

# 实验五：层次聚类

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## 实验要求

### 基本要求

a) 实现single-linkage层次聚类算法； b) 实现complete-linkage层次聚类算法；

### 中级要求

a) 实现average-linkage层次聚类算法； b) 将上述三种算法的性能进行简要对比；

### 高级要求

通过变换聚类簇的个数，测试上述三种算法的性能，并给出分析。

## 1. 生成数据集

数据包含2000个样例，每个样例的前3列表示特征，第4列表示标签。

```
In [1]: from matplotlib import pyplot as plt
import numpy as np
from sklearn.datasets import make_blobs
from sklearn.metrics.cluster import adjusted_rand_score
def create_data(centers, num=100, std=0.7):
    """
    生成用于聚类的数据集
    :param centers: 聚类的中心点组成的数组。如果中心点是二维的，则产生的每个样本都是
    :param num: 样本数
    :param std: 每个簇中样本的标准差
    :return: 用于聚类的数据集。是一个元组，第一个元素为样本集，第二个元素为样本集的真
    """
    X, labels_true = make_blobs(n_samples=num, centers=centers, cluster_std=std)
    return X, labels_true

centers=[[1, 1, 1], [1, 3, 3], [3, 6, 5], [2, 6, 8]]# 用于产生聚类的中心点，聚类中心的维度代表产
X, labels_true= create_data(centers, 2000, 0.5) # 产生用于聚类的数据集，聚类中心点的个数
np.savetxt('D:/Program Files/JupyterSpace/ML/lab5/data.dat', X)
np.savetxt('D:/Program Files/JupyterSpace/ML/lab5/label.dat', labels_true)
print("generate data finish!")

generate data finish!
```

## 2. 初级&中级要求：实现single-linkage、complete-linkage、average-linkage层次聚类算法

```
In [2]: import numpy as np
```

```

MAX_NUM = 1e3

# method
def singleLinkage(X):
    # your code
    Minres = []
    for i in range(len(X[0])):
        Minres.append(MAX_NUM)
    for i in range(len(X[0])):
        if X[0][i] < Minres[i]:
            Minres[i] = X[0][i]
        if X[1][i] < Minres[i]:
            Minres[i] = X[1][i];
    return Minres

def completeLinkage(X):
    # your code
    Maxres = []
    for i in range(len(X[0])):
        Maxres.append(0)
    for i in range(len(X[0])):
        if X[0][i] > Maxres[i]:
            Maxres[i] = X[0][i]
        if X[1][i] > Maxres[i]:
            Maxres[i] = X[1][i];
    return Maxres

def averageLinkage(setList, dest, src, allDist):
    # your code
    Avgres = []
    for clu in range(len(setList)):
        if clu == dest or clu == src:
            Avgres.append(0)
            continue
        sumdistance = 0
        for other_cluster_element in setList[clu]:
            for dest_cluster_element in setList[dest]:
                sumdistance += allDist[dest_cluster_element][other_cluster_element]
            for src_cluster_element in setList[src]:
                sumdistance += allDist[src_cluster_element][other_cluster_element]
        res = sumdistance / ((len(setList[dest]) + len(setList[src])) * len(setList[clu]))
        Avgres.append(res)
    return Avgres

class AgglomerativeClustering:
    def __init__(self):
        # 对每次的合并进行记录
        self.steps = []

    def fit(self, datas, method):
        self.dataCnt = datas.shape[0] # dataCnt为总数据条数
        # 预处理各点之间的距离
        allDist = np.zeros((self.dataCnt, self.dataCnt)) # allDist为各点之间的距离
        for i in range(self.dataCnt):
            for j in range(i):
                allDist[i][j] = allDist[j][i] = np.sum((datas[i] - datas[j])**2)
        setList, clusterCount = [[i] for i in range(self.dataCnt)], self.dataCnt
        # setList为这一类有哪些数据, clusterCount为聚类个数(初始化为数据条数)
        print("calculate distance finish!")

        # 聚类间距离矩阵
        clusterDist = np.zeros((self.dataCnt, self.dataCnt)) + MAX_NUM # clusterDist为聚
        for i in range(clusterCount):
            for j in range(i+1, clusterCount):

```

```

        clusterDist[i][j] = clusterDist[j][i] = allDist[i][j]
    print("calculate cluster distance finish!")

    while clusterCount != 1: # 反复聚类直到只剩下一类
        # 最相似的两个聚类
        res = np.argmin(clusterDist) # res为距离最短的两类(i类和j类)的下标(i, j)
        dest, src = int(res/clusterCount), res%clusterCount # dest为i类, src为j类
        # steps进行一次记录
        self.steps.append((setList[dest][0], setList[src][0])) # steps用于记录每次
        # 聚类间距离矩阵更新
        if (method == averageLinkage):
            modify = method(setList, dest, src, allDist)
        else:
            modify = method(clusterDist[[dest, src]])
        clusterDist[dest] = modify # 所有其他类到新类的距离(一排)
        clusterDist[:, dest] = modify # 所有其他类到新类的距离(一列)
        clusterDist = np.delete(clusterDist, src, axis=0)
        clusterDist = np.delete(clusterDist, src, axis=1)
        clusterDist[dest][dest] = MAX_NUM
        # 聚类更新
        setList[dest] = setList[dest] + setList[src]
        del setList[src]
        clusterCount -= 1 # 总类数-1
        if (self.dataCnt - clusterCount) % (self.dataCnt / 20) == 0:
            print(clusterCount, " clusters left.") # 打印还剩多少类
    print("cluster finish !")

    def label(self, k):
        root = list(range(self.dataCnt))
        def find_root(n):
            if root[root[n]] == root[n]:
                return root[n]
            root[n] = find_root(root[n])
            return root[n]
        for i in range(self.dataCnt-k): # 根据steps记录产生非连通图
            src, dest = self.steps[i]
            root[find_root(dest)] = find_root(src)
        cluster, clusterNum = [0 for i in range(self.dataCnt)], 0
        for i in range(self.dataCnt): # 将根节点标注为新的cluster
            if i == root[i]: # i是根
                clusterNum += 1
                cluster[i] = clusterNum
        for i in range(self.dataCnt): # 将非根节点标注为根节点的cluster
            if i != root[i]: # i不是根
                cluster[i] = cluster[find_root(i)]
        return cluster

```

```

In [3]: from itertools import permutations
def accuracy_cal(labels_true, labels_predict, k):
    labels_resuffle = [0 for i in range(len(labels_true))]
    combination = list(permutations([i for i in range(1, k+1)], k))
    acclis = []
    for i in range(len(combination)): # 每个组合都试一下, 算出acc
        comindex = 0
        for j in range(1, k+1): # 修改每类的label
            for index in range(len(labels_true)): # 把所有label为j的都改成combination
                if labels_true[index] == j:
                    labels_resuffle[index] = combination[i][comindex]
                comindex += 1
        # 计算acc
        acc = 0
        for j in range(len(labels_resuffle)):
            if (labels_resuffle[j] == labels_predict[j]):

```

```

        acc += 1
        acclis.append(acc/len(labels_reshuffle))
#     print("acclis is :{0}".format(acclis))
    return max(acclis)

```

```

In [4]: def plot_data(*data):
        ,,,
        绘制用于聚类的数据集
        :param data: 可变参数。它是一个元组。元组元素依次为：第一个元素为样本集，第二个元
        :return: None
        ,,,

        X, labels_true, labels_predict, cnt=data
        fig=plt.figure()
        ax=fig.add_subplot(1,1,1)
        colors='rgbyckm' # 每个簇的样本标记不同的颜色
        markers='o^sP*DX'
        for i in range(len(labels_true)):
            predict=labels_predict[i]
            ax.scatter(X[i,0],X[i,1],label="cluster %d"%labels_true[i],
                color=colors[predict%len(colors)],marker=markers[labels_true[i]%len(markers)])

        METHOD_APPLY = [singleLinkage,completeLinkage,averageLinkage]

        cnt=0
        for method in METHOD_APPLY:
            model = AgglomerativeClustering()
            model.fit(X,method)
            k=4
            labels_predict = model.label(k)
            acc = adjusted_rand_score(labels_predict, labels_true)
            print("accuracy for {0} is:{1}".format(str(method),acc))
            plot_data(X,labels_true,labels_predict,cnt)
            cnt+=1
            print("-----Segmentation-----")

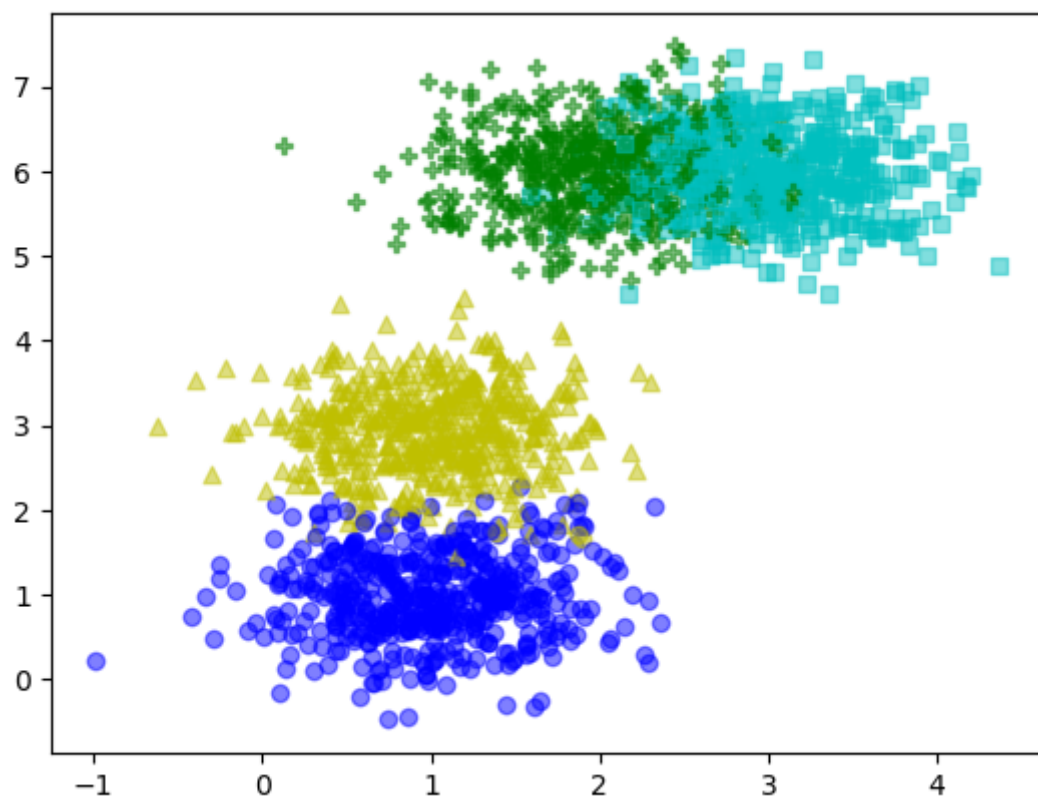
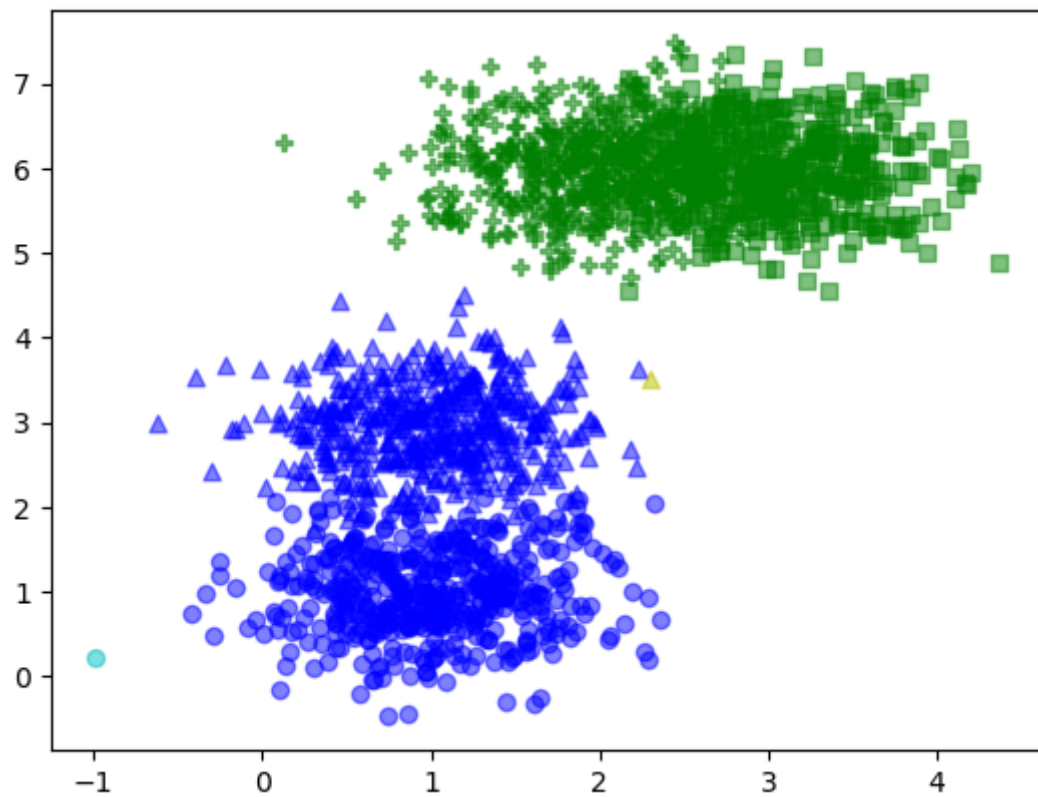
```

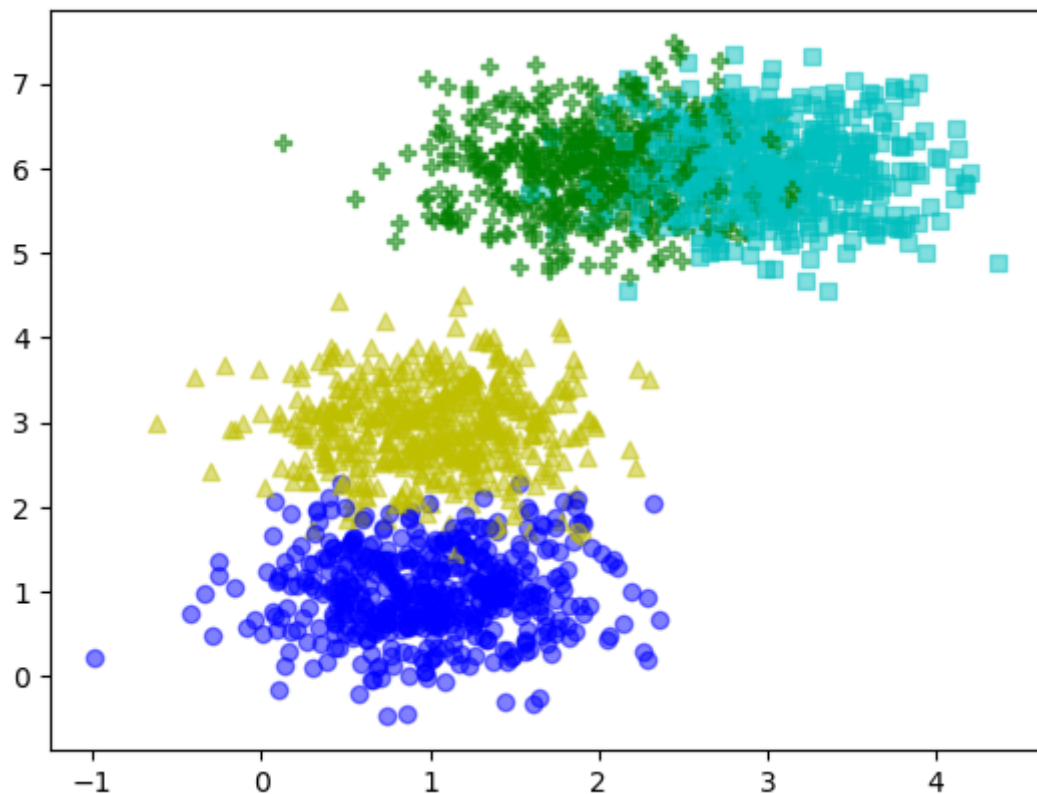
```

calculate distance finish!
calculate cluster distance finish!
1900 clusters left.
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cluster finish !
accuracy for <function singleLinkage at 0x000001D325121EE0> is:0.4991242803198932
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cluster finish !
accuracy for <function completeLinkage at 0x000001D329915EE0> is:0.9906915190011604
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cluster finish !  
accuracy for <function averageLinkage at 0x000001D329915F70> is:0.9933460896657262  
-----Segmentation-----





### 3. 中级要求：性能比较

列出三种算法的accuracy表格如下：

方法	accuracy
singleLinkage	0.4991
completeLinkage	0.9907
averageLinkage	0.9762

可以看出，singleLinkage算法最差，从聚类图中也可看出它常常将大部分数据分为2类或3类，第3/4类只有极少量的点；completeLinkage和averageLinkage算法明显表现很好。原因解释如下：

- singleLinkage容易造成两个聚类明明从整体上看相隔很远，但由于其中个别点的距离较近就被合并了，且合并后这种情况会恶化，类似链式效应，这也是为什么singleLinkage表现出容易合并两个明显不同但存在相交的类。
- 上述两种聚类方法的共同问题就是只考虑了某个极端的属性(如最近距离和最远距离)，而没有考虑类内数据的整体特点。averageLinkage把两个集合中的点两两距离全部放在一起求平均值，考虑了所有点对，更全面，**在改变方差进行实验时会发现averageLinkage相较completeLinkage更稳健。**

### 4. 高级要求

```
In [7]: acclis = [[], [], []]# 二维list，第二维表示不同方法，第一维表示不同clusternum
centerlis = [[1, 1, 1], [1, 3, 3], [3, 6, 5], [2, 6, 8], [2, 5, 6], [3, 4, 7], [4, 7, 8]]
for clunum in range(2, 8):
    centers=[centerlis[i] for i in range(clunum)]# 用于产生聚类的中心点，聚类中心的
    X, labels_true= create_data(centers, 2000, 0.5) # 产生用于聚类的数据集，聚类中心点的
    for methodindex in range(len(METHOD_APPLY)):
```

```
model = AgglomerativeClustering()
model.fit(X, METHOD_APPLY[methodindex])
labels_predict = model.label(clunum)
acclis[methodindex].append(adjusted_rand_score(labels_predict, labels_true))
# acclis[methodindex].append(accuracy_cal(labels_true, labels_predict, clunum)
# print("accuracy for {0} is: {1}, with clunum={2}".format(str(METHOD_APPLY[me
print("-----Segmentation-----")
```



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cluster finish !
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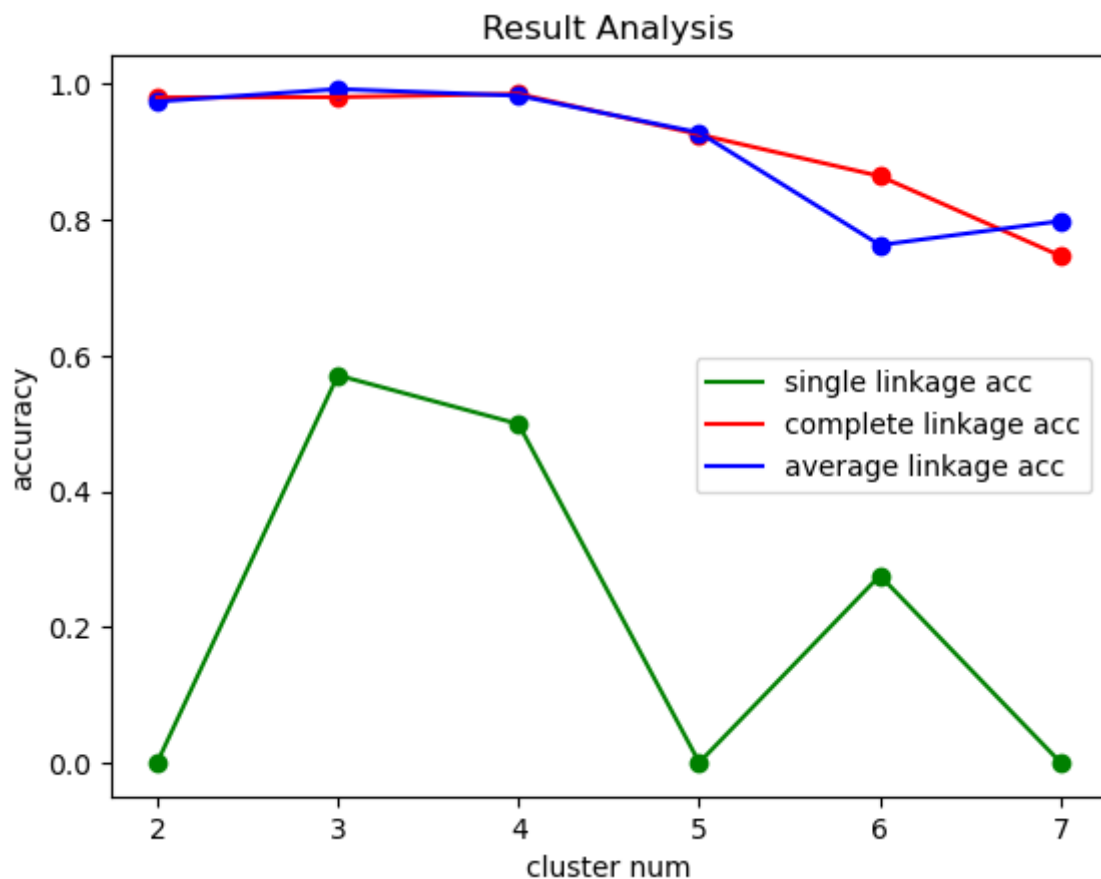
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```

In [8]: k = [i for i in range(2, 8)]
plt.title('Result Analysis')
plt.plot(k, acclis[0], color='green', label='single linkage acc')
plt.plot(k, acclis[1], color='red', label='complete linkage acc')
plt.plot(k, acclis[2], color='blue', label='average linkage acc')
plt.scatter(k, acclis[0], color='green')
plt.scatter(k, acclis[1], color='red')
plt.scatter(k, acclis[2], color='blue')
plt.legend() # 显示图例

plt.xlabel('cluster num')
plt.ylabel('accuracy')
plt.show()

```



- acc: 从acc曲线可以看出, complete linkage和average linkage效果远优于single linkage。average linkage和complete linkage不相上下, 但总体而言average linkage更加稳健。这是因为complete linkage虽然解决了single linkage中链式反应的问题, 但对离群点的处理能力不足, 而average linkage考虑了所有情况, 更加全面, 故效果最好。
- 时间复杂度: 值得一提的是, 运行复杂度方面average linkage高于另外两种算法, 这是因为它每次计算都要考虑全部的组合情况, 而single linkage和complete linkage只需找出两类中较小/大的那个distance即可。

因此, 模型选择时需要在时间和空间上折中。