### K-means clustering

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```
In [114]: # 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/MachineLearningWithPyt
          hon/"
              # Define path del proyecto
                              = "/content/drive/My Drive/Colab Notebooks/TC1002S/N
          otebooksStudents/A01636995"
          else:
              # Define path del proyecto
                              = "/Users/pamelasanchez/Documents/TC1002S/NotebooksS
          tudents/A01636995"
```

Drive already mounted at /content/drive; to attempt to forcibly remoun t, call drive.mount("/content/drive", force\_remount=True).

```
In [120]: # Import the packages that we will be using
import numpy as np  # For array
import pandas as pd  # For data handling
import seaborn as sns  # For advanced plotting
import matplotlib.pyplot as plt  # For showing plots

# Note: specific functions of the "sklearn" package will be imported whe
n needed to show concepts easily
```

## Importing data

```
In [121]: # Dataset url
    url = Ruta + "/datasets/iris/iris.csv"

# Load the dataset
    df = df = pd.read_csv(url, header = None)
    # Column names are added to facilitate the rest of the work
    df = df.rename(columns={0: "Largo_Sepalo"})
    df = df.rename(columns={1: "Ancho_Sepalo"})
    df = df.rename(columns={2: "Largo_Petalo"})
    df = df.rename(columns={3: "Ancho_Petalo"})
    df = df.rename(columns={4: "Especie"})
In [116]: # Get a general 'feel' of the data
# Print the dataframe
```

Out[116]:

df

	Largo_Sepalo	Ancho_Sepalo	Largo_Petalo	Ancho_Petalo	Especie
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	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  $\times$  5 columns

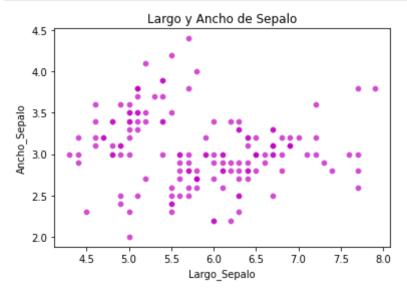
```
In [29]: # get the number of observations and variables
    df.shape

Out[29]: (150, 5)
In [93]: # Drop rows with NaN values if existing
```

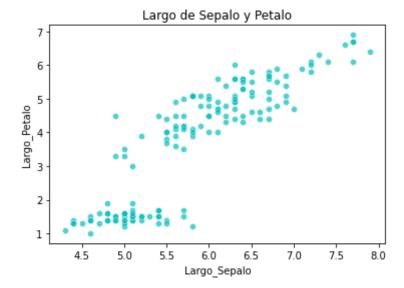
```
In [93]: # Drop rows with NaN values if existing
    df.dropna()
    # Print the new shape
    df.shape
```

Out[93]: (150, 5)

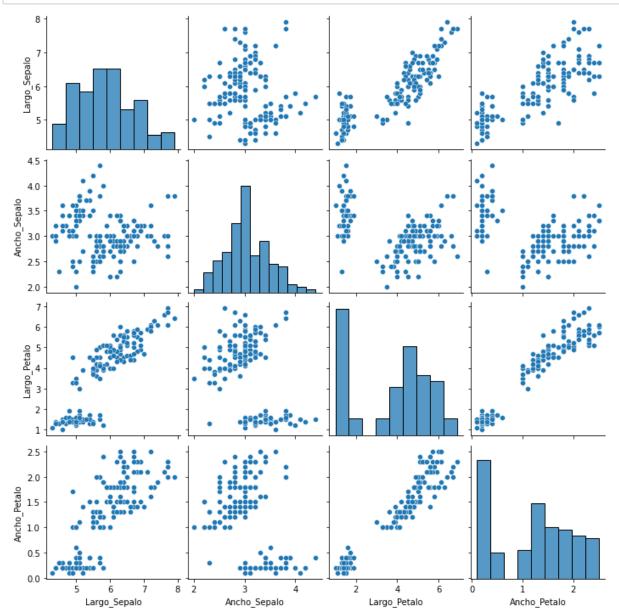
```
In [31]: # Scatterplot of x1 and x2
    sns.scatterplot(data = df, x = "Largo_Sepalo", y = "Ancho_Sepalo", c =
    "m", alpha = 0.7)
    plt.title("Largo y Ancho de Sepalo")
    plt.show()
```



```
In [32]: # Scatterplot of x1 and x3
sns.scatterplot(data = df, x = "Largo_Sepalo", y = "Largo_Petalo", c =
    "c", alpha = 0.7)
plt.title("Largo de Sepalo y Petalo")
plt.show()
```



In [33]: # Pairplot: Scatterplot of all variables
 sns.pairplot(df)
 plt.show()



```
In [122]: # The last column is dropped
    df.drop("Especie", axis=1, inplace = True)
    df
```

Out[122]:

	Largo_Sepalo	Ancho_Sepalo	Largo_Petalo	Ancho_Petalo
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

150 rows × 4 columns

# **Kmeans clustering**

```
In [95]: # Import sklearn KMeans
       from sklearn.cluster import KMeans
       # Define number of clusters
       K = 2# Let's assume there are 2,3,4,5...? clusters/groups
       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
0,
             0,
             0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1,
       1,
             1,
             1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
       1,
             1,
             32)
In [96]: # Add a new column to the dataset with the cluster information
       df.insert(4, "Cluster Information", yestimated, True)
       df.head()
Out[96]:
         Largo Sepalo Ancho Sepalo Largo Petalo Ancho Petalo Cluster Information
                                                       0
        0
               5.1
                        3.5
                                 1.4
                                          0.2
               4.9
                                          0.2
                                                       0
        1
                        3.0
                                 1.4
        2
               4.7
                        3.2
                                          0.2
                                                       0
                                 1.3
        3
               4.6
                        3.1
                                 1.5
                                          0.2
                                                       0
               5.0
                        3.6
                                 1.4
                                          0.2
                                                       0
In [97]: # Label of the estimated clusters
       df.Cluster Information.unique()
Out[97]: array([0, 1], dtype=int32)
In [98]:
       # Cluster centroides
       km.cluster centers
Out[98]: array([[5.00566038, 3.36981132, 1.56037736, 0.29056604],
             [6.30103093, 2.88659794, 4.95876289, 1.69587629]])
```

```
In [99]: # Sum of squared error (sse) of the final model
km.inertia_
Out[99]: 152.3479517603579

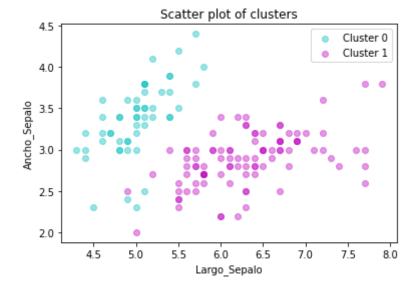
In [100]: # The number of iterations required to converge
km.n_iter_
Out[100]: 4
```

#### Plot estimated clusters

```
In [101]: # Get a dataframe with the data of each clsuter
    df1 = df[df.Cluster_Information == 0]
    df2 = df[df.Cluster_Information == 1]

# Scatter plot of each cluster
    plt.scatter(df1.Largo_Sepalo, df1.Ancho_Sepalo, label = "Cluster 0", c =
    'c', marker = 'o', alpha = 0.4)
    plt.scatter(df2.Largo_Sepalo, df2.Ancho_Sepalo, label = "Cluster 1", c =
    'm', marker = 'o', alpha = 0.4)

    plt.title("Scatter plot of clusters")
    plt.xlabel("Largo_Sepalo")
    plt.ylabel("Ancho_Sepalo")
    plt.legend()
    plt.show()
```

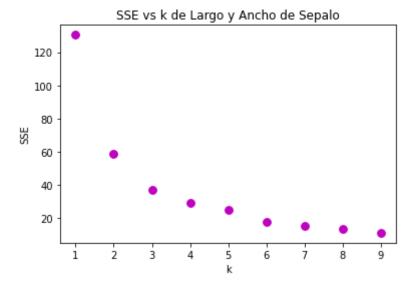


## Selecting K: elbow plot

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

```
In [42]: # Intialize a list to hold sum of squared error (sse)
    sse = []
    # Define values of k
    kNew = range(1,10)
    # For each k
    for i in kNew:
        km = KMeans(n_clusters = i, n_init = "auto")
        km.fit_predict(df[["Largo_Sepalo", "Ancho_Sepalo"]])
        sse.append(km.inertia_)
```

```
In [43]: # Plot sse versus k
plt.scatter(kNew, sse, c = "m", s = 60)
plt.title("SSE vs k de Largo y Ancho de Sepalo")
plt.xlabel("k")
plt.ylabel("SSE")
plt.show()
```



Choose the k after which the sse is minimally reduced. Based on this analysis, it seem that the best k is 3.

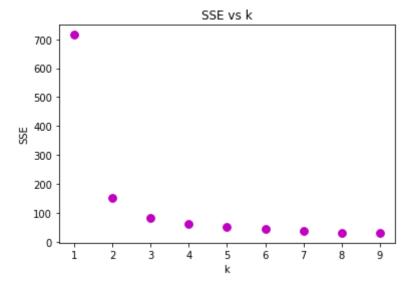
#### \*\*Important remarks\*\*

• Observations? The use of scatter plot is helpful in ML analysis and to choose the best k for the clusters.

## **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

```
In [44]:
         # Repeat the analysis using all six features
         sse = []
         # Define values of k
         kNew = range(1,10)
         # For each k
         for i in kNew:
           km = KMeans(n_clusters = i, n_init = "auto")
           km.fit predict(df)
           sse.append(km.inertia_)
           # Plot sse versus k
         plt.scatter(kNew, sse, c = "m", s = 60)
         plt.title("SSE vs k")
         plt.xlabel("k")
         plt.ylabel("SSE")
         plt.show()
```



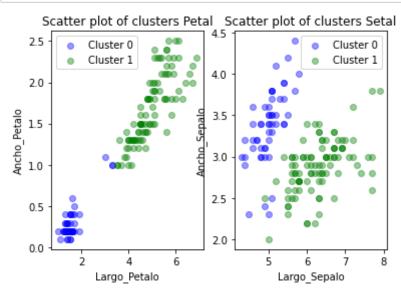
#### **Conclusion**

As seen above, when the SSE is obtained considering all features, the best k is 2. However, when looking at this analysis only from the perspective of two individual features, this answer changes, as seen previously.

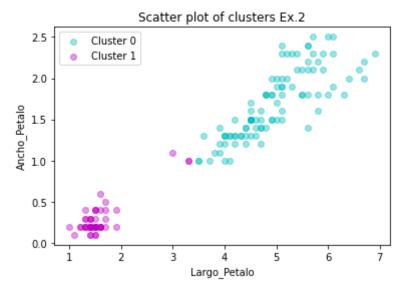
#### Activity: work with the iris dataset

- 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\*\*

```
In [124]:
          # 1.
          # Making the clusters
          df.drop("Cluster_Information", axis = 1)
          K =
          km = KMeans(n clusters = K, n init = "auto")
          yestimated = km.fit predict(df)
          df.insert(4, "Cluster", yestimated, True)
          # Scatter plot of each cluster on Petal
          plt.subplot(1,2,1)
          df1 = df[df.Cluster == 0]
          df2 = df[df.Cluster == 1]
          plt.scatter(df1.Largo Petalo, df1.Ancho Petalo, label = "Cluster 0", c =
          'b', marker = 'o', alpha = 0.4)
          plt.scatter(df2.Largo Petalo, df2.Ancho Petalo, label = "Cluster 1", c =
          'g', marker = 'o', alpha = 0.4)
          plt.title("Scatter plot of clusters Petal")
          plt.xlabel("Largo Petalo")
          plt.ylabel("Ancho_Petalo")
          plt.legend()
          # Scatter plot of each cluster on Setal
          plt.subplot(1,2,2)
          df1 = df[df.Cluster == 0]
          df2 = df[df.Cluster == 1]
          plt.scatter(df1.Largo Sepalo, df1.Ancho Sepalo, label = "Cluster 0", c =
          'b', marker = 'o', alpha = 0.4)
          plt.scatter(df2.Largo Sepalo, df2.Ancho Sepalo, label = "Cluster 1", c =
          'g', marker = 'o', alpha = 0.4)
          plt.title("Scatter plot of clusters Setal")
          plt.xlabel("Largo Sepalo")
          plt.ylabel("Ancho Sepalo")
          plt.legend()
          plt.show()
```



```
In [126]:
          # 2.
          df.drop("Cluster", axis = 1)
          df20 = df.drop(["Largo_Sepalo", "Ancho_Sepalo"], axis = 1)
          # Making the clusters
          K = 2
          km = KMeans(n_clusters = K, n_init = "auto")
          yestimated = km.fit_predict(df20)
          df20.insert(2, "Cluster Information", yestimated, True)
          # Scatter plot of each cluster
          df1 = df20[df20.Cluster_Information == 0]
          df2 = df20[df20.Cluster Information == 1]
          plt.scatter(df1.Largo_Petalo, df1.Ancho_Petalo, label = "Cluster 0", c =
          'c', marker = 'o', alpha = 0.4)
          plt.scatter(df2.Largo Petalo, df2.Ancho Petalo, label = "Cluster 1", c =
          'm', marker = 'o', alpha = 0.4)
          plt.title("Scatter plot of clusters Ex.2")
          plt.xlabel("Largo Petalo")
          plt.ylabel("Ancho_Petalo")
          plt.legend()
          plt.show()
```



```
In [128]:
          # 3.
          df.drop("Cluster", axis = 1)
          df30 = df.drop(["Largo_Petalo", "Ancho_Petalo"], axis = 1)
          # Making the clusters
          K = 2
          km = KMeans(n clusters = K, n init = "auto")
          yestimated = km.fit_predict(df30)
          df30.insert(2, "Cluster Information", yestimated, True)
          # Scatter plot of each cluster
          df1 = df30[df30.Cluster_Information == 0]
          df2 = df30[df30.Cluster Information == 1]
          plt.scatter(df1.Largo_Sepalo, df1.Ancho_Sepalo, label = "Cluster 0", c =
          'b', marker = 'o', alpha = 0.4)
          plt.scatter(df2.Largo Sepalo, df2.Ancho Sepalo, label = "Cluster 1", c =
           'y', marker = 'o', alpha = 0.4)
          plt.title("Scatter plot of clusters Ex.3")
          plt.xlabel("Largo Sepalo")
          plt.ylabel("Ancho_Sepalo")
          plt.legend()
          plt.show()
```

