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In [ ]:
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# Name : Meet Trivedi
# Student Id; N01520331
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Lab 9- K Mean Clustering

Import Libraries

In [1]:

```
import numpy as np
import pandas as pd

import matplotlib.pyplot as plt

import sklearn
from sklearn.preprocessing import scale
import sklearn.metrics as sm
from sklearn.metrics import confusion_matrix, classification_report
from sklearn.cluster import KMeans
from mpl_toolkits.mplot3d import Axes3D
from sklearn import datasets
%matplotlib inline
```

Get the Data(Wine dataset)

In [12]:

```
df = datasets.load_wine()
df
```

Out[12]:

```
{'data': array([[1.423e+01, 1.710e+00, 2.430e+00, ..., 1.040e+00, 3.920e+00,
      1.065e+03],
      [1.320e+01, 1.780e+00, 2.140e+00, ..., 1.050e+00, 3.400e+00,
      1.050e+03],
      [1.316e+01, 2.360e+00, 2.670e+00, ..., 1.030e+00, 3.170e+00,
      1.185e+03],
      [1.327e+01, 4.280e+00, 2.260e+00, ..., 5.900e-01, 1.560e+00,
      8.350e+02],
      [1.317e+01, 2.590e+00, 2.370e+00, ..., 6.000e-01, 1.620e+00,
      8.400e+02],
      [1.413e+01, 4.100e+00, 2.740e+00, ..., 6.100e-01, 1.600e+00,
      5.600e+02]]),
0, 0, 0,
      2, 2]),
'frame': None,
'target_names': array(['class_0', 'class_1', 'class_2'], dtype='<U7'),
'DESCR': '.. _wine_dataset:\n\nWine recognition dataset\n-----
----\n\n**Data Set Characteristics:**\n\n
                                   :Number of Instances: 178 (50
in each of three classes)\n : Number of Attributes: 13 numeric, predictive
                      :Attribute Information:\n \t\t- Alcohol\n \t\t
attributes and the class\n
- Malic acid\n \t\t- Ash\n\t\t- Alcalinity of ash \n \t\t- Magnesium\n\t\t-
Total phenols\n \t\t- Flavanoids\n \t\t- Nonflavanoid phenols\n \t\t- Proant
hocyanins\n\t\t- Color intensity\n \t\t- Hue\n \t\t- OD280/OD315 of diluted
wines\n \t\t- Proline\n\n
                     - class:∖n
                                       - class 0\n
                - class 2\n\t\t\n
                              :Summary Statistics:\n
class 1\n
Min
    Max
         Mean
                SD\n
= ====\n
         Alcohol:
                                 11.0 14.8
                                           13.0
                                                 0.8\n
                                                        Mal
ic Acid:
                    0.74 5.80
                               2.34 1.12\n
                                           Ash:
1.36 3.23
          2.36 0.27\n
                      Alcalinity of Ash:
                                              10.6 30.0
                                                        1
9.5
    3.3\n
                                  70.0 162.0
                                             99.7 14.3\n
                                                         Т
           Magnesium:
                      0.98 3.88
otal Phenols:
                                 2.29 0.63\n
                                             Flavanoids:
0.34 5.08
                      Nonflavanoid Phenols:
                                              0.13 0.66
          2.03 1.00\n
                                                        0.
36 0.12\n
          Proanthocyanins:
                                 0.41 3.58
                                            1.59 0.57\n
                                                        Co
lour Intensity:
                      1.3 13.0
                                 5.1
                                     2.3\n
                                            Hue:
0.48 1.71
          0.96 0.23\n
                      OD280/OD315 of diluted wines: 1.27 4.00
61 0.71\n
          Proline:
                                  278 1680
                                             746
                                                  315\n
:Missing Attribu
                :Class Distribution: class_0 (59), class_1 (71), class_
te Values: None\n
                             :Donor: Michael Marshall (MARSHALL%PL
         :Creator: R.A. Fisher\n
                  :Date: July, 1988\n\nThis is a copy of UCI ML Wine r
U@io.arc.nasa.gov)\n
ecognition datasets.\nhttps://archive.ics.uci.edu/ml/machine-learning-databa
ses/wine/wine.data\n\nThe data is the results of a chemical analysis of wine
s grown in the same\nregion in Italy by three different cultivators. There a
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re thirteen different\nmeasurements taken for different constituents found i n the three types of\nwine.\n\nOriginal Owners: \n\nForina, M. et al, PARVUS - \nAn Extendible Package for Data Exploration, Classification and Correlati on. \nInstitute of Pharmaceutical and Food Analysis and Technologies,\nVia B rigata Salerno, 16147 Genoa, Italy.\n\nCitation:\n\nLichman, M. (2013). UCI Machine Learning Repository\n[https://archive.ics.uci.edu/ml]. Irvine, CA: U niversity of California, \nSchool of Information and Computer Science. \n\n.. topic:: References\n\n (1) S. Aeberhard, D. Coomans and O. de Vel, \n Comp arison of Classifiers in High Dimensional Settings, \n Tech. Rep. no. 92-0 2, (1992), Dept. of Computer Science and Dept. of \n Mathematics and Stati stics, James Cook University of North Queensland. \n (Also submitted to Tec hnometrics). \n\n The data was used with many others for comparing various \n classifiers. The classes are separable, though only RDA \n has achieved 100% correct classification. \n (RDA: 100%, QDA 99.4%, LDA 98.9%, 1NN 96. 1% (z-transformed data)) \n (All results using the leave-one-out technique) \n\n (2) S. Aeberhard, D. Coomans and O. de Vel, \n "THE CLASSIFICATION PE RFORMANCE OF RDA" \n Tech. Rep. no. 92-01, (1992), Dept. of Computer Science e and Dept. of \n Mathematics and Statistics, James Cook University of Nort h Queensland. \n (Also submitted to Journal of Chemometrics).\n',

```
'feature_names': ['alcohol',
   'malic_acid',
   'ash',
   'alcalinity_of_ash',
   'magnesium',
   'total_phenols',
   'flavanoids',
   'nonflavanoid_phenols',
   'proanthocyanins',
   'color_intensity',
   'hue',
   'od280/od315_of_diluted_wines',
   'proline']}
```

In [19]:

```
v_names = df.feature_names
v_names
```

Out[19]:

```
['alcohol',
  'malic_acid',
  'ash',
  'alcalinity_of_ash',
  'magnesium',
  'total_phenols',
  'flavanoids',
  'nonflavanoid_phenols',
  'proanthocyanins',
  'color_intensity',
  'hue',
  'od280/od315_of_diluted_wines',
  'proline']
```

```
In [20]:
x = scale(df.data)
Х
Out[20]:
array([[ 1.51861254, -0.5622498 , 0.23205254, ..., 0.36217728,
         1.84791957,
                     1.01300893],
       [0.24628963, -0.49941338, -0.82799632, ..., 0.40605066,
         1.1134493 , 0.96524152],
       [0.19687903, 0.02123125, 1.10933436, ..., 0.31830389,
         0.78858745, 1.39514818],
       [0.33275817, 1.74474449, -0.38935541, ..., -1.61212515,
        -1.48544548, 0.28057537],
       [0.20923168, 0.22769377, 0.01273209, ..., -1.56825176,
       -1.40069891, 0.29649784],
       [1.39508604, 1.58316512, 1.36520822, ..., -1.52437837,
        -1.42894777, -0.59516041]])
In [21]:
y = pd.DataFrame(df.target)
У
Out[21]:
     0
     0
     0
  2
  3
     0
     0
  4
 173 2
 174 2
 175
     2
 176
     2
```

Apply Kmean Clustering

177 2

178 rows × 1 columns

In [34]:

```
clustering = KMeans(n_clusters=3, random_state=5)
clustering.fit(x)
```

Out[34]:

KMeans(n_clusters=3, random_state=5)

In [35]:

```
wine_df = pd.DataFrame(df.data)
wine_df.columns = ['alcohol','malic_acid', 'ash', 'alcalinity_of_ash', 'magnesium', 'total_
y.columns = ['Targets']
```

In [36]:

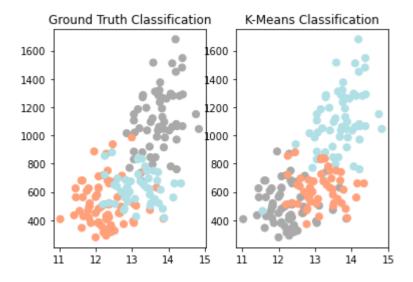
```
color_theme = np.array(['darkgray', 'lightsalmon', 'powderblue', 'darkblue', 'red', 'orange
plt.subplot(1,2,1)

plt.scatter(x=wine_df.alcohol, y=wine_df.proline, c=color_theme[df.target], s=50)
plt.title('Ground Truth Classification')

plt.scatter(x=wine_df.alcohol, y=wine_df.proline, c=color_theme[clustering.labels_], s=50)
plt.title('K-Means Classification')
```

Out[36]:

Text(0.5, 1.0, 'K-Means Classification')



Predictions and Evaluations

In [38]:

```
relabel = np.choose(clustering.labels_, [2, 0, 1]).astype(np.int64)

plt.subplot(1,2,1)

plt.scatter(x=wine_df.alcohol, y=wine_df.proline, c=color_theme[df.target], s=50)

plt.title('Ground Truth Classification')

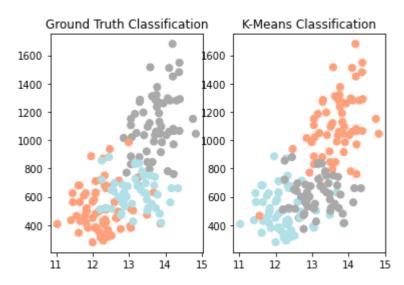
plt.subplot(1,2,2)

plt.scatter(x=wine_df.alcohol, y=wine_df.proline, c=color_theme[relabel], s=50)

plt.title('K-Means Classification')
```

Out[38]:

Text(0.5, 1.0, 'K-Means Classification')



In [39]:

print(classification_report(y, relabel))

	precision	recall	f1-score	support
0	0.00	0.00	0.00	59
1	0.05	0.04	0.05	71
2	0.00	0.00	0.00	48
accuracy			0.02	178
macro avg	0.02	0.01	0.02	178
weighted avg	0.02	0.02	0.02	178

In []: