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PROJECT 1 COLOR COMPRESSION

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1 Project ideas and K-means clustering algorithm

1.1 Project ideas

In computer, a picture can be stored in an array of pixels. In practical, there are many types of pictures. When the computer read a color picture, it will read a picture as an pixel array which is specified by the size of the picture and number of channels in one pixel. And therefore, we have countless number of matrices. Nowadays, the number of channels of common pictures are 3 represents for RGB color with the values in range [0, 255].

K-means clustering is a method of vector quantization, originally from signal processing, that aims to partition n observations into k clusters in which each observation belongs to the cluster with the nearest mean (cluster centers or cluster centroid), serving as a prototype of the cluster.

In this project, we want to compress the color of the picture therefore we have to group some pixels together to generate some clusters. And then we will find the representative pixel for each cluster, which will usually be the center of that cluster. But the special point here that we don't know which criteria to group pixels together from. Therefore we can choose k randomly to compress the color of a picture into k clusters only.

1.2 Description of K-means clustering algorithm

In here, I will summarize of K-means clustering algorithm

Input: Data X and a number of clusters $k_clusters$ you want to group.

Output: All centers M and labels vectors for each pixels.

K-means clustering algorithm has 5 steps to follow:

- Step 1: Pick k clusters randomly outside or inside the pixel array for the initial centers.
- Step 2: Assign each pixels into the clusters with the nearest distance to their *centroids*
- Step 3: If the assignment of each pixels into one cluster in step 2 doesn't change from the previous iteration then we can stop the algorithm.
- Step 4: Update the *centroids* for each clusters by getting the mean of all data-points have been assigned to each clusters in step 2.
- Step 5: Go back to step 2.

2 Functions description

```
def kmeans(img_1d, k_clusters, max_iter, init_centroids='random'):
      # Get 3 dimensions (height, width, num_channels) information of image
      height = img_1d.shape[0]
      width = img_1d.shape[1]
      num_channels = img_1d.shape[2]
      # Convert an array with 3 dimensions into 2 dimensions
      img_1d = img_1d.reshape(height * width, num_channels)
      # Create k_clusters centroids with init_centroids = 'random' or 'in_pixels'
      centroids = [kmeans_init_centers(img_1d, k_clusters, init_centroids)]
      labels = []
      while max_iter:
        labels.append(kmeans_assign_labels(img_1d, centroids[-1]))
12
        new_centroids = kmeans_update_centers(img_1d, labels[-1], k_clusters,
13
                                                                      num_channels)
14
        if has_converged(centroids[-1], new_centroids): break
        centroids.append(new_centroids)
16
        max_iter -= 1
17
      return (centroids, labels)
```

- This function will return the centroids and the labels vectors of each data-points.
- In the first step, we can easily see that the pixel array of a picture read in will have the dimensions of 3, this will be a little challenge to deal with it. So we will make it easier to manipulate by reshape the 3 dimensions into 2 dimensions.

Then we will initialize the centers of $k_clusters$ clusters depend on the parameter $init_centroids$ pass in.

If *init* centroids = 'random' -> centroid has 'c' channels, with 'c' is initial random in [0,255].

If *init* centroids = 'in pixels' -> centroid is a random pixels of original image.

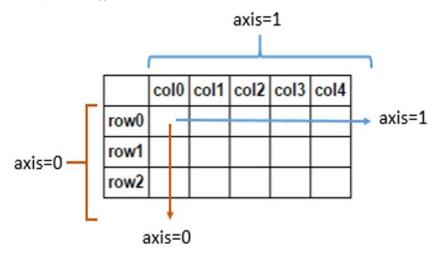
- In step 2, we will assign each pixels into the clusters with the nearest distance to their *centroids* in line 12.
- In step 3, we will check if the assignment of each pixels into one cluster in step 2 doesn't change from the previous iteration then we can stop the algorithm in line 15.
- In step 4, we will append the *new centroids* into the *centroids* array for later check in line 16.
- We will stop the algorithm in *max* iter times or when the converge condition has been reach.
- At the end, the function will return a tuple of 2 values *centroids* and *labels*.

- This function will return $k_clusters$ pixels randomly inside or outside our image array as the center points for each clusters.
- In first condition, if $init_centroids = \text{`in_pixels'}$ so we will use np.random.choice() from numpy module for pick a random array with $k_clusters$ pixels from the img_1d array to make centroids array inside our image array. Therefore, the centroid is a random pixels of original image.
- In second condition, if *init_centroids* = 'random' then we will use *np.random.choice*() function from numpy module for an array picked randomly with size = (*k_clusters*, *num_channels*) and we want all centers are distinguish so we add *replace* = *False*. At the end, the centroid will have 'c' channels, with 'c' is initial random in [0,255].

```
def kmeans_assign_labels(img_1d, centroid):
    distances = []
    for i in range(centroid.shape[0]):
    # Distance between each elements in img_1d vs centroid[i] in Euclidean distance
    distance = np.sqrt(np.sum((img_1d - centroid[i]) ** 2, axis=1))
    distances.append(distance)
    # convert list to np.ndarray
    distances = np.array(distances)
    # Return the indices of clusters where each pixels with minimum distances to all centroids belong to
    return np.argmin(distances.T, axis = 1)
```

• This function will return the indices of clusters where each pixels with minimum distances to all *centroids* belong to or we can call it as *labels* in previous section.

• In this function, I will calculate the distances of all pixels to all centroids and pick the nearest centroid for each pixels. I will calculate the distance in Euclidean distance by using this formula $np.sqrt(np.sum((img_1d-centroid[i])**2, axis=1))$, and parameter axis=1 is for calculate sum in horizontal, as the figure demonstration below.



- Then we will get a *distances* array and we will convert to np.ndarray for later use.
- Now, our distances array have $size = (num_channels, img_1d.shape[0])$. Then we will transpose it to get array with $size = (img_1d.shape[0], num_channels)$.
- At the end, we will use np.argmin() function from numpy module and pass in the transpose form of distances np.ndarray with axis = 1 to get the indices of clusters where each pixels with minimum distances to all centroids belong to.

```
def kmeans_update_centers(img_1d, label, k_clusters, num_channels):
    centroids = np.zeros((k_clusters, num_channels))

for k in range(k_clusters):
    # collect all points assigned to the k-th cluster
    clusterk = img_1d[label == k, :]
    # if clusterk is empty we don't need to update
    if len(clusterk) == 0:
        continue
    # Take average of all datapoints belong to cluster k
    centroids[k, :] = np.mean(clusterk, axis = 0)
    return centroids
```

- This function will return the new *centroids* updated from all clusters get from the latest labels.
- First, we will collect all the *data-points* assigned to the k-th cluster.
- Secondly, if the k-th cluster doesn't have any data-points so we don't need to update the center of that k-th cluster.
- Thirdly, We will take average of all data-points belong to k-th cluster. And we have to iterate these steps for all k clusters.
- Finally, after updated all k clusters we will return the new *centroids*.

```
def has_converged(centers, new_centers):
    # if we have absolute(centers[i] - new_centers[i]) <= atol(=1)
    # => the difference of two RGBs are less than 1
    # we can approve these both RGBs are the same.
    return np.allclose(centers, new_centers, atol = 1)
```

• This function will check the converge condition for the new *centroids* and the old *centroids*, if these two arrays have each relative elements is close to each other and the *errornumber* is less than 1 by using atol = 1 then we can consider that both RGBs colors are the same.

```
def get_new_image(img_1d, k_clusters, centroid, label):
      # Get 3 dimensions (height, width, num_channels) information of image
      height = image_array.shape[0]
      width = image_array.shape[1]
      num_channels = image_array.shape[2]
      # Because the original image_array have 3 dimensions so we have to reshape it to
       2 dimensions
      img_1d = img_1d.reshape(height * width, num_channels)
      # Create new image array by adding the cluster index (=centroid[k]) to each
      pixels
      new_img = np.zeros((img_1d.shape[0], img_1d.shape[1]))
      \# Get the new image array with k\_{clusters} centroids assigned to each pixels.
10
11
      for k in range(k_clusters):
       new_img[label == k, :] += centroid[k]
12
13
      new_img = new_img.reshape(height, width, num_channels)
     return new_img
```

• This function will return the new image array with $k_clusters\ centroids$ assigned to each pixels that we will export to .png and .pdf files at the end.

3 Results

I prepared 2 pictures for testing time.



Figure 1: avatar.jpg



Figure 2: img.jpg

3.1 Testing 1: K = 3

3.1.1 Sample 1

With sample 1, we will get the centroids and labels results as below. My new centroids result:

```
[[ 51.19524933 152.9280978 253.43506519]
[ 52.54914301 56.8495358 92.77106767]
[ [ 232.01616437 220.94881897 220.59589693]]
```

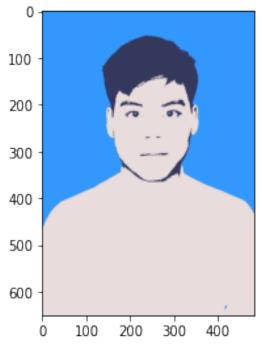


Figure 3: my_avatar_3.jpg

Sklearn centroids result:

```
[[ 51.19613927 152.92673114 253.43151257]
2 [231.93665774 220.82114514 220.46234457]
3 [ 51.30013981 56.10282776 92.4355297 ]]
```

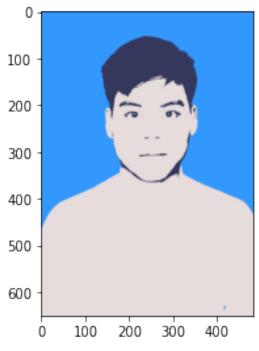


Figure 4: Sklearn_avatar_3.jpg

 \Rightarrow We can see that elements in order [0, 1, 2] of my new centroids are approximately with elements in order [0, 2, 1] in Sklearn centroids. And the picture show between both ways are also the same.

3.1.2 Sample 2

With sample 2, we will get the centroids and labels results as below. My new centroids result:

```
[[152.49458003 70.77234422 230.48494644]
[108.40246167 32.02635536 87.00908169]
[240.40357334 168.593392 213.77850318]]
```

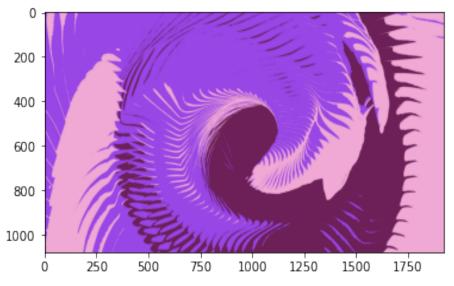


Figure 5: *my_img_3.jpg*

Sklearn centroids result:

[[150.98283514 70.00983381 229.72601646]

```
2 [239.94627034 167.68524163 214.65643656]
3 [109.36530543 32.08481787 85.15615837]]
```

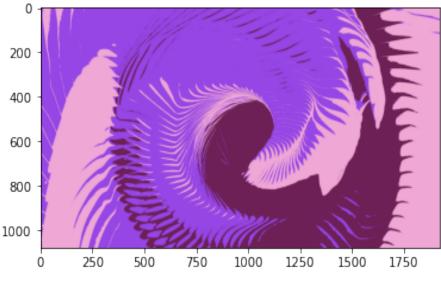


Figure 6: Sklearn_img_3.jpg

 \Rightarrow We can see that elements in order [0, 1, 2] of my new centroids are approximately with elements in order [0, 2, 1] in Sklearn centroids. And the picture show between both ways are also the same.

3.2 Testing 2: K = 5

3.2.1 Sample 1

With sample 1, we will get the centroids and labels results as below. My new centroids result:

```
[[237.17988422 235.71493613 238.71842856]
[183.0349162 137.36658686 130.67936951]
[223.59273408 181.44805243 168.96662921]
[51.13458859 152.93720386 253.49105014]
[28.92059499 44.65924751 88.76326151]]
```

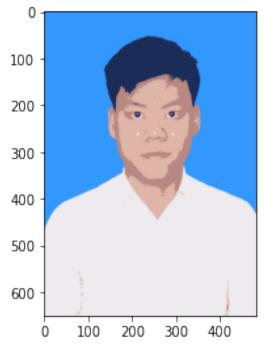


Figure 7: $my_avatar_5.jpg$

Sklearn centroids result:

```
[[237.17835512 235.70043832 238.69961747]
[ 51.13306751 152.93809229 253.49341127]
[ [180.81530172 135.26429598 129.06688218]
[ [222.76587715 180.41203321 167.96858714]
[ [28.59175495 44.52006368 88.74908053]]
```

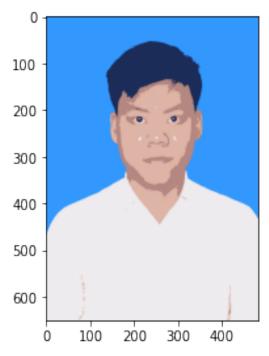


Figure 8: Sklearn_avatar_5.jpg

 \Rightarrow We can see that elements in order [0, 1, 2, 3, 4] of my new centroids are approximately with elements in order [0, 2, 3, 1, 4] in Sklearn centroids. And the picture show between both ways are also the same.

3.2.2 Sample 2

With sample 2, we will get the centroids and labels results as below. My new centroids result:

```
[[225.94005141 76.23554964 92.81608875]

[184.41939849 87.45554131 242.36262026]

[[61.20119247 25.94475322 70.36237693]

[241.88296006 189.97461332 231.20937672]

[105.90379453 49.6280514 204.01769168]]
```

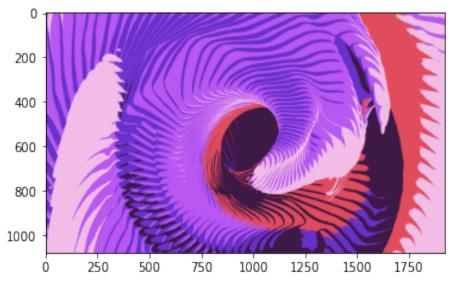


Figure 9: *my_img_5.jpg*

Sklearn centroids result:

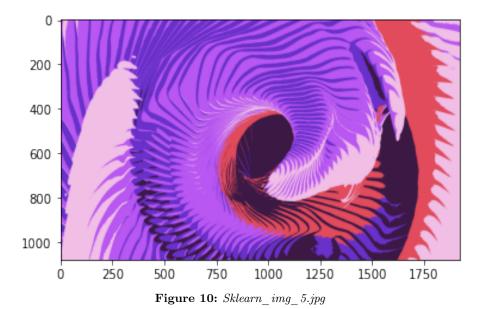
```
[[105.56029721 49.23909135 202.91437094]

[241.96136162 190.19693936 230.95834668]

[225.63839039 75.76976228 92.37659333]

[184.31749334 87.48831962 242.33615897]

[60.22890173 25.90469444 69.19434866]]
```



⇒ We can see that elements in order [0, 1, 2, 3, 4] of my new centroids are approximately with

elements in order [2, 3, 4, 1, 0] in Sklearn centroids. And the picture show between both ways are also the same.

3.3 Testing 3: K = 7

3.3.1 Sample 1

With sample 1, we will get the centroids and labels results as below. My new centroids result:

```
[[200.66189834 155.36184558 144.38218913]
[[154.21294206 109.71162528 110.63694507]
[[225.47337502 223.78750553 229.20712526]
[[51.10398854 152.92473668 253.4950666 ]
[[228.10697128 186.355668 173.86778084]
[[26.45727991 43.66089165 88.68510158]
[[238.73843986 237.30446404 239.98408037]]
```

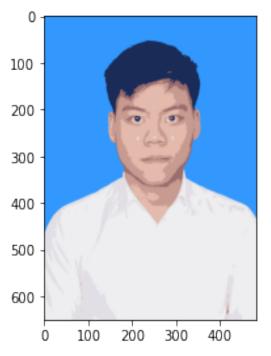


Figure 11: my_avatar_7.jpg

Sklearn centroids result:

```
[[237.22112779 235.78870616 238.79750074]
[ 51.13558876 152.93647132 253.48837191]
[ 69.55573248 61.79578025 89.03423567]
[ [208.0948382 163.46853656 150.79033703]
[ [172.61675158 126.37305828 122.22530135]
[ [ 16.03271923 39.22903464 88.75806927]
[ [230.42063091 190.31753943 179.18567823]]
```

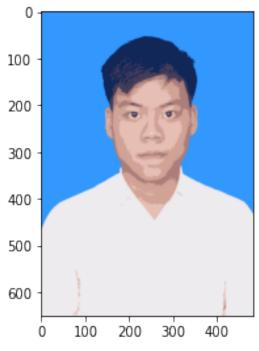
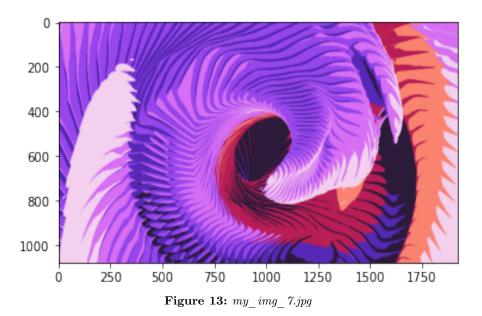


Figure 12: Sklearn_avatar_7.jpg

⇒ We can see that elements of my new centroids are elements in Sklearn centroids are a little bit different between due to small iteration and number of channel. And the picture show between both ways are also still the same.

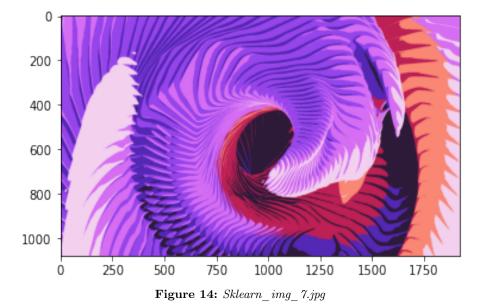
3.3.2 Sample 2

With sample 2, we will get the centroids and labels results as below. My new centroids result:



Sklearn centroids result:

```
[[214.80672576 110.90651762 244.7791835]
[188.20019256 32.19658732 84.24964964]
[84.84550239 40.77940424 180.35044748]
[242.70645544 208.33447862 238.88828772]
[250.6891642 134.3522698 116.17787354]
[45.37386589 26.24445252 57.33711496]
[150.78475293 69.92127439 235.55118058]]
```



⇒ We can see that elements of my new centroids are elements in Sklearn centroids are a little bit different between due to small iteration and number of channel. And the picture show between both ways are also still the same.

4 Comment

Overall, this k-mean clustering program give quite good answer compare to the answer provided by KMeans() of scikit-learn module. And k-mean clustering algorithm are easy to implemented due to

its clearly run-flows. Because at the very last iteration, *centroids* will be updated with a very small error number or maybe not be changed because it will be closer to the center of its cluster time by time. And with a small number of iteration we cannot get the accuracy result in a large data given. And with the randomly chosen each restart we will get different results each time so we have to redo it time by time and choose the most accuracy result as possible.

5 References

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