无监督学习 实验报告

PB17050965 朱子琦

实验环境

- python 3.8
- numpy 1.18.4
- matplotlib 3.2.2

实验内容

数据预处理与结果评测

数据预处理

原始数据各组数据尺度差异很大,很难正确判断是哪一组数据在其中产生影响,因此,先将其按比例缩放到[0,1]范围内,而后再按其到0.5的距离进行离散化,使得其能够分布在[-1,1]范围内

评价函数

兰德系数

根据兰德系数公式统计计算

```
def evaluate_RI(C, K):
evaluate_RI [使用兰德系数评价结果]
Args:
    C ([dict]): [真实结果]
    K ([dict]): [聚类结果]
Returns:
    [num]: [兰德系数值]
a, b, c, d = 0, 0, 0, 0
for i in range(len(C)):
    for j in range(len(C)):
        if i == j:
            continue
        else:
            if C[i] == C[j] and K[i] == K[j]:
                a += 1
            elif C[i] == C[j] and K[i] != K[j]:
            elif C[i] != C[j] and K[i] == K[j]:
            else:
                d += 1
RI = (a + d) / (a + b + c + d)
return RI
```

轮廓系数

在进行轮廓系数计算时,需要确定数据的分类方式,而后根据分类去计算该点到周围各簇以及本簇内其他点的距离,最后根据得到的a(i)和 b(i) 计算对应的SI

```
def evaluate_SI(k,label, data, center):
 def distance(v1, v2):
     return (v2.T @ v2 - 2 * v1.T @ v2 + v1.T @ v1) ** 0.5
 n,m = data.shape
 board = np.zeros([n,n])
 for i in range(n):
     for j in range(n):
         board[i,j] = distance(data[i],data[j])
 def a(i):
     avr, count = 0,0
     for j in range(n):
         if j != i and label[i] == label[j]:
             avr += board[i,j]
             count += 1
     return avr/count
 def b(i):
     d = np.zeros([k])
     count = np.zeros([k])
     for j in range(n):
         d[label[j]] += board[i,j]
         count[label[j]] += 1
     for j in range(k):
         d[j] = d[j] / count[j]
     mind = np.inf
     for j in range(k):
         if j != label[i] and d[j] < mind:</pre>
             mind = d[j]
     return mind
 def s(i):
     ai,bi = a(i),b(i)
     return (bi-ai)/max(ai,bi)
 slist = []
 for i in range(n):
     slist.append(s(i))
 return sum(slist)/n
```

PCA

PCA 是将n维特征映射到k维上,这k维是全新的正交特征也被称为主成分,是在原有n维特征的基础上重新构造出来的k维特征。

对原始数据进行降维处理的过程中,要降低到k维需要得到最大的k个特征向量,通过np.linalg求对应矩阵特征向量,而后根据threhold选出前k个特征向量,用对应的特征向量进行线性变换,得到的矩阵就是降维后的矩阵,并通过函数返回值返回。

```
def PCA(X, threhold=0.8):
PCA [使用PCA算法对数据进行降维处理]
Args:
    X ([np.array([n,m])]): [待降维的数据集np.array([n,m])]]
    threhold (float, optional): [PCA 降维阈值,选取特征累加贡献率达到该阈值的前k个特征]. Defaul
Returns:
    [np.array([n,k])]: [返回降低至k维后的数据集]
if threhold < 0 or threhold > 1:
    print("threhold should be between 0 and 1")
    return X
# 计算输入数据矩阵的特征向量
n, m = X.shape
scatter_matrix = X.T @ X
eig_val, eig_vec = np.linalg.eig(scatter_matrix)
eig_pairs = [(np.abs(eig_val[i]), eig_vec[:, i]) for i in range(m)]
eig_pairs.sort(reverse=True)
# 计算需要达到threhold的k
base, top, k = sum(eig val), 0, m
for i in range(m):
    top += eig_pairs[i][0]
    if top/base > threhold:
        k = i + 1
        break
# 选取前k个特征向量
feature = np.array([ele[1] for ele in eig_pairs[:k]])
data = X @ feature.T
return data
```

K-Means

k-means算法是通过创建k个中心,让数据点选择最近的中心称为簇。每轮迭代更新中心为当前簇的重心,而后进行下一轮迭代,直到每个点所在的簇不再发生改变则停止。

```
def k_means(k, data):
 def distance(v1, v2):
     return (v2.T @ v2 - 2 * v1.T @ v2 + v1.T @ v1) ** 0.5
 def avrcenter(v1, v2, k1, k2):
     return (v1 + v2) / (k1 + k2)
n, m = data.shape
 center = np.zeros([k, m])
 num = np.zeros([k])
label = [1]
next_label = []
for i in range(k):
     center[i] = data[int(n / k * i)]
while next_label != label:
     label = next_label.copy()
     next_label = [None]*n
     for i in range(n):
         d = []
         for j in range(k):
             d.append((distance(center[j], data[i]), j))
         d.sort()
         cluster = d[0][1]
         next_label[i] = cluster
         num[cluster] += 1
     for i in range(k):
         center[i] = np.zeros(m)
     for i in range(n):
         center[next_label[i]] += data[i]
     for i in range(k):
         center[i] = center[i] / num[i]
     num = np.zeros([k])
     t += 1
return k,label,center
```

实验结果

PCA降维的threhold

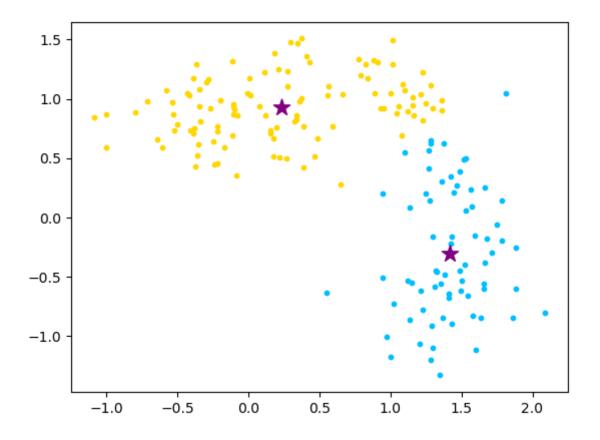
经实验,不同threhold范围内对应的结果维数k,兰德系数RI和轮廓系数SI为

threhold	k	兰德系数RI	轮廓系数SI
0.3759616388123276	1	0.7056433695169174	0.6316664749817953

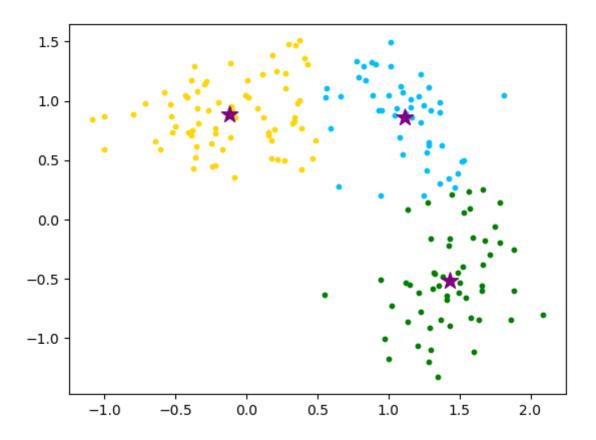
threhold	k	兰德系数RI	轮廓系数SI
0.6261045429970125	2	0.8362216720624643	0.5199313736871968
0.7291660883240724	3	0.9191899955564019	0.505556006687245
0.785168030660275	4	0.9318225099980956	0.4407097834539247
0.838108938465557	5	0.9191899955564019	0.4007694696269683
0.8793872709770619	6	0.9191899955564019	0.36940825202038036
0.9140906731626023	7	0.9264901923443154	0.3475953691837858
0.9210998493733723	8	0.9348695486573986	0.3376958556676636
0.9308947405600075	9	0.9348695486573986	0.3266843483029067
0.9497886943727526	10	0.9348695486573986	0.3171869407121694
0.9677133169251664	11	0.9348695486573986	0.3038264436279519
0.9846275862386208	12	0.9348695486573986	0.3038264436279519
1.0	13	0.9348695486573986	0.3008938518500136

进行PCA处理

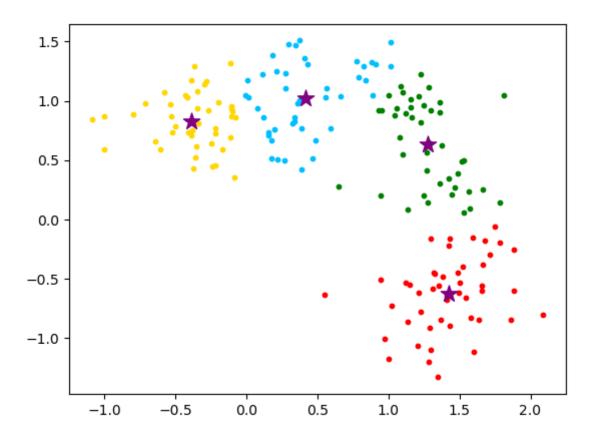
为便于可视化,使用PCA降低至二维 经过PCA后分为2个簇结果如图



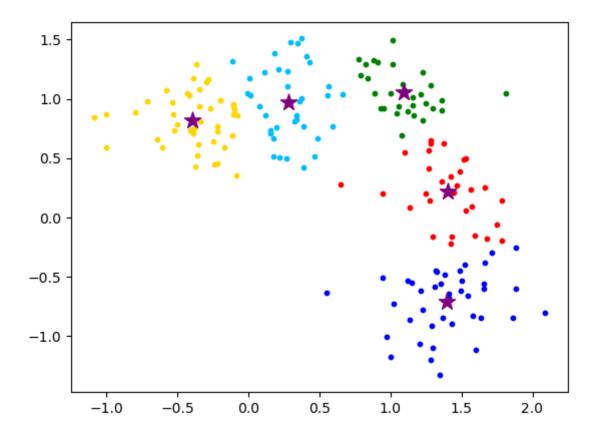
Rand Index:0.6810131403542182 Silhouette Coefficient:0.5405649074932503



Rand Index:0.8362216720624643 Silhouette Coefficient:0.5199313736871968



Rand Index:0.816035040944582 Silhouette Coefficient:0.4488521926995579



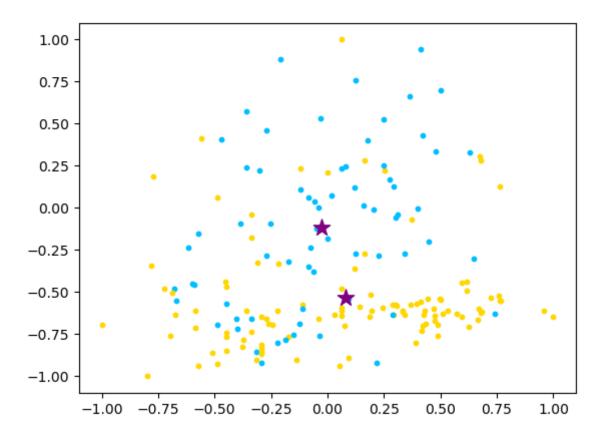
Rand Index:0.7769948581222624

Silhouette Coefficient: 0.43815734585649574

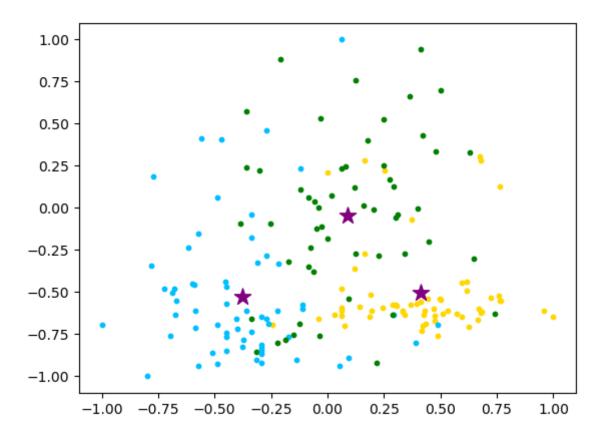
不同簇数下的Si和Ri

k	兰德系数RI	轮廓系数SI
2	0.6810131403542182	0.5405649074932503
3	0.8362216720624643	0.5199313736871968
4	0.816035040944582	0.4488521926995579
5	0.7769948581222624	0.43815734585649574

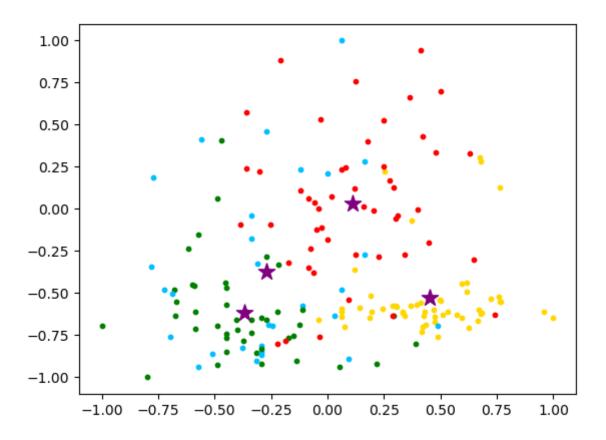
不进行PCA处理



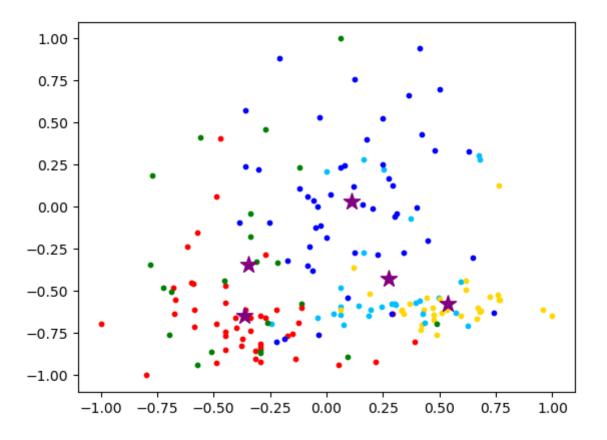
Rand Index:0.6810131403542182 Silhouette Coefficient:0.2987221815974774



Rand Index:0.9348695486573986 Silhouette Coefficient:0.30089385185001355



Rand Index:0.8950041261981845 Silhouette Coefficient:0.25995492195881664



Rand Index:0.8555195835713832 Silhouette Coefficient:0.18794414508559015

k	兰德系数RI	轮廓系数SI
2	0.6810131403542182	0.2987221815974774
3	0.9348695486573986	0.30089385185001355
4	0.8950041261981845	0.25995492195881664
5	0.8555195835713832	0.18794414508559015

可以看出,不进行PCA降维的话得到的图只能反应某几个维度的结果,看起来就非常零散,图像无法作为参考。而考察其兰德系数和轮廓系数,兰德系数要高于PCA处理后的结果,而轮廓系数要低于处理后的结果,这说明不经过PCA处理可以使得结果与真实结果的相似度更高因为其保留的信息更多,但同时分类轮廓也更不明显。