03ClusteringTheory

November 4, 2018

1 分群 (Clustering)

- 1. 目標:
 - 群內相似度很高
 - 群間相似度很低
- 2. 主要影響因子:
 - 相似度的衡量與實作
 - •「群」的定義
 - 演算法
- 3. 應用面:
 - 市場區隔
 - 生物分群
 - 新聞分群

2 距離的定義

1. Euclidean distance(歐基里德距離)

•
$$d(i,j) = \sqrt{|x_{i1} - x_{j1}|^2 + |x_{i2} - x_{j2}|^2 + \dots + |x_{in} - x_{jn}|^2}$$

2. Manhattan (city block) distance(曼哈頓距離)

•
$$d(i,j) = |x_{i1} - x_{j1}| + |x_{i2} - x_{j2}| + \cdots + |x_{in} - x_{jn}|$$

3. Minkowski distance

•
$$d(i,j) = \sqrt[q]{|x_{i1} - x_{j1}|^q + |x_{i2} - x_{j2}|^q + \dots + |x_{in} - x_{jn}|^q}$$

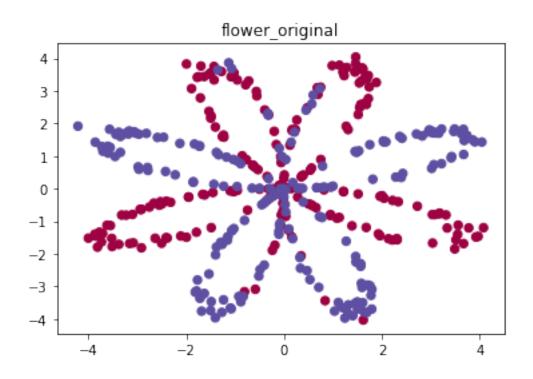
4. Weighted Manhattan distance(加權曼哈頓距離)

•
$$d(i,j) = \sqrt[q]{w1|x_{i1}-x_{j1}|^q+w2|x_{i2}-x_{j2}|^q+\cdots+w3|x_{in}-x_{jn}|^q}$$

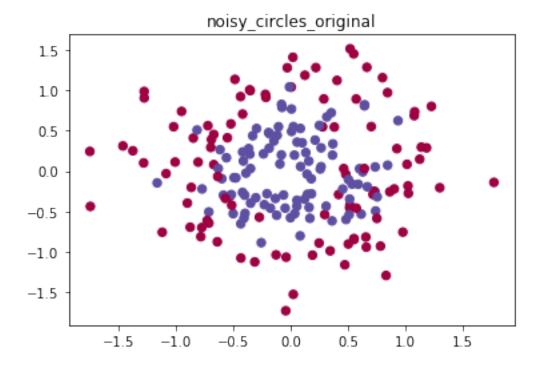
3 IMPORT & DATA

```
In [1]: import matplotlib.pyplot as plt
        from planar_utils import plot_decision_boundary, sigmoid, load_planar_dataset, load_ex
        import numpy as np
        np.random.seed(1212)
        import os
        from collections import Counter
        import pandas as pd
        from load_all_datasets import load
        from sklearn.cluster import KMeans, AgglomerativeClustering, DBSCAN
        from mpl_toolkits.mplot3d import Axes3D
        from sklearn import datasets
        from sklearn.decomposition import PCA
In [2]: # demo 用資料
       datas = load()
        for name, X, Y in datas:
            print(name)
           plt.scatter(X[:, 0], X[:, 1], c=Y, s=40, cmap=plt.cm.Spectral);
           plt.title(name+'_original')
           plt.show()
```

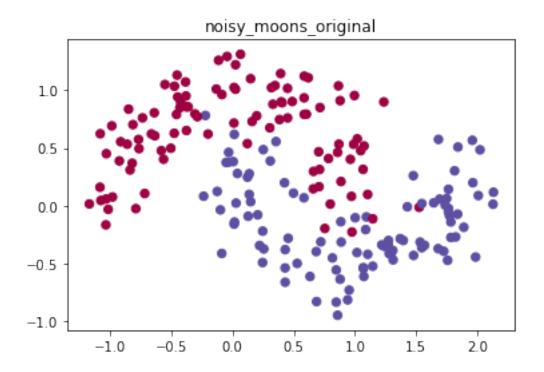
flower



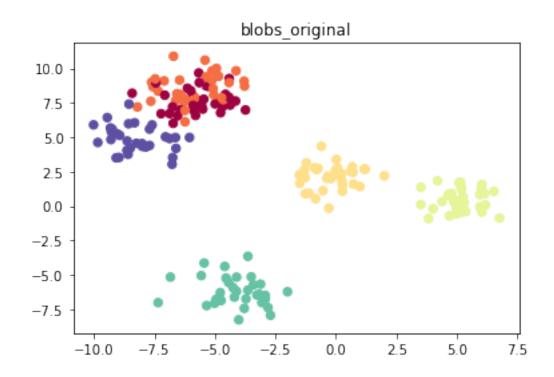
noisy_circles

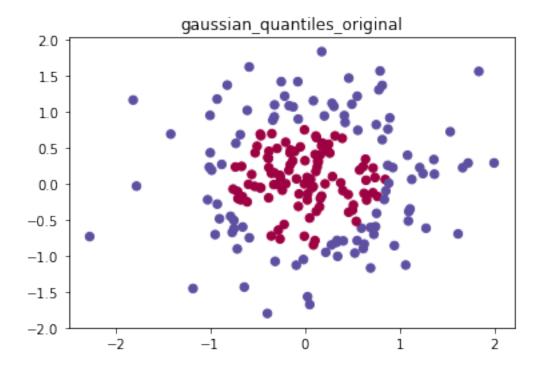


noisy_moons



blobs





```
In [3]: #練習用資料
        iris = datasets.load_iris()
        df_iris = pd.DataFrame(iris.data, columns=iris['feature_names'])
       df_iris['target'] = list(map(lambda x:iris.target_names[x], iris.target))
        df_iris.head()
Out[3]:
                              sepal width (cm)
                                                petal length (cm)
                                                                    petal width (cm)
           sepal length (cm)
        0
                         5.1
                                            3.5
                                                               1.4
                                                                                  0.2
        1
                         4.9
                                            3.0
                                                               1.4
                                                                                  0.2
        2
                         4.7
                                            3.2
                                                               1.3
                                                                                  0.2
        3
                         4.6
                                            3.1
                                                               1.5
                                                                                  0.2
        4
                         5.0
                                                               1.4
                                            3.6
                                                                                  0.2
           target
           setosa
        1
           setosa
        2
           setosa
        3
           setosa
           setosa
```

In [4]: # 簡單資料探索 # 請找出每一個 target (花的類別) 的種類各有幾筆資料 (hint: groupby)

```
#!========!#
df_iris.groupby('target').count()
#!========!#
```

```
Out[4]:
                    sepal length (cm)
                                       sepal width (cm) petal length (cm) \
        target
        setosa
                                    50
                                                      50
                                                                          50
        versicolor
                                    50
                                                      50
                                                                          50
        virginica
                                    50
                                                      50
                                                                          50
                    petal width (cm)
```

target setosa 50 versicolor 50 virginica 50

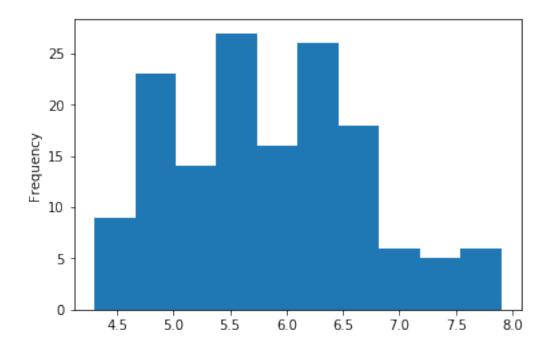
plt.show()

```
In [5]: # 請畫出 ['sepal length (cm)'] 的分布狀況

#!=======your works starts======!#

df_iris['sepal length (cm)'].plot('hist')

#!======your works ends=====!#
```



```
Y = iris_Y
# 請使用 PCA 畫出不同花在三維空間中的分布位置 (請上網尋找 PCA 的程式碼·並複製過來)
#!=======your works starts=======!#
fig = plt.figure(1, figsize=(8, 6))
ax = Axes3D(fig, elev=-150, azim=110)
X_reduced = PCA(n_components=3).fit_transform(X)
ax.scatter(X_reduced[:, 0], X_reduced[:, 1], X_reduced[:, 2], c=Y,
          cmap=plt.cm.Set1, edgecolor='k', s=40)
ax.set_title("PCA with non-log(Fare)")
ax.set_xlabel("1st eigenvector")
ax.w_xaxis.set_ticklabels([])
ax.set_ylabel("2nd eigenvector")
ax.w_yaxis.set_ticklabels([])
ax.set_zlabel("3rd eigenvector")
ax.w_zaxis.set_ticklabels([])
#!===========!#
plt.show()
```

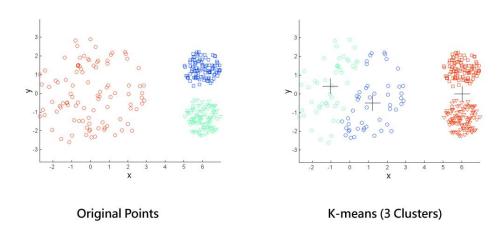
4 ALGORITHM

4.1 1. K-means

4.1.1 steps

- 1. 設定重心:
 - 決定要分幾個群 (e.g. K 群)
 - 隨機產生 K 個重心
- 2. 分群:
 - 找到所有資料點歸屬的重心: 找到分別最靠近這三個點的所有資料點,並做第一個迴圈 的分群
- 3. 尋找重心:
 - 重新計算所有資料點的重心 (element-wise 的平均)
- 4.
- 5. 停止:
 - 重複上述方法,只到不會有資料點因為重新尋找重心而改變群組

kmeans



KmeansCon1.JPG

4.1.2 visualize

1. web:

game

2.

4.1.3 parameters

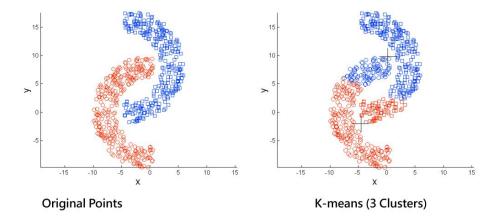
參數 意義

n_cluster 要分成幾個群

random_state 決定初始點的隨機 seed

4.1.4 pros & cons

- 1. 一開始需要決定要分成幾個群
- 2. 優點
 - 效率: 時間複雜度 O(tkn)。n 是資料點的個數 · k 是群的個數 · t 是迴圈的個數 · 班來 說: k, t << n 。
- 3. 缺點
 - 需要指定分成幾群
 - 無法處理具有雜訊 (noisy) 的資料及離群值 (outliers)
 - 無法處理不同大小及不同密度的群
 - 無法發現凸多邊形 (non-convex) 以外的群



KmeansCon2.JPG

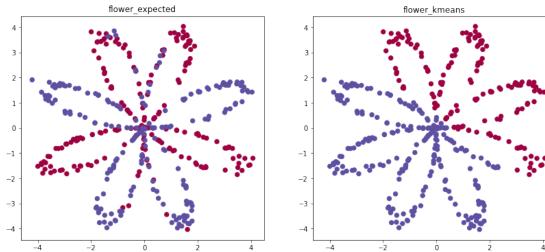
• 通常在 local optimum 停止

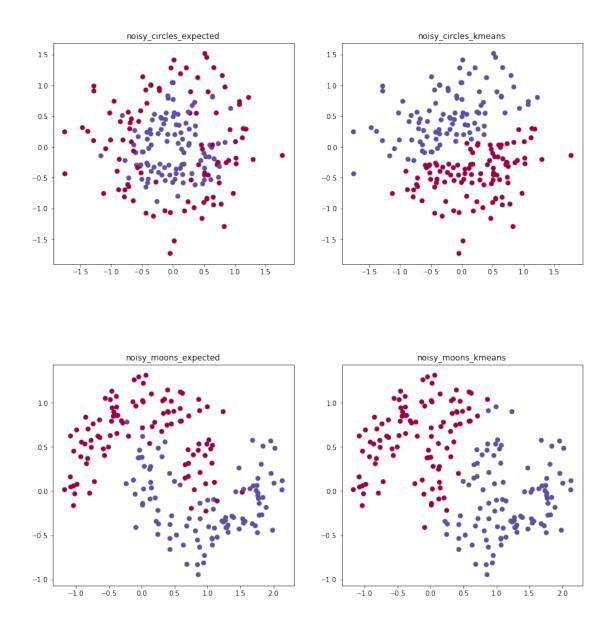
```
In [7]: for name, X, Y in datas:
    plt.figure(figsize=(14,6))

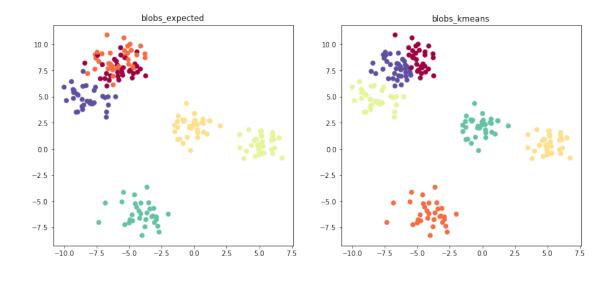
axes1 = plt.subplot(1, 2, 1)
    axes1.scatter(X[:, 0], X[:, 1], c=Y, s=40, cmap=plt.cm.Spectral)
    plt.title(name + "_expected")

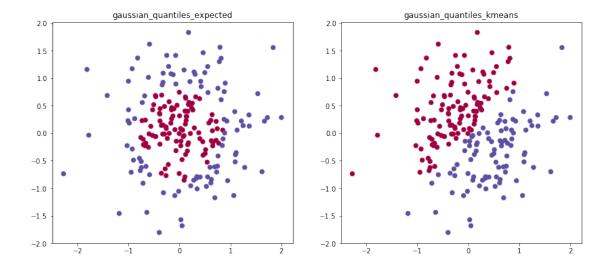
axes2 = plt.subplot(1, 2, 2)
    n_clusters = len(set(Y))
    kmeans = KMeans(n_clusters=n_clusters, random_state=0).fit_predict(X)
    axes2.scatter(X[:, 0], X[:, 1], c=kmeans, s=40, cmap=plt.cm.Spectral);
    plt.title(name+'_kmeans')

plt.show()
```









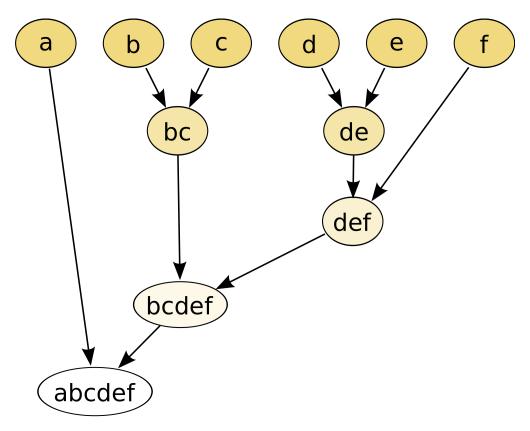
```
In [8]: # 請使用 kmeans 將 iris_X 進行分群 (注意:random_state=0)
#========your works starts=======#

n_clusters = len(set(iris_Y))
kmeans_Y = KMeans(n_clusters=n_clusters, random_state=0).fit_predict(iris_X)
#========your works ends=======#

np.random.seed(1212)
print(kmeans_Y[np.random.randint(149, size=10)])
# # [2 0 0 2 2 1 1 1 1 0]
```

[2 0 0 2 2 1 1 1 1 0]

```
In [9]: # actual
        def plot_iris_predict_acc(predict_Y):
            X = iris_X
            Y = iris_Y
            fig = plt.figure(1, figsize=(16, 6))
            ax = fig.add_subplot(1, 2, 1, projection='3d')
            X reduced = PCA(n components=3).fit transform(X)
            ax.scatter(X_reduced[:, 0], X_reduced[:, 1], X_reduced[:, 2], c=Y,
                       cmap=plt.cm.Set1, edgecolor='k', s=40)
            ax.set_title("PCA with non-log(Fare)")
            ax.set_xlabel("1st eigenvector")
            ax.w_xaxis.set_ticklabels([])
            ax.set_ylabel("2nd eigenvector")
            ax.w_yaxis.set_ticklabels([])
            ax.set_zlabel("3rd eigenvector")
            ax.w_zaxis.set_ticklabels([])
            # predict
            X = iris_X
            Y = predict Y
            fig = plt.figure(1, figsize=(16, 6))
            ax = fig.add_subplot(1, 2, 2, projection='3d')
            X_reduced = PCA(n_components=3).fit_transform(X)
            ax.scatter(X_reduced[:, 0], X_reduced[:, 1], X_reduced[:, 2], c=Y,
                       cmap=plt.cm.Set1, edgecolor='k', s=40)
            ax.set_title("PCA with non-log(Fare)")
            ax.set_xlabel("1st eigenvector")
            ax.w_xaxis.set_ticklabels([])
            ax.set_ylabel("2nd eigenvector")
            ax.w_yaxis.set_ticklabels([])
            ax.set_zlabel("3rd eigenvector")
            ax.w_zaxis.set_ticklabels([])
            plt.show()
        np.random.seed(1212)
        plot_iris_predict_acc(kmeans_Y)
```



hierarchical

4.2 2. Hierarchical Clustering

4.2.1 steps

hierarchical 的分群方法,步驟簡述如下,可參考下圖 (取自維基百科):

- 1. 把每一個點當作一個群組
- 2. 透過掃描過整個資料及尋找出最近的兩個群組,並把這兩個點榜再一起變成一個群組
- 3. 尋找下一個最近的的兩個群組,再綁再一起變成一個群組
- 4.
- 5. 停止: 直到所有資料都被分成一群,或是透過設定參數到分到幾個群時自動停止

4.2.2 parameters

1. table

參數 意義

linkage 如何衡量群與群之間的距離。(註解) n_clusters 分成幾個群

13

2. 註解:

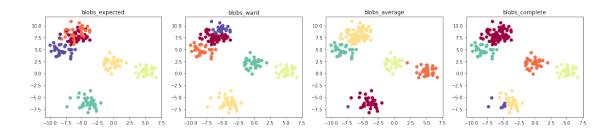
• ward(single): 兩個群中最近的點。

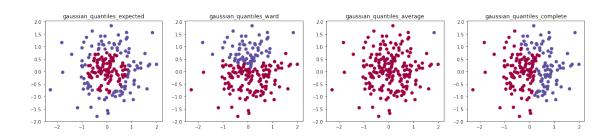
• complete: 兩個群中最遠的點。

● average: 兩個群的重心。

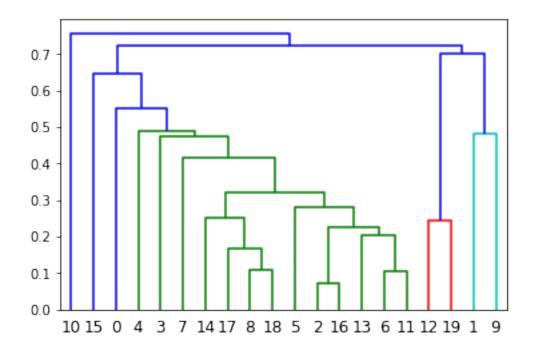
4.2.3 pros & cons

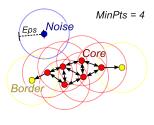
```
axes3 = plt.subplot(1, 4, 3)
      n_clusters = len(set(Y))
      hierarchy = AgglomerativeClustering(linkage=linkage[1], n_clusters=n_clusters).fi
      axes3.scatter(X[:, 0], X[:, 1], c=hierarchy, s=40, cmap=plt.cm.Spectral);
      plt.title(name+'_' + linkage[1])
      axes4 = plt.subplot(1, 4, 4)
      n_clusters = len(set(Y))
      hierarchy = AgglomerativeClustering(linkage=linkage[2], n_clusters=n_clusters).fi
      axes4.scatter(X[:, 0], X[:, 1], c=hierarchy, s=40, cmap=plt.cm.Spectral);
      plt.title(name+'_' + linkage[2])
      plt.show()
                                         1.0
                                         -1.0
                                         -1.5
-1.5 -1.0 -0.5 0.0 0.5 1.0 1.5
                      -1.5 -1.0 -0.5 0.0 0.5 1.0 1.5
                                             -1.5 -1.0 -0.5 0.0 0.5 1.0 1.5
                                                                   -1.5 -1.0 -0.5 0.0 0.5 1.0 1.5
                      -1.0 -0.5 0.0 0.5 1.0
                                   1.5 2.0
                                            -1.0 -0.5 0.0 0.5 1.0
                                                                   -1.0 -0.5 0.0 0.5 1.0
```





In [11]: # Other Method (參考就好)
 from scipy.cluster.hierarchy import dendrogram, linkage
 Z = linkage(X[:20], 'single')
 plt.figure()
 dn = dendrogram(Z)
 plt.show()





DBSCAN

In [12]: # 請用 AgglomerativeClustering(linkage='average') 將 iris_X 進行分群

#=======your works starts======#

n_clusters = len(set(iris_Y))

hierarchy_Y = AgglomerativeClustering(linkage='average', n_clusters=n_clusters).fit_pressure #=======your works ends=======#

plot_iris_predict_acc(hierarchy_Y)





4.3 3. Density Based Clustering (DBSCAN)

4.3.1 steps

顧名思義,這種分群演算法計算的是密度,透過設定多長的半徑內,有出現幾個點,不斷延伸,延伸到無法延伸,所有出現在前面延伸範圍的點分成一個群組,請見下圖 (取自stackexchange)。

4.3.2 parameters

參數 意義 eps 半徑

axes1.scatter(X[:, 0], X[:, 1], c=Y, s=40, cmap=plt.cm.Spectral)

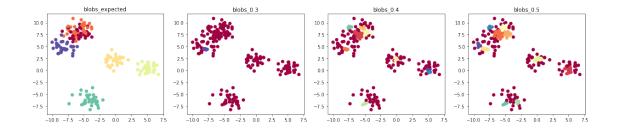
4.3.3 pros & cons

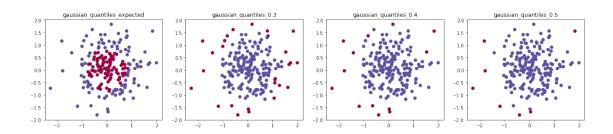
- 1. 常常用來偵測 Noisy Data。
- 2. 無法設定要分成幾個群。

```
axes2 = plt.subplot(1, 4, 2)
      dbscan = DBSCAN(eps=epss[0]).fit_predict(X)
      axes2.scatter(X[:, 0], X[:, 1], c=dbscan, s=40, cmap=plt.cm.Spectral)
      plt.title(name+'_'+str(epss[0]))
      axes3 = plt.subplot(1, 4, 3)
      dbscan = DBSCAN(eps=epss[1]).fit_predict(X)
      axes3.scatter(X[:, 0], X[:, 1], c=dbscan, s=40, cmap=plt.cm.Spectral)
      plt.title(name+'_'+str(epss[1]))
      axes3 = plt.subplot(1, 4, 4)
      dbscan = DBSCAN(eps=epss[2]).fit_predict(X)
      axes3.scatter(X[:, 0], X[:, 1], c=dbscan, s=40, cmap=plt.cm.Spectral)
      plt.title(name+'_'+str(epss[2]))
      plt.show()
                                       1.0
                                       -0.5
                                       -1.0
                                       -1.5
-1.5 -1.0 -0.5 0.0 0.5 1.0 1.5
                                                               -1.5 -1.0 -0.5 0.0 0.5 1.0 1.5
                    -1.0 -0.5 0.0 0.5 1.0 1.5 2.0
                                          -1.0 -0.5 0.0 0.5 1.0 1.5 2.0
```

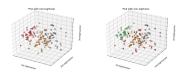
0.0 -0.5

-1.0 -1.5





plot_iris_predict_acc(dbscan_Y)



In []: