programming-4

May 1, 2024

Assignment 4

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[19]: import os
      import warnings
      warnings.filterwarnings("ignore")
      dog_image_directory = r'C:\Users\injam\Desktop\DM_Assignment_1\Cropped'
      from skimage import filters
      from skimage import io, color
      from skimage import exposure
      import numpy as np
      def calculate_angle(dx, dy):
          return np.mod(np.arctan2(dy, dx), np.pi)
      hist_images = []
      labels = []
      for index, breed in enumerate(os.listdir(dog_image_directory)):
          image_folder = os.path.join(dog_image_directory, breed)
          for image in os.listdir(image_folder):
              img = io.imread(os.path.join(image_folder, image.strip()))
              img = color.rgb2gray(img)
              sobel_img = calculate_angle(filters.sobel_h(img), filters.sobel_v(img))
              hist, _ = exposure.histogram(sobel_img, nbins=36)
              hist_images.append(hist / np.sum(hist))
              labels.append(index)
      hist_images = np.array(hist_images)
      labels = np.array(labels)
```

```
[20]: from sklearn.decomposition import PCA
model=PCA(2)
pca2 = model.fit_transform(hist_images)
```

```
[25]: from sklearn.cluster import KMeans, BisectingKMeans, SpectralClustering from sklearn.metrics import silhouette_score, fowlkes_mallows_score
```

```
clustering_models = {
          "KMeans Random": KMeans(init="random", n_clusters=4),
          "KMeans KMeans++": KMeans(init="k-means++", n_clusters=4),
          "BisectingKmeans": BisectingKMeans(init="random", n_clusters=4),
          "SpectralClustering": SpectralClustering(n_clusters=4)
      }
      for method, model in clustering models.items():
          fitted_model = model.fit(pca2)
          print(method)
          print("fowlkes :" + str(fowlkes_mallows_score(labels, fitted_model.
       ⇒labels )))
          print("silhouette :" + str(silhouette_score(pca2, fitted_model.labels_)))
          print("\n")
     KMeans Random
     fowlkes :0.30353266800098666
     silhouette :0.4185683795750631
     KMeans KMeans++
     fowlkes :0.31520188962993373
     silhouette: 0.4135128261083564
     BisectingKmeans
     fowlkes :0.333017685831437
     silhouette :0.4332733726639791
     SpectralClustering
     fowlkes :0.34834824513062457
     silhouette :-0.029977113490016673
[22]: from sklearn.cluster import DBSCAN
      db = DBSCAN(eps=0.015, min_samples=2).fit(pca2)
      # Number of clusters in labels, ignoring noise if present.
      n_clusters_ = len(set(labels)) - (1 if -1 in labels else 0)
      n_noise_ = list(labels).count(-1)
      print("Estimated number of clusters: %d" % n_clusters_)
```

Estimated number of clusters: 4

print("Estimated number of noise points: %d" % n_noise_)

```
Estimated number of noise points: 0
     0.1 \text{ eps}=0.015, \min \text{ samples}=2
[23]: print("fowlkes :" +str(fowlkes_mallows_score(labels,db.labels_)))
      print("silhoutte :"+str(silhouette_score(pca2,db.labels_)))
     fowlkes :0.48807103785693184
     silhoutte: 0.4802708103721345
[24]: from sklearn.cluster import AgglomerativeClustering
      linkage=["ward", "complete", "average", "single"]
      for link in linkage:
          clustering=AgglomerativeClustering(n_clusters=4,linkage=link)
          pred=clustering.fit(pca2)
          print(link)
          print("folwkes :"+str(fowlkes_mallows_score(labels,pred.labels_)))
          print(" silhouette :"+str(silhouette_score(pca2,pred.labels_))+"\n")
     ward
     folwkes :0.3146235000818094
      silhouette: 0.40256571902377786
     complete
     folwkes: 0.40659482650984735
      silhouette: 0.401907724039627
     average
     folwkes : 0.4965405634587954
      silhouette :0.6257171216011216
     single
     folwkes :0.4977891732358187
      silhouette :0.6869276296961174
                                                                              agglomerative
     0.1.1 fowlkes
                         \mathbf{score}
                                    \mathbf{best}
                                              \mathbf{to}
                                                      worst
            single, Average, dbscan, complete, spectral, bisecting, k-means++, ward, random
```

 \mathbf{to}

single, average, dbscan, bisecting, random, k-means++, ward, complete, spectral

worst

agglomerative

best

0.1.2 silhouette

[]:

score