KMeans Clustering

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Lake Picture

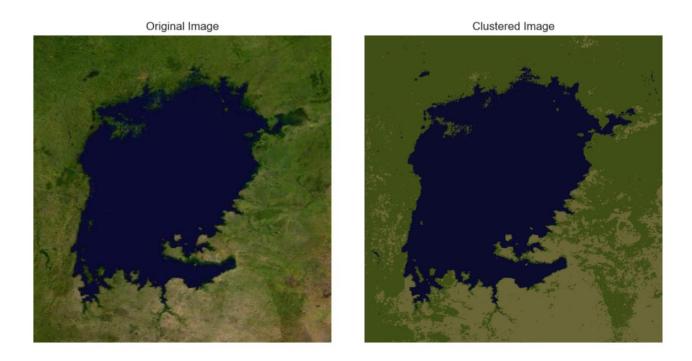


Figure 1. Original and Clustered Image with 3 clusters

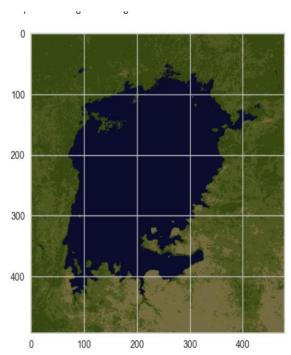


Figure 2. Clustered Image (K=3)

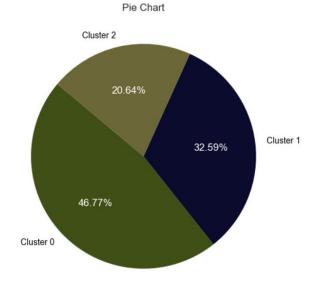


Figure 3. Distribution of Clusters (K=3)

	Red	Green	Blue	Total_counts
0	64	79	22	110905
1	11	12	45	77278
2	107	103	56	48937

Figure 4. RGB Counts of each cluster (K=3)

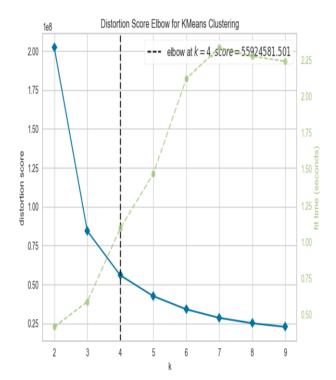


Figure 5. Find optimal K using Elbow method

```
For n_clusters = 2 The average silhouette_score is : 0.6895546114519658
For n_clusters = 3 The average silhouette_score is : 0.6463363467170485
For n_clusters = 4 The average silhouette_score is : 0.5728389709569488
For n_clusters = 5 The average silhouette_score is : 0.5556914900234113
For n_clusters = 6 The average silhouette_score is : 0.535214860837284
```

Figure 6. Silhouette score for different K values

Silhouette analysis for KMeans clustering on sample data with n_clusters = 2

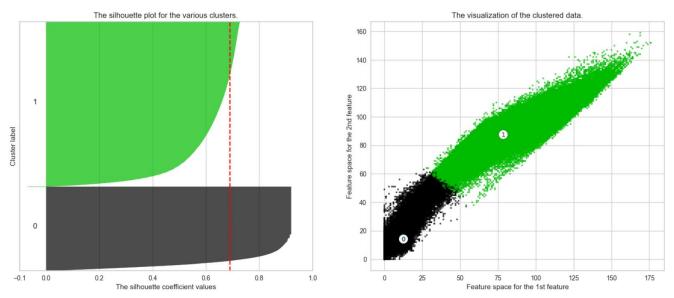


Figure 7. Silhouette analysis with K=2

Silhouette analysis for KMeans clustering on sample data with n_clusters = 3

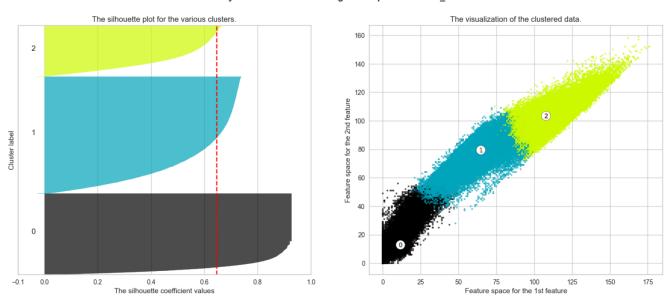


Figure 8. Silhouette analysis with K=3

Silhouette analysis for KMeans clustering on sample data with n_clusters = 4

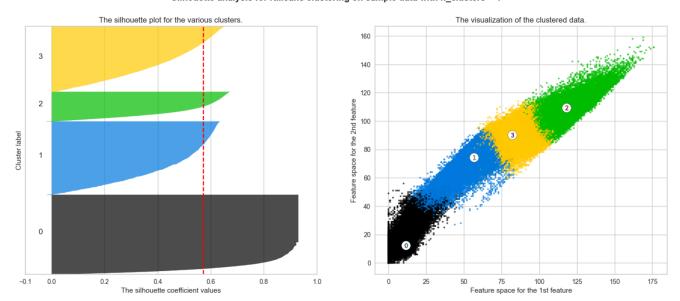


Figure 9. Silhouette analysis with K=4

Silhouette analysis for KMeans clustering on sample data with n_clusters = 5

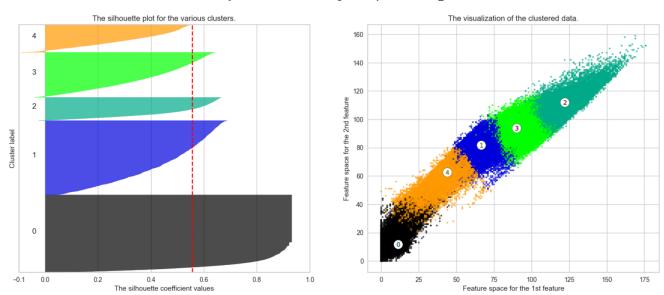


Figure 10. Silhouette analysis with K=5

Silhouette analysis for KMeans clustering on sample data with n_clusters = 6

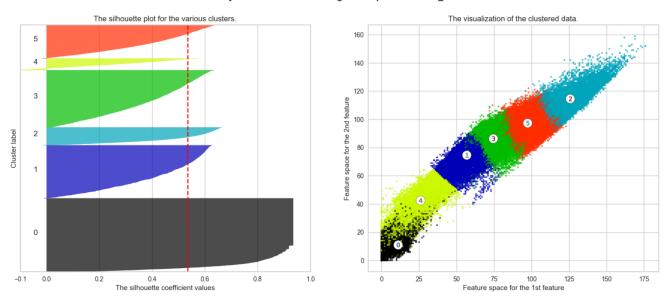


Figure 11. Silhouette analysis with K=6

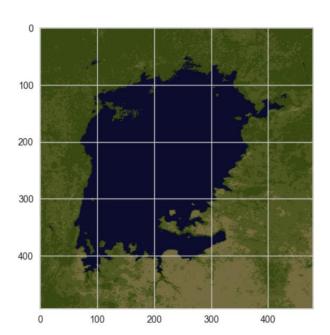


Figure 12. Clustered picture with 4 clusters

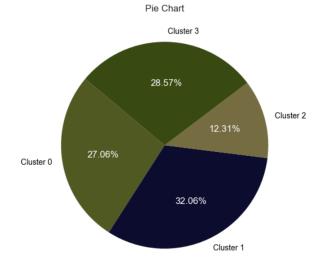


Figure 13. Cluster distribution (K=4)

	Red	Green	Blue	Total_counts
0	81	89	35	64162
1	11	12	46	76025
2	117	109	65	29197
3	56	73	17	67736

Figure 14. RBG counts of all 4 clusters

B. Granite Picture



Figure 15. Original Picture

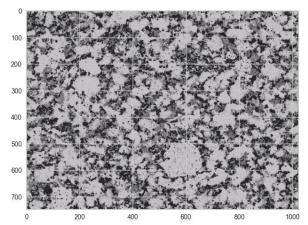


Figure 16. Clustered Image

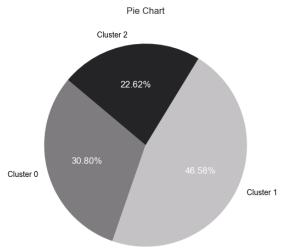


Figure 17. Distribution of Clusters (K=3)

	Red	Green	Blue	Total_counts
0	126	124	127	236206
1	197	194	197	357272
2	36	36	38	173498

Figure 18. Total RGB pixels in each cluster

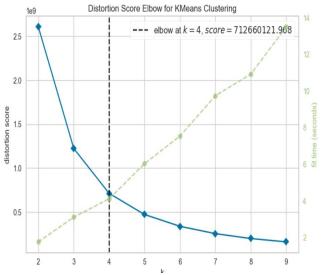


Figure 19. Optimal k using elbow method

C. CODE

FOR LAKE:

```
import cv2
import matplotlib.pyplot as plt
import matplotlib.cm as cm
import numpy as np
from sklearn.cluster import KMeans
from sklearn.metrics import
silhouette_samples, silhouette_score
import numpy as np
import pandas as pd
from yellowbrick.cluster import
KElbowVisualizer
"""## **Read Image**""
# Read the image using cv2
img path = 'MicrosoftTeams-image.png'
img = cv2.imread(img_path)
# Convert BGR image to RGB (matplotlib
uses RGB)
img_rgb = cv2.cvtColor(img,
cv2.COLOR BGR2RGB)
# Display the image using matplotlib
plt.imshow(img rgb)
plt.axis('off')
plt.show()
img_rgb.shape #Image shape
original shape = img.shape
# Reshape the image to a 2D array while
preserving the number of channels
reshaped_img = img_rgb.reshape(-1, 3)
reshaped_img.shape #Reshaped image shape
"""## **KMeans model with 3 clusters**""
kmeans = KMeans(n_clusters = 3,
random_state = 42, n_init = 10)
```

```
kmeans.fit(reshaped_img) #Train KMeans
cluster centroids =
kmeans.cluster_centers_
# Convert cluster centroids to integers
cluster centroids =
np.uint8(cluster_centroids)
# Create an image with the cluster
centroids
clustered_img =
cluster_centroids[kmeans.labels_]
# Reshape the clustered image back to its
original shape
clustered_img =
clustered_img.reshape(img_rgb.shape)
# Display the original image and the
clustered image using matplotlib
plt.figure(figsize=(10, 5))
plt.subplot(1, 2, 1)
plt.imshow(img_rgb)
plt.title('Original Image')
plt.axis('off')
plt.subplot(1, 2, 2)
plt.imshow(clustered_img)
plt.title('Clustered Image')
plt.axis('off')
plt.tight_layout()
plt.show()
cluster_centroids
"""## **Show Clustered Image**""
c=np.uint8(kmeans.cluster_centers_)
seg img = c[kmeans.labels ]
seg_img2 = seg_img.reshape((img.shape))
plt.imshow(seg_img2)
labels = kmeans.labels
```

```
# Count the number of pixels in each
cluster counts = np.bincount(labels,
minlength=3)
# Calculate the percentage of pixels in
each cluster
total pixels = labels.shape[0]
cluster_percentages = (cluster_counts /
total_pixels) * 100
# Print the cluster percentages
for cluster_idx, percentage in
enumerate(cluster_percentages):
    print(f"Cluster {cluster idx}:
{percentage:.2f}%")
plt.figure(figsize=(6, 6))
cluster colors =
kmeans.cluster_centers_.astype(int)
# Create the pie chart
pie = plt.pie(cluster_percentages,
labels=[f'Cluster {i}' for i in
range(3)], colors=cluster_colors / 255,
              autopct='%1.2f%%',
startangle=140)
# Set the text color inside the pie chart
to blue
for text in pie[1]:
    text.set_color('black')
# Set the text color outside the pie
chart to black
plt.setp(pie[2], color='white')
# Set the title
plt.title('Pie Chart')
# Display the pie chart
plt.show()
cluster_counts #Total number of pixels in
each clusters
```

```
#Create DataFrame containing RGB values
and total pixels of each clusters
data = {
        'Red': cluster_centroids[:, 0],
        'Green': cluster_centroids[:, 1],
        'Blue': cluster_centroids[:, 2],
        'Total_counts': cluster_counts}
df = pd.DataFrame(data)
df
# Calculate lake area using the previous
Dataframe
lake_area = (df.loc[1, 'Total_counts'] /
df['Total counts'].sum()) * 200000
lake area
# Define KMeans algorithm initialization
parameters
kmeans kwargs = {
    "init": "random",
Initialization method for centroids
    "n init": 10,
                             # Number of
time the k-means algorithm will be run
with different centroid seeds
    "random state": 42,
                             # Random
seed for reproducibility
# Initialize an empty list to store the
sum of squared distances for each k value
sse = []
# Loop through different values of k
(number of clusters)
for k in range(2, 11):
   # Create a KMeans model with the
current k value and specified
initialization parameters
   model = KMeans(n_clusters=k,
**kmeans kwargs)
   # Fit the KMeans model to the
reshaped image data
   model.fit(reshaped_img)
```

```
# Append the inertia (sum of squared
distances) of the fitted model to the
    sse.append(model.inertia_)
# Plot the Elbow curve to determine the
optimal number of clusters (k)
plt.plot(range(2, 11), sse)
plt.xticks(range(1, 11))
                           # Set x-axis
ticks to match the range of k values
plt.xlabel("k")
                            # Label for
x-axis
plt.ylabel("WCSS")
                           # Label for
v-axis
plt.title("Elbow Method for Optimal
k") # Title for the plot
plt.show()
                           # Display the
# Create a KElbowVisualizer instance with
the KMeans model and specify the range of
visualizer = KElbowVisualizer(model,
k=(2, 10)
# Fit the visualizer to the reshaped
image data
visualizer.fit(reshaped img)
# Display the elbow visualization
visualizer.show()
# Commented out IPython magic to ensure
Python compatibility.
X = reshaped img
range_n_clusters = [2, 3, 4, 5, 6]
for n_clusters in range_n_clusters:
    # Create a subplot with 1 row and 2
columns
    fig, (ax1, ax2) = plt.subplots(1, 2)
    fig.set size inches(18, 7)
    # The 1st subplot is the silhouette
```

```
ax1.set_xlim([-0.1, 1])
    # The (n clusters+1)*10 is for
inserting blank space between silhouette
    # plots of individual clusters, to
demarcate them clearly.
    ax1.set_ylim([0, len(X) + (n_clusters
+ 1) * 10])
    # Initialize the clusterer with
n clusters value and a random generator
    # seed of 10 for reproducibility.
    clusterer =
KMeans(n_clusters=n_clusters,
n_init="auto", random_state=10)
    cluster labels =
clusterer.fit predict(X)
    # The silhouette_score gives the
average value for all the samples.
    # This gives a perspective into the
density and separation of the formed
    # clusters
    silhouette_avg = silhouette_score(X,
cluster_labels)
    print(
        "For n_clusters =",
        n clusters,
        "The average silhouette score is
        silhouette_avg,
    # Compute the silhouette scores for
each sample
    sample silhouette values =
silhouette samples(X, cluster labels)
    y lower = 10
    for i in range(n clusters):
        # Aggregate the silhouette scores
for samples belonging to
        # cluster i, and sort them
        ith_cluster_silhouette_values =
sample silhouette values[cluster labels
== i]
```

```
ith_cluster_silhouette_values.sor
t()
        size cluster i =
ith_cluster_silhouette_values.shape[0]
        y_{upper} = y_{lower} +
size_cluster_i
        color = cm.nipy spectral(float(i)
/ n clusters)
        ax1.fill_betweenx(
            np.arange(y_lower, y_upper),
            ith_cluster_silhouette_values
            facecolor=color,
            edgecolor=color,
            alpha=0.7,
        # Label the silhouette plots with
their cluster numbers at the middle
        ax1.text(-0.05, y_lower + 0.5 *
size cluster i, str(i))
        # Compute the new y_lower for
next plot
        y_lower = y_upper + 10 # 10 for
the 0 samples
    ax1.set_title("The silhouette plot
for the various clusters.")
    ax1.set xlabel("The silhouette
coefficient values")
    ax1.set ylabel("Cluster label")
    # The vertical line for average
silhouette score of all the values
    ax1.axvline(x=silhouette avg,
color="red", linestyle="--")
    ax1.set yticks([]) # Clear the yaxis
labels / ticks
    ax1.set_xticks([-0.1, 0, 0.2, 0.4,
0.6, 0.8, 1])
```

```
# 2nd Plot showing the actual
clusters formed
    colors =
cm.nipy spectral(cluster labels.astype(fl
oat) / n_clusters)
    ax2.scatter(
        X[:, 0], X[:, 1], marker=".",
s=30, lw=0, alpha=0.7, c=colors,
edgecolor="k"
    # Labeling the clusters
    centers = clusterer.cluster_centers_
    # Draw white circles at cluster
centers
    ax2.scatter(
        centers[:, 0],
        centers[:, 1],
        marker="o",
        c="white",
        alpha=1,
        s = 200,
        edgecolor="k",
    for i, c in enumerate(centers):
        ax2.scatter(c[0], c[1],
marker="$%d$" % i, alpha=1, s=50,
edgecolor="k")
    ax2.set_title("The visualization of
the clustered data.")
    ax2.set xlabel("Feature space for the
1st feature")
    ax2.set ylabel("Feature space for the
2nd feature")
    plt.suptitle(
        "Silhouette analysis for KMeans
clustering on sample data with n clusters
= %d"
          % n clusters,
        fontsize=14,
        fontweight="bold",
```

```
plt.show()
"""## **KMeans with 4 clusters**""
kmeans_new = KMeans(n_clusters = 4,
random_state = 42, n_init = 10)
kmeans_new.fit(reshaped_img) #Train the
new Kmeans model
new centroids =
kmeans_new.cluster_centers_
# Convert new centroids to integers
new_centroids = np.uint8(new_centroids)
# Create an image with the new cluster
centroids
new_clustered_img =
new centroids[kmeans new.labels ]
# Reshape the new clustered image back to
its original shape
reshaped new clustered img =
new_clustered_img.reshape(img_rgb.shape)
# Display the original image and the new
clustered image using matplotlib
plt.figure(figsize=(10, 5))
plt.subplot(1, 2, 1)
plt.imshow(img rgb)
plt.title('Original Image')
plt.axis('off')
plt.subplot(1, 2, 2)
plt.imshow(reshaped_new_clustered_img)
plt.title('Clustered Image')
plt.axis('off')
plt.tight_layout()
plt.show()
"""## **Display Clustered Image**"""
```

```
center =
np.uint8(kmeans_new.cluster_centers_)
seg image = center[kmeans new.labels ]
seg_image2 = seg_image.reshape(img.shape)
plt.imshow(seg_image2)
labels_new = kmeans_new.labels_
# Count the number of pixels in each
cluster
cluster_counts_new =
np.bincount(labels_new, minlength=4)
# Calculate the percentage of pixels in
each cluster
total pixels new = labels new.shape[0]
cluster_percentages_new =
(cluster_counts_new / total_pixels_new) *
100
# Print the cluster percentages
for cluster idx, percentage in
enumerate(cluster_percentages_new):
    print(f"Cluster {cluster_idx}:
{percentage:.2f}%")
plt.figure(figsize=(6, 6))
cluster colors new =
kmeans new.cluster centers .astype(int)
# Create the pie chart
pie = plt.pie(cluster_percentages_new,
labels=[f'Cluster {i}' for i in
range(4)], colors=cluster_colors_new /
255.
              autopct='%1.2f%%',
startangle=140)
# Set the text color inside the pie chart
to blue
for text in pie[1]:
    text.set_color('black')
# Set the text color outside the pie
chart to black
plt.setp(pie[2], color='white')
```

```
# Set the title
plt.title('Pie Chart')
# Display the pie chart
plt.show()
cluster counts new #Contains total number
of pixels in each clusters
#Dataframe containing RGB values and
total number of pixels of each cluster
data_new = {
        'Red': new_centroids[:, 0],
        'Green': new_centroids[:, 1],
        'Blue': new centroids[:, 2],
        'Total counts':
cluster counts new}
df_new = pd.DataFrame(data_new)
df new
lake_area_new = (df_new.loc[1,
'Total_counts'] /
df_new['Total_counts'].sum()) * 200000
lake_area_new
```

FOR GRANITE:

```
import cv2
import matplotlib.pyplot as plt
import matplotlib.cm as cm
import numpy as np
from sklearn.cluster import KMeans
from sklearn.metrics import
silhouette_samples, silhouette_score
import numpy as np
import pandas as pd
from yellowbrick.cluster import
KElbowVisualizer

"""## **Read Image**""

# Read the image using cv2
img_path = 'MicrosoftTeams-image (1).png'
img = cv2.imread(img_path)
```

```
# Convert BGR image to RGB (matplotlib
uses RGB)
img_rgb = cv2.cvtColor(img,
cv2.COLOR BGR2RGB)
# Display the image using matplotlib
plt.imshow(img_rgb)
plt.axis('off')
plt.show()
img_rgb.shape #Image shape
original_shape = img.shape
# Reshape the image to a 2D array while
preserving the number of channels
reshaped img = img rgb.reshape(-1, 3)
reshaped_img.shape #Reshaped image shape
"""## **KMeans model with 3 clusters**""
kmeans = KMeans(n clusters = 3,
random_state = 42, n_init = 10)
kmeans.fit(reshaped_img) #Train KMeans
cluster centroids =
kmeans.cluster centers
cluster centroids =
np.uint8(cluster centroids)
# Create an image with the cluster
centroids
clustered img =
cluster_centroids[kmeans.labels_]
# Reshape the clustered image back to its
original shape
clustered img =
clustered img.reshape(img rgb.shape)
# Display the original image and the
clustered image using matplotlib
plt.figure(figsize=(10, 5))
```

```
plt.subplot(1, 2, 1)
plt.imshow(img_rgb)
plt.title('Original Image')
plt.axis('off')
plt.subplot(1, 2, 2)
plt.imshow(clustered_img)
plt.title('Clustered Image')
plt.axis('off')
plt.tight_layout()
plt.show()
cluster_centroids
"""## **Show Clustered Image**"""
c=np.uint8(kmeans.cluster_centers_)
seg img = c[kmeans.labels ]
seg_img2 = seg_img.reshape((img.shape))
plt.imshow(seg img2)
labels = kmeans.labels_
# Count the number of pixels in each
cluster
cluster counts = np.bincount(labels,
minlength=3)
# Calculate the percentage of pixels in
each cluster
total pixels = labels.shape[0]
cluster_percentages = (cluster_counts /
total pixels) * 100
# Print the cluster percentages
for cluster_idx, percentage in
enumerate(cluster percentages):
    print(f"Cluster {cluster_idx}:
{percentage:.2f}%")
plt.figure(figsize=(6, 6))
cluster colors =
kmeans.cluster_centers_.astype(int)
```

```
# Create the pie chart
pie = plt.pie(cluster_percentages,
labels=[f'Cluster {i}' for i in
range(3)], colors=cluster colors / 255,
              autopct='%1.2f%%',
startangle=140)
# Set the text color inside the pie chart
to blue
for text in pie[1]:
    text.set_color('black')
# Set the text color outside the pie
chart to black
plt.setp(pie[2], color='white')
# Set the title
plt.title('Pie Chart')
# Display the pie chart
plt.show()
cluster_counts #Total number of pixels in
each clusters
#Create DataFrame containing RGB values
and total pixels of each clusters
data = {
        'Red': cluster centroids[:, 0],
        'Green': cluster_centroids[:, 1],
        'Blue': cluster centroids[:, 2],
        'Total_counts': cluster_counts}
df = pd.DataFrame(data)
df
# Define KMeans algorithm initialization
parameters
kmeans_kwargs = {
    "init": "random",
Initialization method for centroids
    "n init": 10,
time the k-means algorithm will be run
with different centroid seeds
    "random state": 42,
seed for reproducibility
```

```
# Initialize an empty list to store the
sum of squared distances for each k value
sse = []
# Loop through different values of k
(number of clusters)
for k in range(2, 11):
    # Create a KMeans model with the
current k value and specified
initialization parameters
    model = KMeans(n_clusters=k,
**kmeans_kwargs)
    # Fit the KMeans model to the
reshaped image data
    model.fit(reshaped_img)
    # Append the inertia (sum of squared
distances) of the fitted model to the
    sse.append(model.inertia_)
# Plot the Elbow curve to determine the
optimal number of clusters (k)
plt.plot(range(2, 11), sse)
plt.xticks(range(1, 11))
                             # Set x-axis
ticks to match the range of k values
plt.xlabel("k")
                            # Label for
x-axis
plt.ylabel("WCSS")
                            # Label for
v-axis
plt.title("Elbow Method for Optimal
k") # Title for the plot
plt.show()
                           # Display the
# Create a KElbowVisualizer instance with
the KMeans model and specify the range of
visualizer = KElbowVisualizer(model,
k=(2, 10)
# Fit the visualizer to the reshaped
image data
visualizer.fit(reshaped img)
```

```
# Display the elbow visualization
visualizer.show()
# Commented out IPython magic to ensure
Python compatibility.
X = reshaped img
range_n_clusters = [2, 3, 4, 5, 6]
for n clusters in range_n_clusters:
    # Create a subplot with 1 row and 2
columns
    fig, (ax1, ax2) = plt.subplots(1, 2)
    fig.set_size_inches(18, 7)
    # The 1st subplot is the silhouette
plot
    ax1.set_xlim([-0.1, 1])
    # The (n clusters+1)*10 is for
inserting blank space between silhouette
    # plots of individual clusters, to
demarcate them clearly.
    ax1.set_ylim([0, len(X) + (n_clusters
+ 1) * 10])
    # Initialize the clusterer with
n clusters value and a random generator
    # seed of 10 for reproducibility.
    clusterer =
KMeans(n_clusters=n_clusters,
n init="auto", random state=10)
    cluster labels =
clusterer.fit_predict(X)
    # The silhouette score gives the
average value for all the samples.
    # This gives a perspective into the
density and separation of the formed
    # clusters
    silhouette_avg = silhouette_score(X,
cluster labels)
    print(
        "For n_clusters =",
        n clusters,
```

```
"The average silhouette_score is
        silhouette_avg,
    # Compute the silhouette scores for
each sample
    sample silhouette values =
silhouette samples(X, cluster labels)
    y_lower = 10
    for i in range(n_clusters):
        # Aggregate the silhouette scores
for samples belonging to
        # cluster i, and sort them
        ith cluster silhouette values =
sample_silhouette_values[cluster_labels
== i]
        ith cluster silhouette values.sor
t()
        size cluster i =
ith_cluster_silhouette_values.shape[0]
        y_upper = y_lower +
size_cluster_i
        color = cm.nipy spectral(float(i)
 n clusters)
        ax1.fill_betweenx(
            np.arange(y_lower, y_upper),
            ith_cluster_silhouette_values
            facecolor=color,
            edgecolor=color,
            alpha=0.7,
        # Label the silhouette plots with
their cluster numbers at the middle
        ax1.text(-0.05, y lower + 0.5 *
size_cluster_i, str(i))
        # Compute the new y_lower for
next plot
```

```
y_lower = y_upper + 10 # 10 for
the 0 samples
    ax1.set title("The silhouette plot
for the various clusters.")
    ax1.set_xlabel("The silhouette
coefficient values")
    ax1.set_ylabel("Cluster label")
    # The vertical line for average
silhouette score of all the values
    ax1.axvline(x=silhouette_avg,
color="red", linestyle="--")
    ax1.set_yticks([]) # Clear the yaxis
labels / ticks
    ax1.set_xticks([-0.1, 0, 0.2, 0.4,
0.6, 0.8, 1])
    # 2nd Plot showing the actual
clusters formed
    colors =
cm.nipy_spectral(cluster_labels.astype(fl
oat) / n clusters)
    ax2.scatter(
        X[:, 0], X[:, 1], marker=".",
s=30, lw=0, alpha=0.7, c=colors,
edgecolor="k"
    # Labeling the clusters
    centers = clusterer.cluster centers
    # Draw white circles at cluster
centers
    ax2.scatter(
        centers[:, 0],
        centers[:, 1],
        marker="o",
        c="white",
        alpha=1,
        s = 200,
        edgecolor="k",
    for i, c in enumerate(centers):
```

```
ax2.scatter(c[0], c[1],
marker="$%d$" % i, alpha=1, s=50,
edgecolor="k")
    ax2.set_title("The visualization of
the clustered data.")
    ax2.set xlabel("Feature space for the
1st feature")
    ax2.set ylabel("Feature space for the
2nd feature")
    plt.suptitle(
        "Silhouette analysis for KMeans
clustering on sample data with n_clusters
= %d"
          % n clusters,
        fontsize=14,
        fontweight="bold",
plt.show()
.....
"""## **KMeans with 4 clusters**""
kmeans_new = KMeans(n_clusters = 4,
random state = 42, n init = 10)
kmeans new.fit(reshaped img) #Train the
new Kmeans model
new centroids =
kmeans_new.cluster_centers_
# Convert new centroids to integers
new centroids = np.uint8(new centroids)
# Create an image with the new cluster
centroids
new clustered img =
new_centroids[kmeans_new.labels_]
# Reshape the new clustered image back to
its original shape
reshaped_new_clustered_img =
new clustered img.reshape(img rgb.shape)
```

```
# Display the original image and the new
clustered image using matplotlib
plt.figure(figsize=(10, 5))
plt.subplot(1, 2, 1)
plt.imshow(img_rgb)
plt.title('Original Image')
plt.axis('off')
plt.subplot(1, 2, 2)
plt.imshow(reshaped_new_clustered_img)
plt.title('Clustered Image')
plt.axis('off')
plt.tight_layout()
plt.show()
"""## **Display Clustered Image**"""
center =
np.uint8(kmeans new.cluster centers )
seg image = center[kmeans new.labels ]
seg image2 = seg_image.reshape(img.shape)
plt.imshow(seg image2)
labels new = kmeans new.labels
# Count the number of pixels in each
cluster
cluster counts new =
np.bincount(labels new, minlength=4)
# Calculate the percentage of pixels in
each cluster
total pixels new = labels new.shape[0]
cluster_percentages_new =
(cluster_counts_new / total_pixels_new) *
100
# Print the cluster percentages
for cluster idx, percentage in
enumerate(cluster_percentages_new):
    print(f"Cluster {cluster_idx}:
{percentage:.2f}%")
```

```
plt.figure(figsize=(6, 6))
cluster_colors_new =
kmeans_new.cluster_centers_.astype(int)
# Create the pie chart
pie = plt.pie(cluster_percentages_new,
labels=[f'Cluster {i}' for i in
range(4)], colors=cluster_colors_new /
255,
              autopct='%1.2f%%',
startangle=140)
# Set the text color inside the pie chart
to blue
for text in pie[1]:
    text.set_color('black')
# Set the text color outside the pie
chart to black
plt.setp(pie[2], color='white')
# Set the title
plt.title('Pie Chart')
# Display the pie chart
plt.show()
#cluster counts new #Contains total
number of pixels in each clusters
#Dataframe containing RGB values and
data_new = {
        'Red': new centroids[:, 0],
        'Green': new centroids[:, 1],
        'Blue': new_centroids[:, 2],
        'Total counts':
cluster counts new}
df new = pd.DataFrame(data new)
df_new
```



Adhip Bhattarai is a dedicated individual pursuing a Bachelor's degree in Computer Engineering at Tribhuvan University. With a strong passion for machine learning and data science, he is constantly exploring the latest advancements in these fields.

Although he may not have notable accomplishments just yet, Adhip's enthusiasm and drive for learning and applying cutting-edge technologies make him a promising and ambitious individual in the world of computer engineering.



Bishal Rijal is a dedicated individual currently studying Bachelor's in Computer and Technology at Tribhuvan University. Bishal's enthusiasm for research and innovation has led him to undertake various projects and engage in practical applications of his knowledge. He

continually seeks to deepen in understanding of the subject matter, staying up-to-date with the latest advancements and trends. With his relentless determination, inquisitive mindset, and expertise in machine learning and data science, Bishal Rijal is poised to make significant contributions to the ever-evolving field of technology.