Week 11 Quiz

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Due Tues Nov 22nd, 11:59pm

The MNIST digits dataset is a dataset commonly used to demonstrate image recognition.

The dataset is composed of a set of images of handwritten digits from 0 to 9. There are 1797 images, each 8x8 pixels. If we flatten each image we get a dataset of 1797 observations, each with 64 features, each belonging to one of 10 classes.

In this quiz we'll see how well these images cluster using KMeans clustering. We'll compare the KMeans clustering assingments to clusters generated by HAC with Ward linkage.

Instructions

Replace the Name and UNI in cell above and the notebook filename

Replace all '_' below using the instructions provided.

When completed,

- 1. make sure you've replaced Name and UNI in the first cell and filename
- 2. Kernel -> Restart & Run All to run all cells in order
- 3. Print Preview -> Print (Landscape Layout) -> Save to pdf
- 4. post pdf to GradeScope

Setup Environment

```
In [1]: import numpy as np
   import pandas as pd
  import seaborn as sns
```

```
import matplotlib.pyplot as plt
%matplotlib inline
```

Load the Dataset

```
In [2]: # From sklearn datasets import load_digits.

# Load the dataset into 'digits' using load_digits

digits = load_digits()

# Extract digits['data'] to X_digits. No need to reshape as each image has already been flattened to 1x64

X_digits = digits['data']

# Extract the labels in digits['target'] to y_digits

y_digits = digits['target']

# Assert that the shape of X_digits is 1797 rows, 64 columns

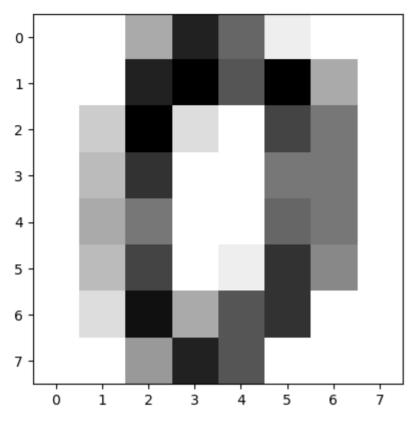
assert X_digits.shape == (1797,64)

In [3]: # We can use plt.imshow() to display one of the images as an example.

# 'digits['images']' is a list of images of size 8x8 pixels.

# We can plot the first image using plt.imshow with cmap=plt.cm.gray_r

# You should see a black '0' on a white background.
plt.imshow(digits['images'][0], cmap=plt.cm.gray_r);
```



```
In [4]: # First we'll reduce our dataset from 64 to 2 dimensions using PCA for plotting

# Import PCA from sklearn
from sklearn.decomposition import PCA

# Instantiate a pca object that will result in 2 components being returned.

# Use n_components=2, random_state=512

# Store as pca
pca = PCA(n_components=2, random_state=512)

# Transform X_digits to 2D using fit_transform.

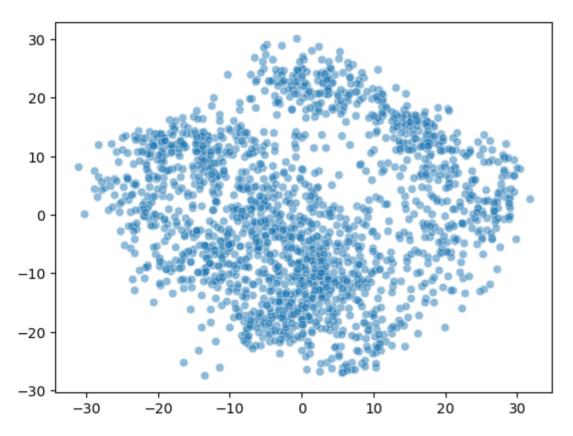
# Store as X_2D
X_2D = pca.fit_transform(X_digits)

# assert that the dataset has been reduced to 2 dimensions (2 columns)
assert X_2D.shape[1] == 2
```

```
In [5]: # Create a scatterplot of X_2D

# Use sns.scatterplot to plot X_2D with
# the first column of X_2D on the x-axis
# the second column of X_2D on the y-axis
# alpha=.5 to make points slightly transparent
sns.scatterplot(x = X_2D[:,0], y = X_2D[:,1], alpha=.5)
```

Out[5]: <AxesSubplot: >



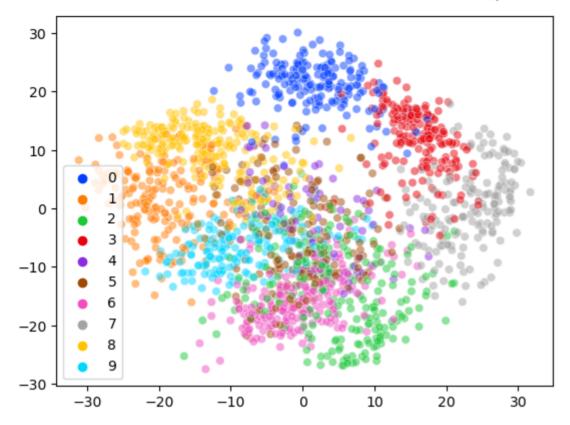
```
In [6]: # Cluster the full dataset X_digits using KMeans

# Import KMeans from sklearn
from sklearn.cluster import KMeans

# Intantiate a KMeans object
# which will generate 10 clusters
# use init='k-means++'
```

```
and random state=512
          all other parameters as default (including init='k-means++')
        # Store as km
        km = KMeans(n clusters = 10, init='k-means++', random state=512)
        # Use .fit predict() on X digits to both fit our k-means model and generate cluster assignments.
        # Store the result as cluster assignments km
        cluster assignments km = km.fit predict(X digits)
        # print the first 15 cluster assignments in cluster assignments km
        # They should all be integers between 0 and 9 inclusive
        cluster assignments km[:15]
        array([0, 2, 2, 1, 7, 8, 3, 6, 2, 8, 0, 4, 9, 1, 7], dtype=int32)
Out[6]:
In [7]: # Plot X 2D as a scatterplot colored by their cluster assignments
        # Use sns.scatterplot to plot X 2D again with
        # the first column of X 2D on the x-axis
        # the second column of X 2D on the y-axis
        # alpha=.5 to make points slightly transparent
        # hue=cluster assignments km to color each point by it's cluster assignment
        # palette='bright' to use a brighter color palette than the seaborn default
        sns.scatterplot(x = X 2D[:,0], y = X 2D[:,1], alpha=.5, hue=cluster assignments km, palette='bright')
        # Note that the legend shows cluster assignments, not digit label
        # Also note that, since we are clustering in 64 dimensional space, the clusters will
        # appear to overlap in 2 dimensional space
```

Out[7]: <AxesSubplot: >



```
In [8]: # What are the labels for images placed in cluster 9?

# Since we actually have labels for these images we can see which
# digits were placed in cluster 9 without looking at the images themselves.

# Use y_digits and cluster_assignments_km to print the labels for
# images assigned to cluster 9
# You should see that most of the labels are digit 2 with some 1s and 8s
y_digits[cluster_assignments_km == 9]
```

```
Out[8]:
            2, 1, 2, 1, 2, 3, 1, 2, 2, 2, 1, 2, 2, 2, 1, 2, 1, 1, 1, 1, 1, 2, 2,
            2, 1, 2, 1, 2, 1, 2, 1, 2, 2, 1, 2, 1, 2, 2, 1, 2, 1, 1, 1, 1, 1, 1,
            2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 8, 2, 2, 2, 2, 2, 2, 2, 2, 2,
            In [9]: # Evaluate clustering using Homogeneity
       # sklearn has a metric homogeneity score which measures the homogeneity of clusters
       # It returns a value between 0 and 1, where a higher value means more homogeneous.
       # Compare the clustering assignments of Kmeans found above with an HAC model using Ward linkage
       # Refer to the documentation here:
       # https://scikit-learn.org/stable/modules/generated/sklearn.metrics.homogeneity score.html
       # import homogeneity score from sklearn.metrics
       from sklearn.metrics import homogeneity score
       # import AgglomerativeClustering from sklearn.cluster
       from sklearn.cluster import AgglomerativeClustering
       # generate cluster assignments of X digits with AgglomerativeClustering
       # using n clusters=10
       # and linkage='ward'
       # all other parameters as default
       # store as cluster assignments hac
       cluster assignments hac = AgglomerativeClustering(n clusters=10, linkage='ward').fit predict(X digits)
       #Print the homogeneity scores for the clustering assigned by KMeans
       print(f'homogeneity score for KMeans: {homogeneity score(y digits,cluster assignments km):0.2f}')
       #Print the homeogeneity score for the clustering assinged by Agglomerative Clustering
       print(f'homogeneity score for HAC : {homogeneity score(y digits,cluster assignments hac):0.2f}')
       homogeneity score for KMeans: 0.74
       homogeneity score for HAC : 0.86
In [ ]:
```