- First rename this file by replacing "ID" with your "last_name_actual_id"
- HWF-Python. Use this template to implement the following tasks:

Tasks:

1. Read image data (ilk-3b-1024.tif) at the locations provided in the file (ilk-3b-1024-10k-pnts.csv)

ilk-3b-1024.tif contains 3 spectral bands, Width = 1024, Height = 1024

####

Generating training data: For this, use the sample location data from file ilk-3b-1024-10k-pnts.csv. This file contains 10k points, with format (ID, X, Y). Now, for each X, Y coordinate read pixel data (a vector of length 12) as a pandas dataframe.

removed attribute selection task, cheer up!

- 2. For initial clustering you can use KMeans() function from sklearn library. To get yourself familiar, do clustering with K=10 and plot cluster data.
- 3. Implement the "Algorithm 1" in the paper "Learning the k in k-means"
- 4. Generate visualizations (plots) as asked in the cells.

Read each cell and write code as needed

Organize your clode logically, create new cells as needed

You are allowed to search for code snippet on the web for plotting, sklearn, etc., but code should be your own. You are not allowed to directly search for actual implementaion code of the the paper and use code for the same that may be freely available on the web and/or github repositories.

I have already removed attribue selection component to reduce your implementation time, so don't take any shortcuts! Enjoy doing your first data science project.

```
! pip install rasterio
     Requirement already satisfied: rasterio in /usr/local/lib/python3.10/dist-packages (1.3.10)
     Requirement already satisfied: affine in /usr/local/lib/python3.10/dist-packages (from rasterio) (2.4.0)
     Requirement already satisfied: attrs in /usr/local/lib/python3.10/dist-packages (from rasterio) (23.2.0)
     Requirement already satisfied: certifi in /usr/local/lib/python3.10/dist-packages (from rasterio) (2024.2.2)
     Requirement already satisfied: click>=4.0 in /usr/local/lib/python3.10/dist-packages (from rasterio) (8.1.7)
     Requirement already satisfied: cligj>=0.5 in /usr/local/lib/python3.10/dist-packages (from rasterio) (0.7.2)
     Requirement already satisfied: numpy in /usr/local/lib/python3.10/dist-packages (from rasterio) (1.25.2)
     Requirement already satisfied: snuggs>=1.4.1 in /usr/local/lib/python3.10/dist-packages (from rasterio) (1.4.7)
     Requirement already satisfied: click-plugins in /usr/local/lib/python3.10/dist-packages (from rasterio) (1.1.1)
     Requirement already satisfied: setuptools in /usr/local/lib/python3.10/dist-packages (from rasterio) (67.7.2)
     Requirement already satisfied: pyparsing>=2.1.6 in /usr/local/lib/python3.10/dist-packages (from snuggs>=1.4.1->rasterio) (3.1.2)
# import necessary libraries; don't change this cell, if you need any additional libraries, use next cell
import numpy as np
import rasterio as rio
from rasterio.plot import show
from rasterio.plot import show_hist
from rasterio.windows import Window
from\ rasterio.windows\ import\ from\_bounds
import pandas as pd
import random
import matplotlib.pyplot as plt
import seaborn as sns
import sklearn
from sklearn.cluster import KMeans
import sys
np.random.seed(100)
# if you need additional libraries, include them here
# make to include all your additional liberies here and not any other cells
from sklearn.decomposition import PCA
from scipy.stats import norm, anderson
from sklearn.preprocessing import StandardScaler
```

Step 1: Prepare data (Generate training data)

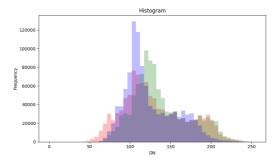
```
# Read image: data/ilk-3b-1024.tif
# Hint: explore rasterio package
img_path = "ilk-3b-1024.tif"
rgb_img = rio.open(img_path)
```

once successfully opened image, you can get meta data such as size, projection, etc.

read rasterio documentation to learn more about the libarary and its functionality

```
print(rgb_img.profile)
     {'driver': 'GTiff', 'dtype': 'uint8', 'nodata': None, 'width': 1024, 'height': 1024, 'count': 3, 'crs': CRS.from_epsg(26916), 'trans
0.0, -1.0, 4620006.0), 'blockysize': 2, 'tiled': False, 'interleave': 'pixel'}
rgb img.shape
     (1024, 1024)
# visualize image data and its histogram (write your code here to generate figure similar to the one below
# Hint: you can explore matplotlib subplots, raterio.plot, etc
img_arr = rgb_img.read()
#Splitting image into rgb channels
red = img_arr[0]
green = img arr[1]
blue = img_arr[2]
fig, axes = plt.subplots(1,2,figsize=(15,5))
#Plotting the Image
show(rgb_img, ax=axes[0])
axes[0].set_title('Image')
#Plotting histograms for R,G,B values
axes[1].hist(red.flatten(), bins=50, range=[0,255], color='red', alpha=0.25)
axes[1].hist(green.flatten(), bins=50, range=[0,255], color='green', alpha=0.25)
axes[1].hist(blue.flatten(), bins=50, range=[0,255], color='blue', alpha=0.25)
axes[1].set_title("Histogram")
axes[1].set_xlabel("DN")
axes[1].set_ylabel("Frequency")
plt.tight_layout()
plt.show()
```





Likewise, open csv file for reading sample locations

```
# read csv file containing sample locations: data/ilk-3b-1024-10k-pnts.csv
# hint, you can use pandas functions csv_path = "ilk-3b-1024-10k-pnts.csv"
xydata = pd.read_csv(csv_path)
```

once successfuly read csv file, examine the data to make sure everything is fine ## display top 5 rows here xydata.head()

	id	Х	Υ	\blacksquare
0	0	395141.5756	4619891.465	11.
1	1	395267.6855	4619541.988	
2	2	395177.9343	4619966.368	
3	3	395444.7146	4619615.272	
4	4	395236.0146	4619433.194	

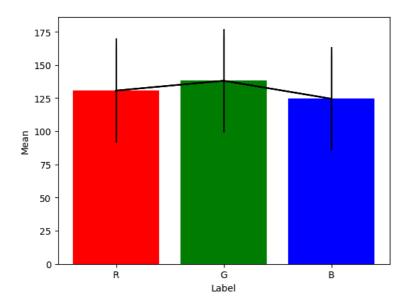
print(xydata.head())

	id	X	Υ
0	0	395141.5756	4619891.465
1	1	395267.6855	4619541.988
2	2	395177.9343	4619966.368
3	3	395444.7146	4619615.272
4	4	395236.0146	4619433.194

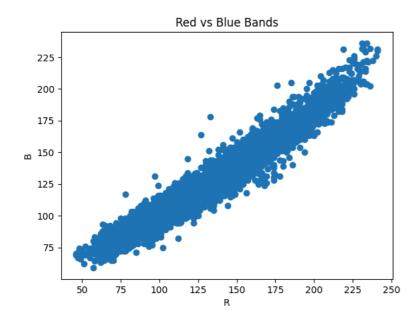
xydata.describe()

	id	х	Υ	
count	10000.00000	10000.000000	1.000000e+04	11.
mean	4999.50000	395042.964026	4.619498e+06	
std	2886.89568	291.171868	2.927231e+02	
min	0.00000	394536.668900	4.618991e+06	
25%	2499.75000	394789.177300	4.619241e+06	
50%	4999.50000	395041.552000	4.619502e+06	
75%	7499.25000	395292.699275	4.619753e+06	
max	9999.00000	395548.958000	4.620000e+06	

```
len(xydata)
     10000
\# now you can read data from raster image for every x,y coordinate from the csv file
# Hint: explore rasterio image reading functions
# The code you write here should finally produce a panadas dataframe, say df
points = []
with rio.open(img_path) as image:
 for i in range(len(xydata)):
   row, col = image.index(xydata['X'][i], xydata['Y'][i])
    rgb = image.read([1,2,3], window=((row, row+1), (col, col+1)))
   points.append(rgb.flatten())
cols = ['R', 'G', 'B']
# convert the data you read (say into variable d) to a dataframe
df = pd.DataFrame(points, columns=cols)
# First name the columns in the dataframe as R, G, B.
# Then display image data (r, g, b) for first 5 points
df.head()
          R
               G
                    В
                        \blacksquare
     0
        74
              86
                   86
     1 106 120 105
     2 155 163 154
     3
        76
              92
                  86
     4 209 213 187
 # compute mean and standard deviation of each band,
# generate bargraphs, and show standard deviation on the bar plot
m = df.mean()
sd = df.std()
print("Means:\n", m)
print("SD:\n", sd)
     Means:
         130.6914
         138.0920
         124.5246
     dtype: float64
     SD:
          39.133841
     R
         34.703386
     G
         31.192081
     В
     dtype: float64
# write code to plot the means and standard deviations here
plt.bar(['R', 'G', 'B'], m, color=['red', 'green', 'blue'])
for i, mean in enumerate(m):
 plt.errorbar(['R', 'G', 'B'], m, yerr=sd[i], color='black', capsize=0)
plt.xlabel('Label')
plt.ylabel('Mean')
plt.show()
```



```
# visualize rgb data as scatter plot; select bands 1 and 3
# note image attributes are called bands (e.g., red, green, blue bands)
plt.scatter(df['R'], df['B'])
plt.title('Red vs Blue Bands')
plt.xlabel('R')
plt.ylabel('B')
plt.show()
```



Done with Step 1

Step 2: To get yourself familar with KMeans clustering, play with KMeans() clustering from scikit package

```
# k-means clustering using KMeans() from scikit-learn with K = 10

K = 10

# use km0 to store mode
km0 = KMeans(n_clusters=K)

# use y_km0 for storing predicted lables
y_km0 = km0.fit_predict(df)
```

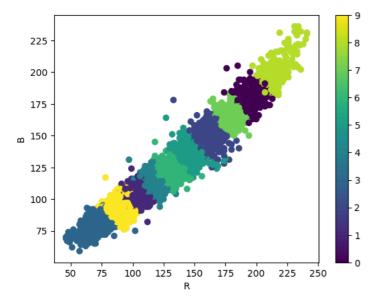
```
# write code to print number of samples in each cluster

clusterId, numPoints = np.unique(y_km0, return_counts=True)
# clusterId, numPoints
pointsPerCluster = []
for i in range(len(clusterId)):
    ele = [clusterId[i], numPoints[i]]
    pointsPerCluster.append(ele)

print(pointsPerCluster)

    [[0, 802], [1, 1595], [2, 842], [3, 768], [4, 1688], [5, 903], [6, 1207], [7, 816], [8, 288], [9, 1091]]
# visualize clusters as scatter plot

plt.scatter(df['R'], df['B'], c=y_km0)
plt.colorbar()
plt.xlabel('R')
plt.ylabel('B')
plt.show()
```



generate clustered image and display by writing code in the following cells

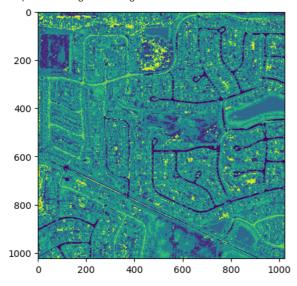
```
# read full image data
img = rgb_img.read()

# display image to check if you read the data correctly
show(img, cmap='viridis')
```

```
0
       200
       400
       600
       800
      1000
                    200
                              400
                                                 800
            0
                                       600
                                                          1000
     <Axes: >
# note that the img data you read is ndarray, verify it
type(img)
     numpy.ndarray
# and its shape should be (3, 1024, 1024)
print(img.shape)
     (3, 1024, 1024)
# However, the clustering model prediction expects that the input to be r,g,b vectors
# So write code to convert image into rgb vectors
rgb_vec = img.transpose(1,2,0).reshape(-1,3)
# if you converted it correctly, you should get the shape of vector as (1048576, 3)
rgb_vec.shape
     (1048576, 3)
\mbox{\tt\#} its good to verify that format of r, g, b, and actual values
# for example, first image pixel value should be (124, 129, 112)
rgb_vec[0]
     array([124, 129, 112], dtype=uint8)
# predict label for each pixel (vector) by calling the predict method of KMeans() clustering
imgkm = km0.predict(rgb_vec)
     /usr/local/lib/python3.10/dist-packages/sklearn/base.py:439: UserWarning: X does not have valid feature names, but KMeans was fitted
       warnings.warn(
# Check the shape of the output, should be same as 1024 * 1024 = 1048576
imgkm.shape
     (1048576,)
# you need to convert the is 1-D r,g,b vector back to image of 1024,1024 (note that its a single band image)
imgkm_out = imgkm.reshape(1024, 1024)
# check the shape
imgkm_out.shape
     (1024, 1024)
type(imgkm_out)
     numpy.ndarray
```

```
# plot the clustered image
plt.imshow(imgkm_out)
```

<matplotlib.image.AxesImage at 0x7eb7aaf9be50>

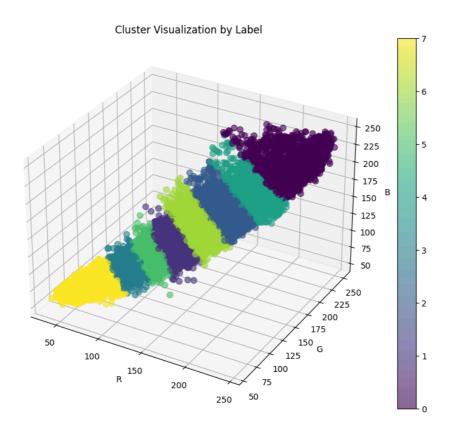


End of Step 2

Step 3: Implement the "Algorithm 1" in the paper "Learning the k in k-means"

```
# implemention code for Algorithm 1
\ensuremath{\text{\#}} Read the algorithm and the description in particular section 2.1
# You are allowed to use any Python library statstical functions like mean, standard deviation, variance, etc.
# create extra cells as needed, don't forget to comment each new cell you create
def check_for_normality(df, alpha=0.0001):
  res = anderson(df)
  n = len(df)
  stephens_correction = (1 + 4/n - 25/(n*n))
  res_corrected = res.statistic * stephens_correction
  res_alpha = res.significance_level/100.0
  cvs = res.critical_values
  nearest_ind = np.argmin(np.abs(res_alpha-alpha))
  nearest_alpha = res_alpha[nearest_ind]
  nearest_cv = cvs[nearest_ind]
  return res_corrected < nearest_cv
def child_centers_by_pca(cur_center, df):
  pca = PCA(n_components=1)
  pca.fit(df)
  s = pca.components_[0]
  eigenvalue = pca.explained_variance_[0]
  m = s*np.sqrt((2*eigenvalue)/np.pi)
  c1 = cur_center + m
  c2 = cur center - m
  return c1, c2
```

```
# your algorith 1 clode goes here ...
def gmeans(df, alpha=0.0001, max_iters=250):
  init_centers = [np.mean(df, axis=0)]
  for i in range(max_iters):
    updated_centers = []
    for center in init_centers:
     c1, c2 = child_centers_by_pca(center, df)
      v = c1-c2
     projection = np.dot(df-np.mean(df, axis=0), v) / np.linalg.norm(v)
      scaler = StandardScaler()
     projection = scaler.fit_transform(projection.reshape(-1,1)).flatten()
      if check_for_normality(projection):
       updated_centers.append(center)
       km = KMeans(n_clusters=2, init=np.array([c1, c2])).fit(df)
       updated_centers.extend(km.cluster_centers_)
    if set(map(tuple, init_centers)) == set(map(tuple, updated_centers)):
    init_centers = updated_centers
  return KMeans(n_clusters=len(init_centers), init=init_centers).fit(df)
opt_clusters = gmeans(rgb_vec)
# print optimal number of clusters found with the algorithm
print("Optimal number of clusters is:", opt_clusters.n_clusters)
     Optimal number of clusters is: 8
# print number of samples in each cluster
clusterId, numPoints = np.unique(opt_clusters.predict(rgb_vec), return_counts=True)
pointsPerCluster = []
for i in range(len(clusterId)):
 ele = [clusterId[i], numPoints[i]]
  pointsPerCluster.append(ele)
print("Number of points per cluster:", pointsPerCluster)
     Number of points per cluster: [[0, 51292], [1, 161131], [2, 95003], [3, 180153], [4, 110230], [5, 235938], [6, 107646], [7, 107183]
# generate scatter plot and color using cluster labels
figure = plt.figure(figsize=(10, 8))
ax = figure.add_subplot(111, projection='3d')
scatter_plot = ax.scatter(rgb_vec[:, 0], rgb_vec[:, 1], rgb_vec[:, 2], c=opt_clusters.labels_, s=50, alpha=0.6)
ax.set_title('Cluster Visualization by Label')
ax.set_xlabel('R')
ax.set_ylabel('G')
ax.set_zlabel('B')
figure.colorbar(scatter_plot, ax=ax)
plt.show()
```



 $\mbox{\#}$ Apply the final model to predict labels for each pixel in the image, $\mbox{\#}$ that is, generate a clustered image

preds = opt_clusters.predict(rgb_vec)
image_clusters = preds.reshape(1024, 1024)

display the clustered image here.
plt.imshow(image_clusters)
plt.show()

