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In [2]:
import matplotlib.pyplot as plt
from sklearn import datasets
from sklearn.cluster import KMeans
import pandas as pd
import numpy as np
from sklearn import preprocessing
from sklearn.mixture import GaussianMixture
%matplotlib inline
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']
model=KMeans(n clusters=3)
model.fit(X)
plt.figure(figsize=(15,10))
colormap=np.array(['red','lime','black'])
plt.subplot(2,2,1)
plt.scatter(X.petal_length,X.petal_width,c=colormap[Y.Targets],s=60,marker='*')
plt.title('Real clusters')
plt.xlabel('petal length')
plt.vlabel('petal width')
plt.subplot(2,2,2)
plt.scatter(X.petal length,X.petal width,c=colormap[model.labels ],s=60)
plt.title('K means clustering')
plt.xlabel('petal length')
plt.vlabel('petal width')
scales=preprocessing.StandardScaler()
scales.fit(X)
xsa=scales.transform(X)
xs=pd.DataFrame(xsa,columns=X.columns)
gmm=GaussianMixture(n_components=3)
gmm.fit(xs)
gmm_y=gmm.predict(xs)
plt.subplot(2,2,3)
colorm=np.array(['red','lime','black'])
plt.scatter(X.petal length, X.petal width, c=colorm[gmm y], s=60)
plt.title('EM clustering')
plt.xlabel('petal length')
plt.ylabel('petal width')
print('Observation: The GMM using EM algorithm based clustering matched the true labels than the KMeans')
```

Observation: The GMM using EM algorithm based clustering matched the true labels than the KMeans



