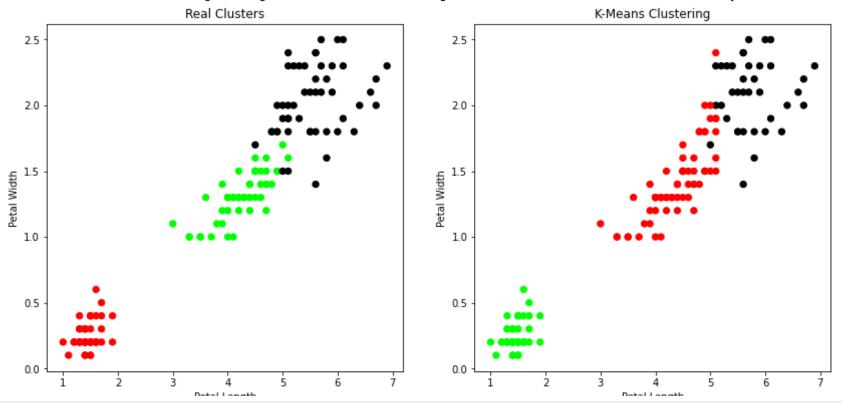
```
import matplotlib.pyplot as plt
In [2]:
         from sklearn import datasets
         from sklearn.cluster import KMeans
         import pandas as pd
         import numpy as np
         # import some data to play with
         iris = datasets.load iris()
         X = pd.DataFrame(iris.data)
         X.columns = ['Sepal Length', 'Sepal Width', 'Petal Length', 'Petal Width']
         y = pd.DataFrame(iris.target)
         y.columns = ['Targets']
         # Build the K Means Model
         model = KMeans(n clusters=3)
         model.fit(X)
         # model.labels : Gives cluster no for which samples belongs to
         # # Visualise the clustering results
         plt.figure(figsize=(14,14))
         colormap = np.array(['red', 'lime', 'black'])
         # Plot the Original Classifications using Petal features
         plt.subplot(2, 2, 1)
         plt.scatter(X.Petal Length, X.Petal Width, c=colormap[y.Targets], s=40)
         plt.title('Real Clusters')
         plt.xlabel('Petal Length')
         plt.vlabel('Petal Width')
         # Plot the Models Classifications
         plt.subplot(2, 2, 2)
         plt.scatter(X.Petal Length, X.Petal Width, c=colormap[model.labels], s=40)
         plt.title('K-Means Clustering')
         plt.xlabel('Petal Length')
         plt.ylabel('Petal Width')
         # General EM for GMM
         from sklearn import preprocessing
         # transform your data such that its distribution will have a
         # mean value 0 and standard deviation of 1.
```

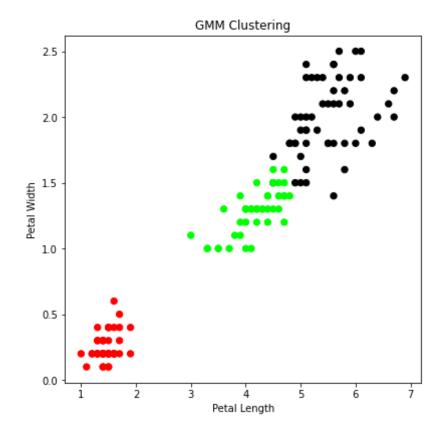
```
scaler = preprocessing.StandardScaler()
scaler.fit(X)
xsa = scaler.transform(X)
xs = pd.DataFrame(xsa, columns = X.columns)
from sklearn.mixture import GaussianMixture
gmm = GaussianMixture(n_components=3)
gmm.fit(xs)
gmm_y = gmm.predict(xs)
plt.subplot(2, 2, 3)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[gmm_y], s=40)
plt.title('GMM Clustering')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
print('Observation: The GMM using EM algorithm based clustering matched the true labels more closely than the Kmeans
```

Matplotlib is building the font cache; this may take a moment.

Observation: The GMM using EM algorithm based clustering matched the true labels more closely than the Kmeans.



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In []: