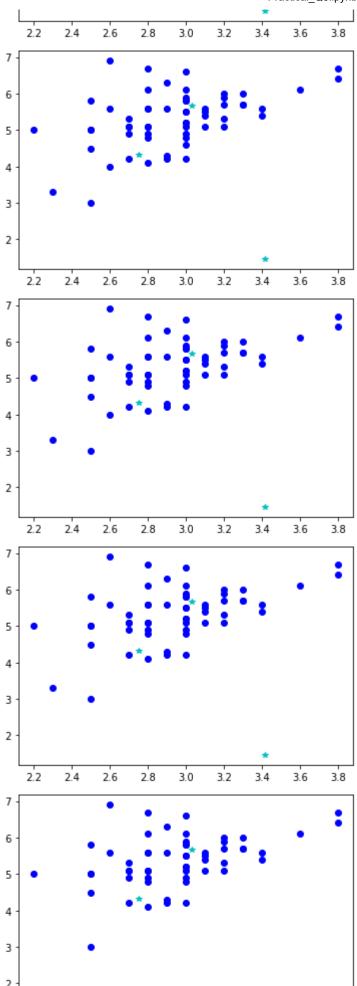


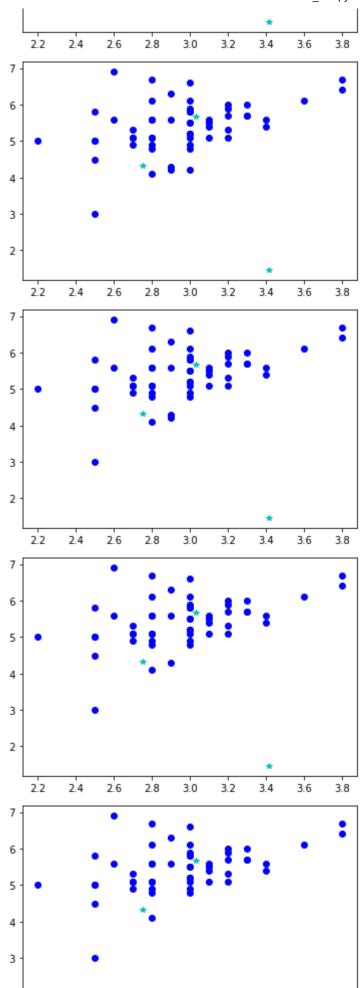


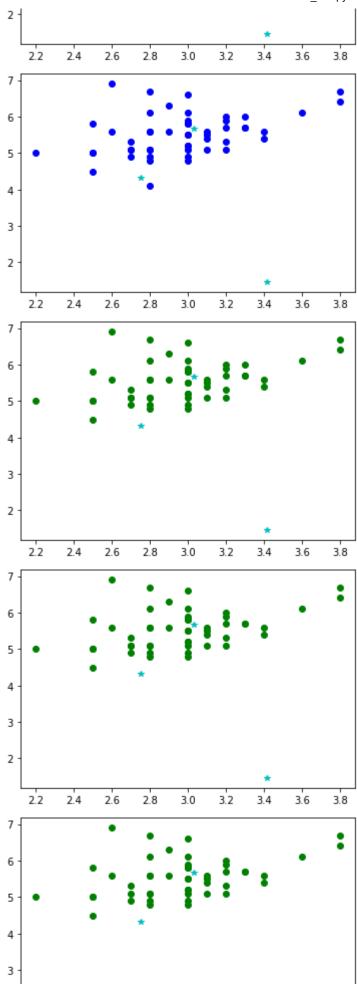


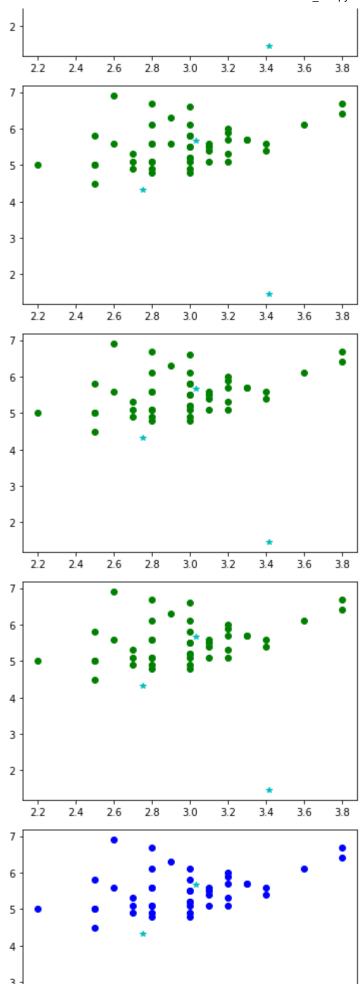


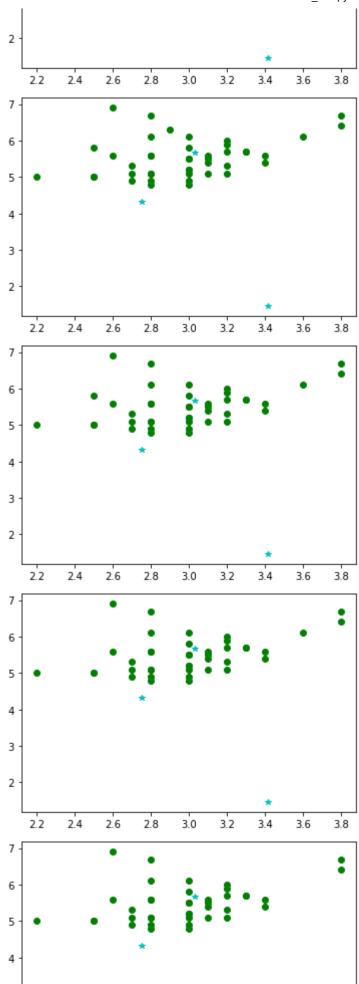


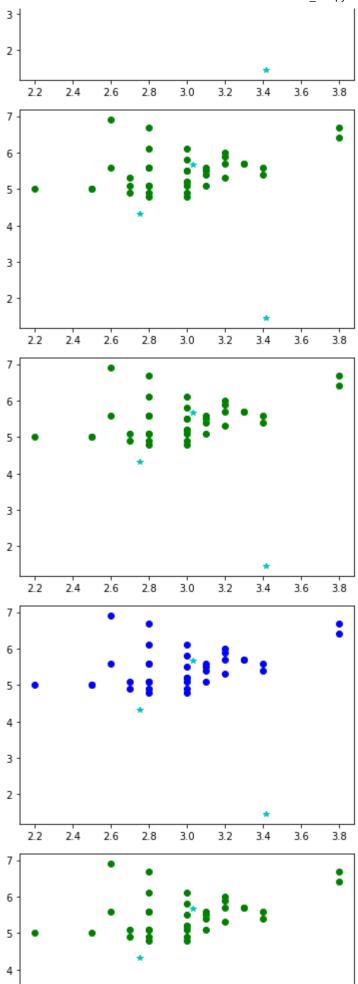


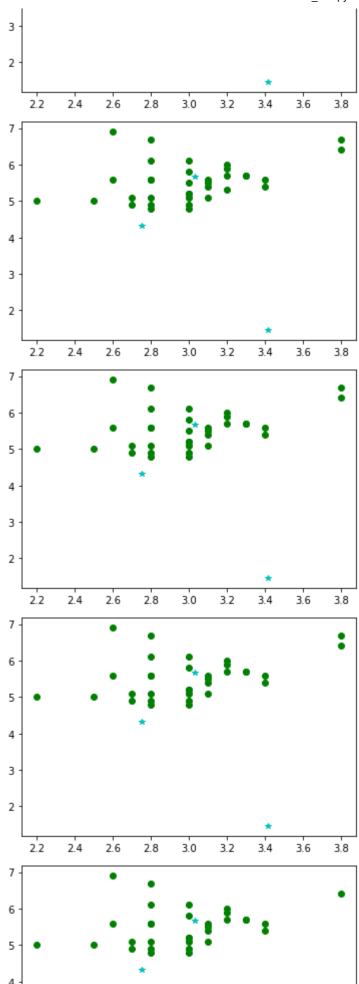


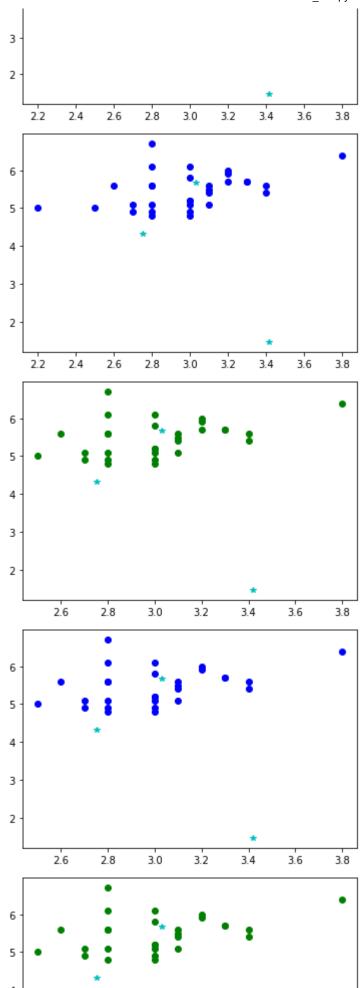


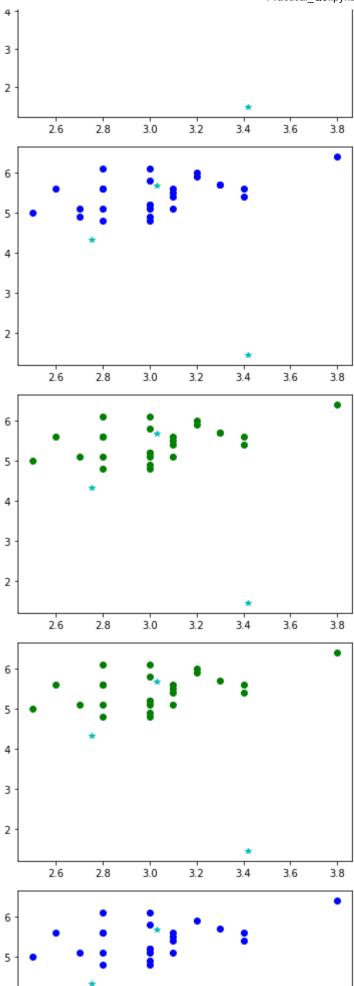


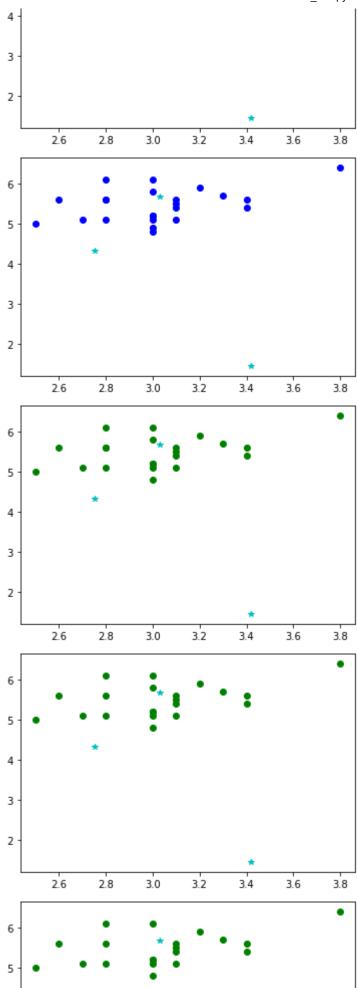


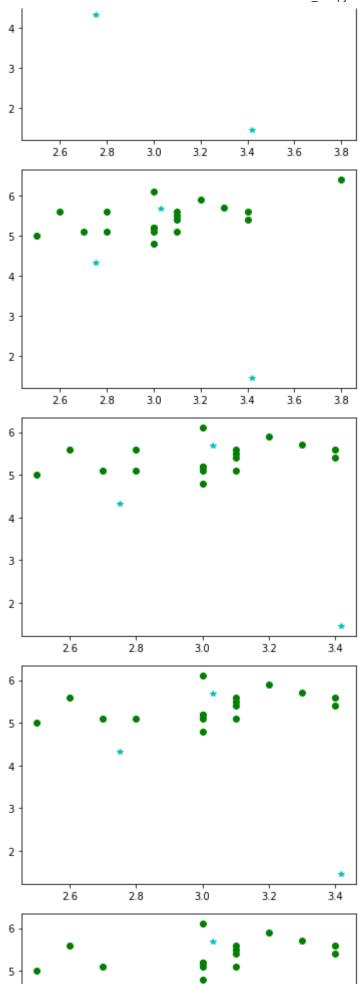


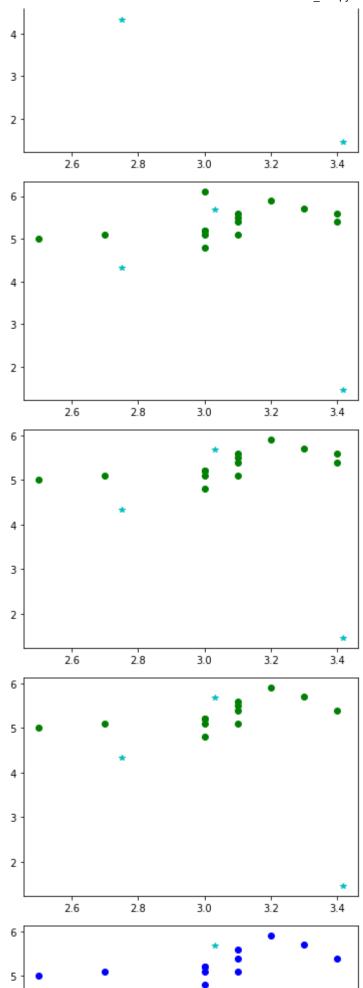


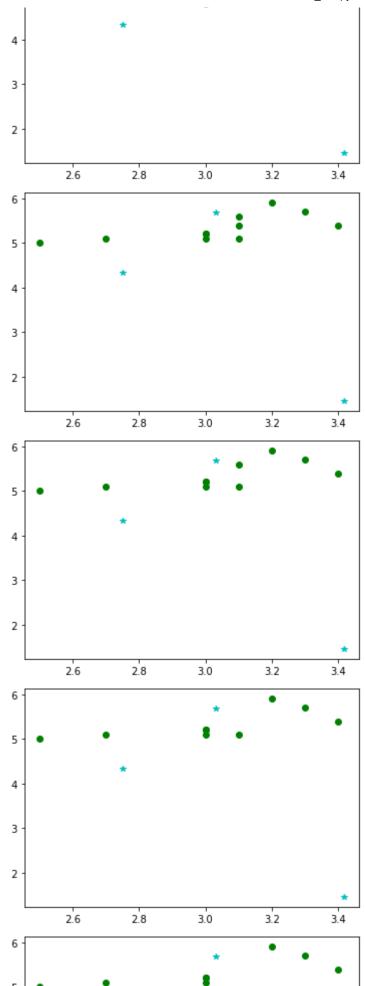


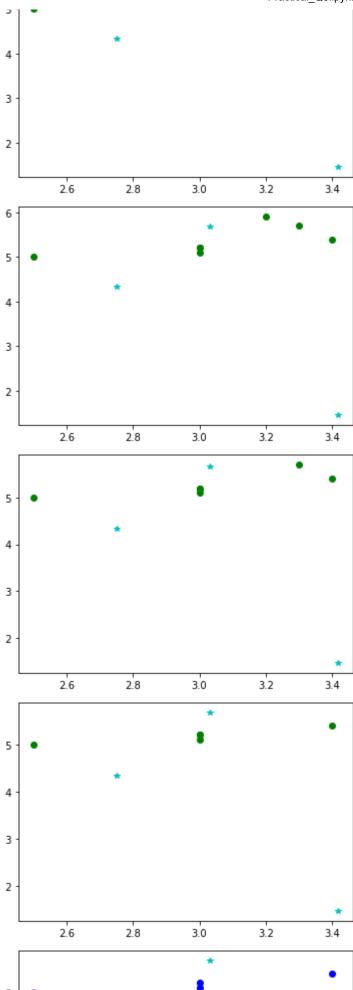


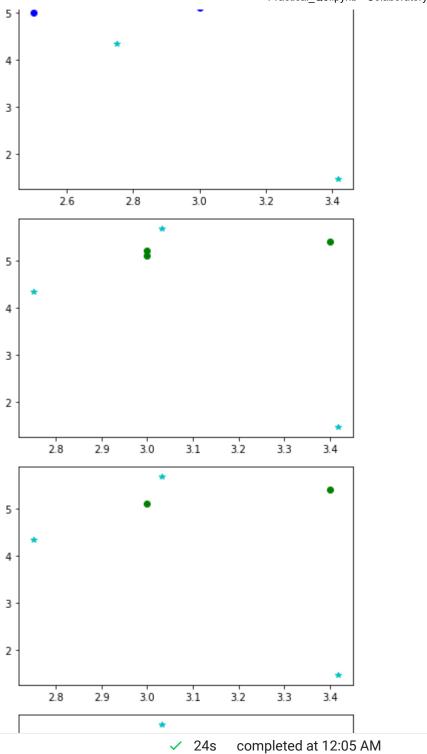












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