Lab4: K-means

1K-means算法原理

1.1 算法概述

k-means是基于划分的动态聚类算法,通过相似度的计算将样本划分为对应类,同时依靠样本均值来计算聚类中心,通过迭代来遍历可能的簇划分,利用贪心策略求得近似解,容易陷入局部最优。本质上是"物以类聚"。

1.2 算法步骤

- 1. 确定分类数目k
- 2. 初始化聚类中心 $\mu_1, \mu_2, \mu_3 \dots \mu_k$
- 3. 计算样本到各个聚类中心的距离,选择最小距离 (最高相似度) 给样本分类
- 4. 使用类中样本均值重新计算聚类中心
- 5. 重复step3, step4直到重新计算距离并分类后,没有样本的类别发生改变

1.3 关键问题

1.3.1 k取值问题

初始设定分类数目k会极大程度影响分类效果。k过小导致模型欠拟合,k过大则有可能类别过密。因此确定一个合适的k值很重要。通过观察可以粗略确定可视化后图片像素点聚类分割的效果。为了定量确定k的聚类效果,可以采用以下两种评估方式:

- SSE: 聚类内部相似度均值
- 轮廓系数 (Silhouette Coefficient)

$$s = \frac{1}{n} \sum \frac{b-a}{max(a,b)}$$

a表示向量到所有它属于的簇中其它点的距离,即簇内不相似度;b表示向量到距离最近一簇内的所有点的平均距离,即簇间不相似度

1.3.2 聚类中心初始化问题

聚类的最终效果严重依赖于初始聚类的中心点的选择。如果选择不恰当,将会非常容易陷入局部最优解。因此,此处使用Kmeans++手段优化。

Kmeans++的基本思想是初始的聚类中心距离越远越好。算法步骤如下:

- 1. 从数据集 χ 中随机选择一个样本点作为第一个聚类中心点 C_1
- 2. 计算 χ 中剩余样本点x距离当前已有聚类中心点的最短距离D(x)和这个样本点被选为聚类中心的概率 $P(x)=rac{D(x)^2}{\sum_{x\in X}D(x)^2}$

3. 选择样本点作为下一个聚类中心,重复step2直到选择了k个聚类中心

2 核心代码

代码使用的库如下

```
In []: from sklearn import preprocessing
   import PIL.Image as Image
   import numpy as np
   import sys
   import pandas as pd
   import random
   import matplotlib.pyplot as plt
```

2.1 图像处理

图片为24位图像,每个像素点由三原色即RGB组成,每种颜色取值0~255,因此像素点的颜色值由3个通道表示。把图片转化成可以处理的数据,使用numpy库将图像的RGB表示处理成 R^3 矩阵。

为什么要令矩阵中元素值介于0~1之间?对输入图像进行归一化操作。将0~255数值映射到0~1中,使得数据处理更加统一方便,便于模型快速收敛。

```
In []: def load_pic(file_path):
    """
    Generate numpy matrix according to input file path.
    Return values include list, pic_width & pic_height.
    """
    with open(file_path, 'rb') as f: # open picture file
        data = []
        position = []
        image = Image.open(f)
        width, height = image.size
        for x in range(width):
            r,g,b = image.getpixel((x,y)) # get rgb at(x,y)
            position.append((x,y))
            data.append([r/255.0,g/255.0,b/255.0]) # normalize
    print("Picuture successfully loaded.")
    return data, width, height, position
```

2.2 kmeans

计算距离的方式采用欧氏距离。

```
In []: def cal_dist(dataset, centroids):
    '''Calculate Euclidean distance '''
    dist_list = []
    for data in dataset:
        # fill the data k times to minus centroids
        diff = np.tile(data,(len(centroids),1))-centroids
        dist_square = np.sum(np.square(diff),axis=1)
```

dist_list.append(dist_square)

```
return np.array(dist_list)

def cal_centroids(dataset,centroids):
    '''calculate new centroids'''
    dist_list = cal_dist(dataset,centroids)
    # calculate the centroids index of min distance
    min_idx = np.argmin(dist_list,axis=1)
    new_centroids = pd.DataFrame(dataset).groupby(min_idx).mean().values

# calculate the changes
    changes = new_centroids-centroids

return changes,new_centroids

In []: def Kmeans(X,k, position):
    centroids = random.sample(X,k) # randomly sample k centroids
```

```
changes, centroids = cal_centroids(X,centroids)
# continue updating centroids until changes equals zero
# or stop when epoch reaches 100000
epoch = 1
while np.any(changes != 0) or epoch > 50:
    changes, centroids = cal_centroids(X,centroids)
    print("Finished: k=%(k)i, epoch=%(e)i"% {"k":k,"e":epoch})
   epoch += 1
# calculate clusters according to centroids
cluster = []
rgb_cluster = [] # calculate rgb clusters to estimate results(SSE)
for i in range(k):
   cluster.append([])
   rgb_cluster.append([])
dist_list = cal_dist(X,centroids)
min_idx = np.argmin(dist_list,axis=1)
for i,j in enumerate(min_idx):
    cluster[j].append(position[i])
   rgb_cluster[j].append(X[i])
print("Finish Kmeans: K=%i"% k)
return centroids,cluster,rgb_cluster
```

2.3 可视化结果

```
In [ ]:
    def draw_result(pic_path,results,width,height):
        plt.title("Picture Segment using Kmeans") # title for the pic
        plt.subplot(231)
        plt.imshow(Image.open(pic_path)) # display original pic
        plt.axis('off') # shut down the axis
        plt.xticks([])
        plt.yticks([]) # shut down x,y values

        colors = [(0, 0, 255),(255, 0, 0),(0, 255, 0),(60, 0, 220),(167, 255, 167)]

        n = 2
        for res in results:
            new_pic = Image.new("RGB",(width,height))
            for i in range(len(res[0])):
```

```
In []: path = "./flowers.jpg"
  pic_data, width, height, position = load_pic(path)

res = []
  all_rgb = []
  for k in range(2,6):
        centroids, cluster, rgb_cluster = Kmeans(pic_data,k,position)
        res.append((centroids,cluster))
        all_rgb.append(rgb_cluster)

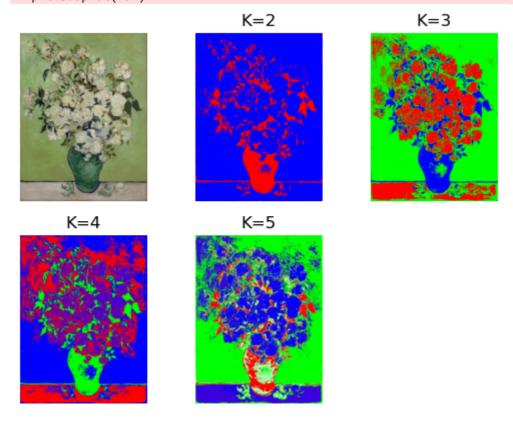
draw_result(path,res,width,height)
```

Picuture successfully loaded. Finished: k=2, epoch=1 Finished: k=2, epoch=2 Finished: k=2, epoch=3 Finished: k=2, epoch=4 Finished: k=2, epoch=5 Finished: k=2, epoch=6 Finished: k=2, epoch=7 Finished: k=2, epoch=8 Finished: k=2, epoch=9 Finished: k=2, epoch=10 Finish Kmeans: K=2 Finished: k=3, epoch=1 Finished: k=3, epoch=2 Finished: k=3, epoch=3 Finished: k=3, epoch=4 Finished: k=3, epoch=5 Finished: k=3, epoch=6 Finished: k=3, epoch=7 Finished: k=3, epoch=8 Finished: k=3, epoch=9 Finished: k=3, epoch=10 Finished: k=3, epoch=11 Finished: k=3, epoch=12 Finish Kmeans: K=3 Finished: k=4, epoch=1 Finished: k=4, epoch=2 Finished: k=4, epoch=3 Finished: k=4, epoch=4 Finished: k=4, epoch=5 Finished: k=4, epoch=6 Finished: k=4, epoch=7 Finished: k=4, epoch=8 Finished: k=4, epoch=9 Finished: k=4, epoch=10 Finished: k=4, epoch=11 Finished: k=4, epoch=12 Finish Kmeans: K=4 Finished: k=5, epoch=1 Finished: k=5, epoch=2 Finished: k=5, epoch=3 Finished: k=5, epoch=4 Finished: k=5, epoch=5 Finished: k=5, epoch=6 Finished: k=5, epoch=7 Finished: k=5, epoch=8 Finished: k=5, epoch=9 Finished: k=5, epoch=10 Finished: k=5, epoch=11 Finished: k=5, epoch=12 Finished: k=5, epoch=13 Finished: k=5, epoch=14 Finished: k=5, epoch=15 Finished: k=5, epoch=16 Finished: k=5, epoch=17 Finished: k=5, epoch=18 Finished: k=5, epoch=19 Finished: k=5, epoch=20 Finished: k=5, epoch=21

Finished: k=5, epoch=22

Finished: k=5, epoch=23 Finished: k=5, epoch=24 Finished: k=5, epoch=25 Finished: k=5, epoch=26 Finished: k=5, epoch=27 Finished: k=5, epoch=28 Finished: k=5, epoch=29 Finished: k=5, epoch=30 Finished: k=5, epoch=31 Finished: k=5, epoch=32 Finished: k=5, epoch=33 Finished: k=5, epoch=34 Finished: k=5, epoch=35 Finished: k=5, epoch=36 Finished: k=5, epoch=37 Finished: k=5, epoch=38 Finished: k=5, epoch=39 Finished: k=5, epoch=40 Finished: k=5, epoch=41 Finished: k=5, epoch=42 Finished: k=5, epoch=43 Finished: k=5, epoch=44 Finished: k=5, epoch=45 Finished: k=5, epoch=46 Finished: k=5, epoch=47 Finished: k=5, epoch=48 Finish Kmeans: K=5

C:\Users\A.R.L\AppData\Local\Temp\ipykernel_50076\2987248863.py:3: MatplotlibDepr
ecationWarning: Auto-removal of overlapping axes is deprecated since 3.6 and will
be removed two minor releases later; explicitly call ax.remove() as needed.
plt.subplot(231)



3 测试评估

由上述图片可见, k=3和k=4时拟合效果相对较好。相似点划分较为清晰, 能较好的区分物体前景和背景。

4参考资料

- 1. https://blog.csdn.net/qq_30759585/article/details/106269933
- 2. https://zhuanlan.zhihu.com/p/608317023
- 3. https://blog.csdn.net/qq_41498261/article/details/100727309
- 4. https://blog.csdn.net/qq_43741312/article/details/97128745