k-means-HVoe

August 27, 2023

0.0.1 Test a DHBW Seminarwork in W2022

Seminar task DW13 for DWH lecture in WS2022

```
[1]: import pandas as pd # we need the dataframe structure from pandas
     import numpy as np # import numpy for random, mean, sqrt function
     from sklearn import datasets # import data
     import matplotlib.pyplot as plt # import for plotting the data
     import matplotlib.patches as mpatches # import for plotting the data
     import copy # import function to get a deep copy of important values (otherwise,
     →values could be overwritten)
     %matplotlib inline
     iris = datasets.load_iris() # load iris data set
     # load array into pandas dataframe, name columns according to what value they
     \rightarrowrepresent
     initial_iris_data = pd.DataFrame(iris.data, columns=['Sepal Length', 'Sepal_
      →Width', 'Petal Width', 'Petal Length']) # import the values from the iris,
      →dataset and name the columns according to the values they represent
     print(initial iris data)
     # load which data set is connected to which species of iris
     target = pd.DataFrame(iris.target, columns=['Target']) # load (scientific)_
      → labeling of iris data
     #check versions of libraries
     print('pandas version is: {}'.format(pd.__version__))
     print('numpy version is: {}'.format(np.__version__))
     import sklearn
     print('sklearn version is: {}'.format(sklearn.__version__))
     # Import libary time to check execution date+time
     import time
```

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

```
[150 rows x 4 columns]
pandas version is: 1.0.1
numpy version is: 1.18.1
sklearn version is: 0.22.1
```

Actual date & time of the notebook: 26.08.2023 20:04:40

```
[2]: # Initialization: create 3 centroids (for 3 clusters)
     def createRandomCentroids(k, seed):
         np.random.seed(seed = seed)
         centroids = {
             i+1: [
                  initial_iris_data['Sepal Length'].min() + np.random.random() *__
      →(initial_iris_data['Sepal Length'].max() - initial_iris_data['Sepal Length'].
      \rightarrowmin()),
                  initial_iris_data['Sepal Width'].min() + np.random.random() *_
      →(initial_iris_data['Sepal Width'].max() - initial_iris_data['Sepal Width'].
      \rightarrowmin()),
                  initial_iris_data['Petal Width'].min() + np.random.random() *__
      →(initial_iris_data['Petal Width'].max() - initial_iris_data['Petal Width'].
      \rightarrowmin()),
                  initial_iris_data['Petal Length'].min() + np.random.random() *__
      →(initial_iris_data['Petal Length'].max() - initial_iris_data['Petal Length'].
      \rightarrowmin())
```

```
for i in range(k) # create an array of length k with index 1 to index k_{\square} \rightarrow entries which represent the cluster centroids. The entries themself are arrays \rightarrow (in this case of length 4) } return centroids
```

```
[3]: # draw initial cluster centers on two graphs
     def drawInitialCentroids(data, centroids):
         plt.figure(figsize=(12, 3))
         colors = np.array(['red', 'green', 'blue'])
         # plot the datavalues sepal length in comparison to their sepal width
         plt.subplot(1, 2, 1)
         plt.scatter(data['Sepal Length'], data['Sepal Width'], c="black")
         for i in centroids.keys():
             plt.scatter(centroids[i][0], centroids[i][1], s=100, color=colors[i - 1])
         plt.title('Sepal Length vs Sepal Width')
         # plot the datavalues petal length in comparison to their petal width
         plt.subplot(1, 2, 2)
         plt.scatter(data['Petal Length'], data['Petal Width'], c="black")
         for i in centroids.keys():
             plt.scatter(centroids[i][3], centroids[i][2], s=100, color=colors[i - 1])
         plt.title('Petal Length vs Petal Width')
```

```
[5]: # draws the current state of the clustering. With the clustercenters beeing ⊔

⇒alpha 1 and the points alpha 0.5

def drawFigure(data, centroids):

plt.figure(figsize=(12,3))
```

```
colors = np.array(['red', 'green', 'blue'])
        iris_targets_legend = np.array(iris.target_names)
        red_patch = mpatches.Patch(color='red', label='Setosa')
        green_patch = mpatches.Patch(color='green', label='Versicolor')
        blue_patch = mpatches.Patch(color='blue', label='Virginica')
        plt.subplot(1, 2, 1)
        plt.scatter(data['Sepal Length'], data['Sepal Width'],
      for i in centroids.keys():
            plt.scatter(centroids[i][0], centroids[i][1], s = 100, color=colors[i -u
      →1])
        plt.title('Sepal Length vs Sepal Width')
        plt.legend(handles=[red_patch, green_patch, blue_patch])
        plt.subplot(1,2,2)
        plt.scatter(data['Petal Length'], data['Petal Width'],

→c=colors[data['cluster'] - 1], alpha = 0.5)
        for i in centroids.keys():
            plt.scatter(centroids[i][3], centroids[i][2], s= 100, color=colors[i -u
      →1])
        plt.title('Petal Length vs Petal Width')
        plt.legend(handles=[red_patch, green_patch, blue_patch])
[6]: # calc new centroids using weight (using the mean function of each value, that
     \hookrightarrow is assigned to the cluster)
     def calcNewCentroids(df, centroids):
        for i in centroids.keys():
            centroids[i][0] = np.mean(df[df['cluster'] == i]['Sepal Length'])
            centroids[i][1] = np.mean(df[df['cluster'] == i]['Sepal Width'])
            centroids[i][2] = np.mean(df[df['cluster'] == i]['Petal Width'])
            centroids[i][3] = np.mean(df[df['cluster'] == i]['Petal Length'])
[7]: # compare the prediction of the clustering against the actual values and return
     →a success percentage
     def calcAccuracy(prediction, target):
        correct = 0
        for i in prediction.index:
            if prediction[i] == target[i] + 1:
                correct += 1
        return correct / prediction.count()
[8]: # iterate until stable (until centroids do not change)
     iterationCount = 0
     def kMeans(data, numberOfClusters):
        bestAccuracy = 0
```

```
bestPrediction = None
    bestCentroids = None
    for i in range(50):
        centroids = createRandomCentroids(numberOfClusters, i)
        while True:
            oldCentroids = copy.deepcopy(centroids) # copy old centroids
 → (otherwise they would be overwritten)
            assignPointsToCluster(data, centroids)
            calcNewCentroids(data, centroids)
             # if the centroids have not changed in the current run of k-means
 \hookrightarrow that means that clustering is at the current best and will not change any \sqcup
 \rightarrow further
            if centroids == oldCentroids:
        accuracy = calcAccuracy(data['cluster'], target['Target']) # calculate_
 →accuracy of current run
        if accuracy > bestAccuracy:
            print(f"{i}: {accuracy}")
             # copy all the important values from the currently most sucessfull \sqcup
 \rightarrowrun for returning it at the end of the function
            bestPrediction = data.copy(deep = True)
            bestCentroids = copy.deepcopy(centroids)
            bestAccuracy = accuracy
    drawFigure(bestPrediction, bestCentroids) # draw the best clustering we can_\sqcup
 \rightarrow achive by using k-means
    return bestPrediction
bestPrediction = kMeans(initial_iris_data, 3) # run k means with iris data, with_
 →3 clusters and save output of the function to bestPrediction
```

0: 0.24 1: 0.58



