K-means clustering

The notebook aims to study and implement a k-means clustering using "sklearn". A synthetic dataset will be used to identify clusters automatically using the K-means method.

Acknowledgments

• Inquiries: mauricio.antelis@tec.mx

Importing libraries

```
# Define where you are running the code: colab or local
RunInColab
              = True # (False: no | True: yes)
# If running in colab:
if RunInColab:
   # Mount your google drive in google colab
   from google.colab import drive
   drive.mount('/content/drive')
   # Find location
   #!pwd
   #!1s
   #!ls "/content/drive/My Drive/Colab Notebooks/MachineLearningWithPython/"
   # Define path del proyecto
                = "/content/drive/My Drive/content/drive/My Drive/"
   # Define path del proyecto
   Ruta
    Mounted at /content/drive
# Import the packages that we will be using
import numpy as np
                              # For array
                    # For data handling
import pandas as pd
import seaborn as sns
                               # For advanced plotting
# Note: specific functions of the "sklearn" package will be imported when needed to show concepts easily
```

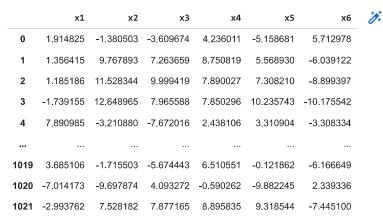
Importing data

```
# Dataset url
url = "drive/My Drive/datasets/SyntheticData4Clustering_X.csv"
# Load the dataset
df = pd.read_csv(url)
```

Undertanding and preprocessing the data

1. Get a general 'feel' of the data

```
# Print the dataframe df
```



get the number of observations and variables
observations = df.count()

print("Numero de observaciones: ", observations)

```
Numero de observaciones: x1 1024
x2 1024
x3 1024
x4 1024
x5 1024
x6 1024
dtype: int64
```

2. Drop rows with any missing values

```
# Drop rows with NaN values if existing
total_rows=len(df.axes[0]) #===> Axes of 0 is for a row
print("Numero de filas: "+str(total_rows))
```

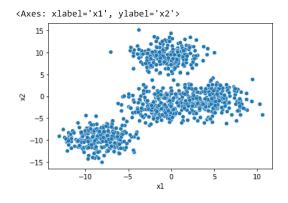
Print the new shape

Numero de filas: 1024

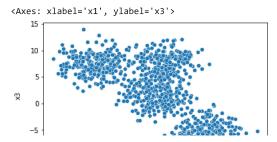
3. Scatterplot

Scatterplot of x1 and x2

sns.scatterplot(data=df, x="x1", y="x2")



Scatterplot of x1 and x3
sns.scatterplot(data=df, x="x1", y="x3")



Difficult to plot independetly all combinations, let's use pairplot

Pairplot: Scatterplot of all variables

g=.sns.pairplot(df,.corner=True,.diag_kind="kde")
g.map_lower(sns.kdeplot,levels=4,clors="2")
plt.show()

```
/usr/local/lib/python3.9/dist-packages/seaborn/distributions.py:1185: UserWarning: The following kwargs
 cset = contour_func(
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  cset = contour func(
/usr/local/lib/python3.9/dist-packages/seaborn/distributions.py:1185: UserWarning: The following kwargs
```

It looks like there are 3 or 4 clusters/groups

Note that we do not know in advance the class/cluster/group to which each point belongs to: we need to apply unsupervised learning;



Kmeans clustering

Kmeans clustering

```
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                          . .
# Import sklearn KMeans
from sklearn.cluster import KMeans
# Define number of clusters
K = 3 # Let's assume there are 2,3,4,5...? clusters/groups
#Crear el objeto o el modelo de MachineKMeans
km = KMeans(n_clusters =K,n_init="auto")
# Do K-means clustering (assing each point in the dataset to a cluster)
yestimated = km.fit predict(df)
# Print estimated cluster of each point in the dataset
yestimated
                       array([2, 1, 1, ..., 1, 2, 2], dtype=int32)
                                                                                                                                                                 56 A.O.
# Add a new column to the dataset with the cluster information
df2 = df.assign(yestimated=yestimated)
df2
```

```
x1
                            x2
                                      х3
                                                                       x6 yestimated
            1 914825 -1 380503 -3 609674
                                           4.236011
                                                     -5.158681
                                                                 5.712978
       0
                                                                                    1
            1.356415 9.767893 7.263659 8.750819
                                                      5.568930
                                                                                    0
       1
                                                                -6.039122
            4.405400 44.500044 0.000440 7.000007
                                                     7 000040
                                                                 0 000007
# Laber of the estimated clusters
df2.yestimated.unique()
    array([1, 2, 0], dtype=int32)
# Cluster centroides
km.cluster_centers_
    array([[ 1.85043266, -1.34592151, -2.11883656, 4.5718429 , -0.79519547,
             -0.55114018],
            [-8.3650671 , -9.59550917, 7.40711607, 3.77249056, -9.44226128,
            2.67666451],
[-0.44229417, 9.13121533, 7.61409814, 7.22984721, 8.13001382,
             -7.6264221 ]])
# Sum of squared error (sse) of the final model
km.inertia_
    44295.1263266536
# The number of iterations required to converge
km.n_iter_
    2
```

Important remarks

- · The number of each cluster is randomly assigned
- The order of the number in each cluster is random

Plot estimated clusters

Plot estimated clusters

```
# Get a dataframe with the data of each clsuter

df_1 = df2[df2.yestimated==0]
df_2 = df2[df2.yestimated==1]
df_3 = df2[df2.yestimated==2]

# Scatter plot of each cluster

plt.scatter(df_1.x1, df_1.x2, label="Cluster 0", c='r',marker='o',s=32,alpha=0.3)
plt.scatter(df_2.x1, df_2.x2, label="Cluster 1", c='g',marker='o',s=32,alpha=0.3)
plt.scatter(df_3.x1, df_3.x2, label="Cluster 2", c='b',marker='o',s=32,alpha=0.3)

plt.title("Scatter plot for each clutter")
plt.xlabel('x1')
plt.xlabel('x2')
plt.legend()
plt.show()
```

```
Scatter plot(for each clutter

15 Cluster 0
Cluster 1
Cluster 1
Cluster 2
```

→ Selecting K: elbow plot

```
Check the acurracy of the model using k-fold cross-validation
```

```
# Intialize a list to hold sum of squared error (sse)

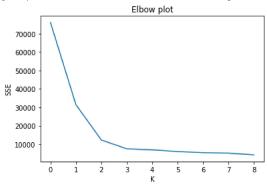
sse = []

# Define values of k
k_rng=range(1,10)

# For each k
for k in k_rng:
    km = KMeans(n_clusters=k,n_init="auto")
    km.fit_predict(df[['x1','x2']])
    sse.append(km.inertia_)

# Plot sse versus k
plt.title('Elbow plot')
plt.xlabel('K')
plt.ylabel('SSE')
plt.plot(sse)
```

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Choose the k after which the sse is minimally reduced

Important remarks

Observations?

Final remarks

- · K-Means clustering algorithm is perhaps the simplest and most popular unsupervised learning algorithm
- The number of clusters have to be defined by the user (i.e., by you ii)
- The number assigned to each cluster is randomly assigned from set 0, 1, 2
- · If there is no information about the number of clusters k, then use the elbow plot method to choose the best number of clusters k
- · The order of the number in each cluster is random
- The sklearn package provides the tools for data processing suchs as k-means

Activity:

- 1. Repeat this analysis using other pair of features, e.g., x3 and x6
- 2. Repeat this analysis using all six features, e.g., x1, x2,..., x6

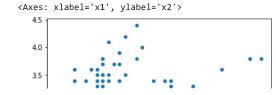
3. Provide conclusions

Activity: work with the iris dataset

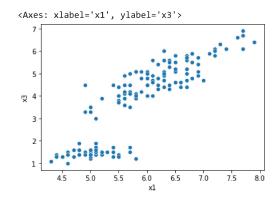
- 1. Do clustering with the iris flower dataset to form clusters using as features the four features
- 2. Do clustering with the iris flower dataset to form clusters using as features the two petal measurements: Drop out the other two features
- 3. Do clustering with the iris flower dataset to form clusters using as features the two sepal measurements: Drop out the other two features
- 4. Which one provides the better grouping? Solve this using programming skills, e.g., compute performance metrics

```
url_iris = "drive/My Drive/A01641179/datasets/iris/iris.csv"
df_iris = pd.read_csv(url_iris, header = None)
df_iris = df_iris.rename(columns={0: "x1"})
df_iris = df_iris.rename(columns={1: "x2"})
df_iris = df_iris.rename(columns={2: "x3"})
df_iris = df_iris.rename(columns={3: "x4"})
df_iris = df_iris.rename(columns={4: "x5"})
df_iris
                                          1
           x1 x2 x3 x4
                                    x5
       0 5.1 3.5 1.4 0.2
                             Iris-setosa
          4.9 3.0 1.4 0.2
                             Iris-setosa
         4.7 3.2 1.3 0.2
                             Iris-setosa
          4.6 3.1 1.5 0.2
                             Iris-setosa
          5.0 3.6 1.4 0.2
                             Iris-setosa
     145 6.7 3.0 5.2 2.3 Iris-virginica
     146 6.3 2.5 5.0 1.9 Iris-virginica
     147 6.5 3.0 5.2 2.0 Iris-virginica
     148 6.2 3.4 5.4 2.3 Iris-virginica
     149 5.9 3.0 5.1 1.8 Iris-virginica
     150 rows × 5 columns
observations_iris = df_iris.count()
print("Numero de observaciones: ", observations_iris)
    Numero de observaciones: x1
    x2
          150
    х3
          150
    х4
          150
     x5
          150
    dtype: int64
total_rows_iris=len(df_iris.axes[0]) #===> Axes of 0 is for a row
print("Numero de filas: "+str(total_rows_iris))
    Numero de filas: 150
```

sns.scatterplot(data=df_iris, x="x1", y="x2")



sns.scatterplot(data=df_iris, x="x1", y="x3")



g= sns.pairplot(df_iris, corner=True, diag_kind="kde")
g.map_lower(sns.kdeplot,levels=4,clors="2")
plt.show()

```
/usr/local/lib/python3.9/dist-packages/seaborn/distributions.py:1185: UserWarning: The following kwargs
      cset = contour func(
# Import sklearn KMeans
from sklearn.cluster import KMeans
# Define number of clusters
K = 4 # Let's assume there are 2,3,4,5...? clusters/groups
#Crear el objeto o el modelo de MachineKMeans
km_iris = KMeans(n_clusters =K,n_init="auto")
# Do K-means clustering (assing each point in the dataset to a cluster)
yestimated_iris = km_iris.fit_predict(df_iris.iloc[:,0:4])
# Print estimated cluster of each point in the dataset
yestimated_iris
    1, 1, 1, 1, 1, 1, 0, 0, 0, 3, 0, 3, 0, 3, 0, 3, 3, 3, 3, 0, 3, 0,
          3, 3, 0, 3, 0, 3, 0, 0, 0, 0, 0, 0, 0, 3, 3, 3, 3, 0, 3, 0, 0, 0,
          3, 3, 3, 0, 3, 3, 3, 3, 0, 3, 3, 2, 0, 2, 2, 2, 2, 3, 2, 2, 2,
          0, 0, 2, 0, 0, 2, 2, 2, 2, 0, 2, 0, 2, 0, 2, 2, 0, 0, 2, 2, 2, 2,
          2, 0, 0, 2, 2, 2, 0, 2, 2, 0, 2, 2, 2, 0, 0, 2, 0], dtype=int32)
        df2_iris = df_iris.assign(yestimated_iris=yestimated_iris)
df2_iris
                                                     1
          x1 x2 x3 x4
                                 x5 yestimated_iris
      0 5.1 3.5 1.4 0.2
                          Iris-setosa
                                                 1
        4.9 3.0 1.4 0.2
                           Iris-setosa
                                                 1
      2 4.7 3.2 1.3 0.2
                           Iris-setosa
        4.6 3.1 1.5 0.2
                          Iris-setosa
                                                 1
         5.0 3.6 1.4 0.2
                          Iris-setosa
                                                 1
     145 6.7 3.0 5.2 2.3 Iris-virginica
                                                 2
     146 6.3 2.5 5.0 1.9 Iris-virginica
                                                 0
     147 6.5 3.0 5.2 2.0 Iris-virginica
                                                 0
                                                 2
     148 6.2 3.4 5.4 2.3 Iris-virginica
     149 5.9 3.0 5.1 1.8 Iris-virginica
                                                 0
    150 rows × 6 columns
# Laber of the estimated clusters
df2_iris.yestimated_iris.unique()
    array([1, 0, 3, 2], dtype=int32)
# Cluster centroides
km_iris.cluster_centers_
                   , 2.855
                                          , 1.625
    array([[6.2525
                               , 4.815
                   , 3.428
                             , 1.462
           [5.006
                                          , 0.246
                     , 3.1
           [6.9125
                                , 5.846875 , 2.13125 ],
           [5.53214286, 2.63571429, 3.96071429, 1.22857143]])
# Sum of squared error (sse) of the final model
km iris.inertia
    57.22847321428572
# The number of iterations required to converge
km_iris.n_iter_
```

```
# Get a dataframe with the data of each clsuter

df_1_iris = df2_iris[df2_iris.yestimated_iris==0]

df_2_iris = df2_iris[df2_iris.yestimated_iris==1]

df_3_iris = df2_iris[df2_iris.yestimated_iris==2]

df_4_iris = df2_iris[df2_iris.yestimated_iris==2]

# Scatter plot of each cluster

plt.scatter(df_1_iris.x1, df_1_iris.x2, label="Cluster 0", c='r',marker='o',s=32,alpha=0.3)

plt.scatter(df_2_iris.x1, df_2_iris.x2, label="Cluster 1", c='g',marker='o',s=32,alpha=0.3)

plt.scatter(df_3_iris.x1, df_3_iris.x2, label="Cluster 2", c='b',marker='o',s=32,alpha=0.3)

plt.scatter(df_4_iris.x1, df_4_iris.x2, label="Cluster 2", c='r',marker='o',s=32,alpha=0.3)

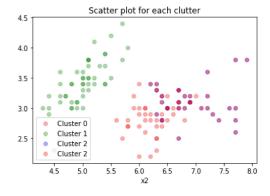
plt.title("Scatter plot for each clutter")

plt.xlabel('x1')

plt.xlabel('x2')

plt.legend()

plt.show()
```



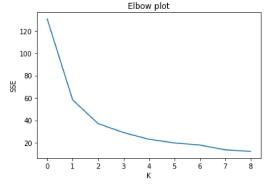
```
# Intialize a list to hold sum of squared error (sse)
sse = []

# Define values of k
k_rng=range(1,10)

# For each k
for k in k_rng:
    km_iris = KMeans(n_clusters=k,n_init="auto")
    km_iris.fit_predict(df_iris[['x1','x2']])
    sse.append(km_iris.inertia_)

# Plot sse versus k
plt.title('Elbow plot')
plt.xlabel('K')
plt.ylabel('SSE')
plt.plot(sse)
```

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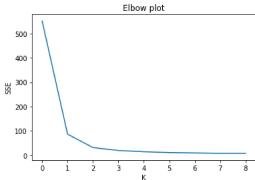
```
# Intialize a list to hold sum of squared error (sse)
sse = []
```

Define values of k

```
k_rng=range(1,10)
# For each k
for k in k_rng:
    km_iris = KMeans(n_clusters=k,n_init="auto")
    km_iris.fit_predict(df_iris[['x3','x4']])
    sse.append(km_iris.inertia_)

# Plot sse versus k
plt.title('Elbow plot')
plt.xlabel('K')
plt.ylabel('SSE')
plt.plot(sse)
```

[<matplotlib.lines.Line2D at 0x7fc4fdd4edc0>]



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