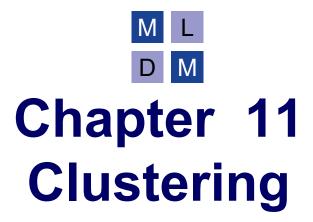
# MIMA Group



#### **Outline**

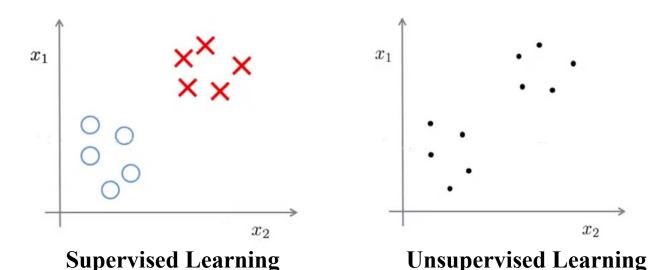


- Definition
- Distance Calculation
- Algorithms
  - K-means
  - Mixture of Gaussian
  - DBSCAN
  - AGNES
- Performance Measure

#### **Definition**



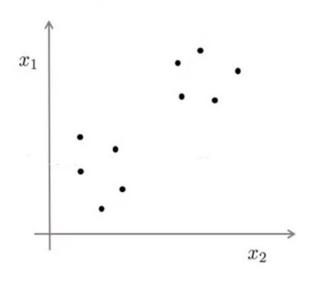
- Supervised learning VS. Unsupervised learning
- In supervised learning, we know something(label or value) about data( $X=\{x1,...,xn\}$ ,  $Y=\{y1,...,yn\}$ known), learn y=f(x)
- In unsupervised learning, we know nothing about the data(X known, Y unknown).

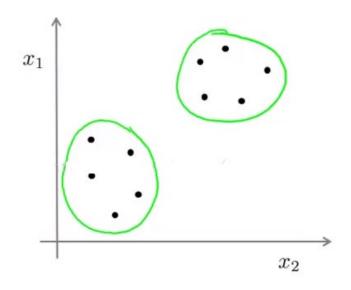


#### **Definition**



Clustering is the task of grouping a set of objects in such a way that objects in the same group(intragroup) are more similar to each other than to those in other groups(inter-group).





## **Distance Calculation**



Minkowski distance

$$dist_{mk}(x_i, x_j) = \left(\sum_{u=1}^{n} |x_{iu} - x_{ju}|^p\right)^{\frac{1}{p}}$$

$$x_i = (x_{i1}; x_{i2}; ... x_{in}), x_j = (x_{j1} x_{j2}; ... x_{jn})$$

- Euclidean distance (p=2)
- Manhattan distance (p=1)

$$x_1 = [2,1]$$

$$x_2 = [3,3]$$
Euclidean distance
$$d = (|2-3|^1 + |1-3|^1)^1 = 3$$

$$d = (|2-3|^2 + |1-3|^2)^{1/2} = \sqrt{5}$$

## Other Similarity Metrics



■ Chebyshev Distance D=max( $|x_1 - x_2|, |y_1 - y_2|$ )

D=max(
$$|x_1 - x_2|, |y_1 - y_2|$$
)

Cosine

$$\cos(\theta) = \frac{\sum_{i=1}^{n} (x_i * y_i)}{\sqrt{\sum_{i=1}^{n} (x_i)^2} * \sqrt{\sum_{i=1}^{n} (y_i)^2}}$$

- Hamming Distance
  - $d(x,y)=\sum X[i] \oplus Y[i]$
- Jaccard Distance

$$\frac{|A \cup B| - |A \cap B|}{|A \cup B|}$$

Correlation coefficient and Correlation distance

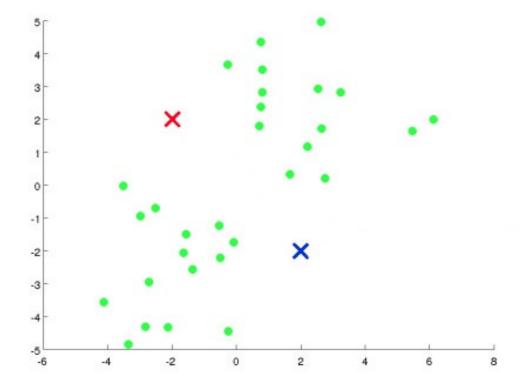
# **Algorithm--K-means**



- Randomly select K samples as the centroids of clustering.
- Calculate the distances of each sample from the K centroid points.
- Select the cluster centroid  $c_i$  (i = 1, 2 ... K) with the smallest distance to divide the cluster.
- Re-determine the centroids.
- Iterate until it converges.



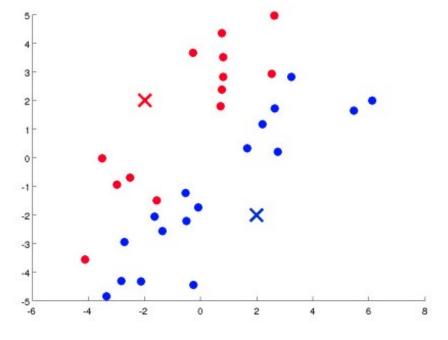
- By experience, set k=2.
- Randomly select two points as cluster centroids.





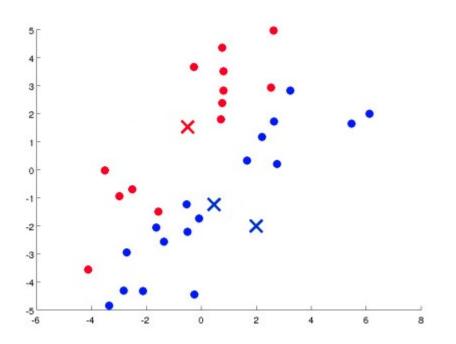
- For all green samples, compute distances from the blue point and red point.
- Choose the smallest distance and assign it to one cluster.

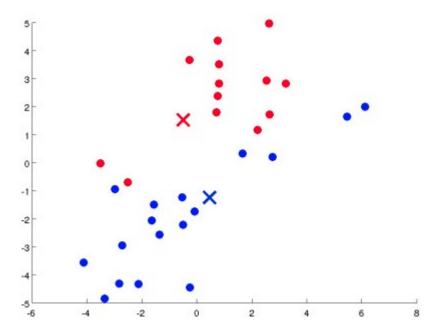
#### 簇划分





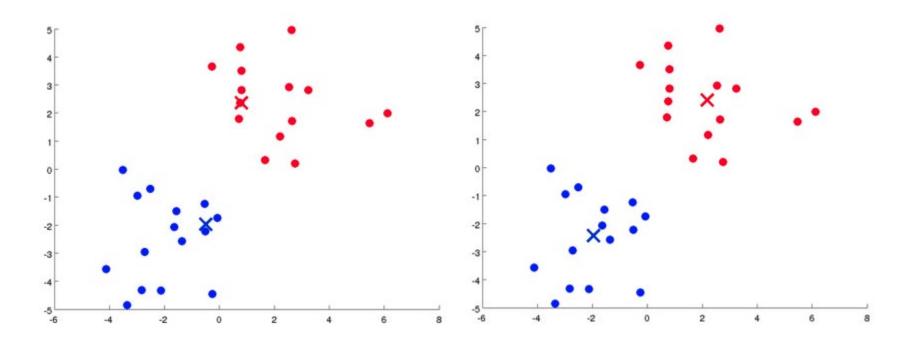
- Compute the average for two clusters.
- Re-determine cluster centroids.



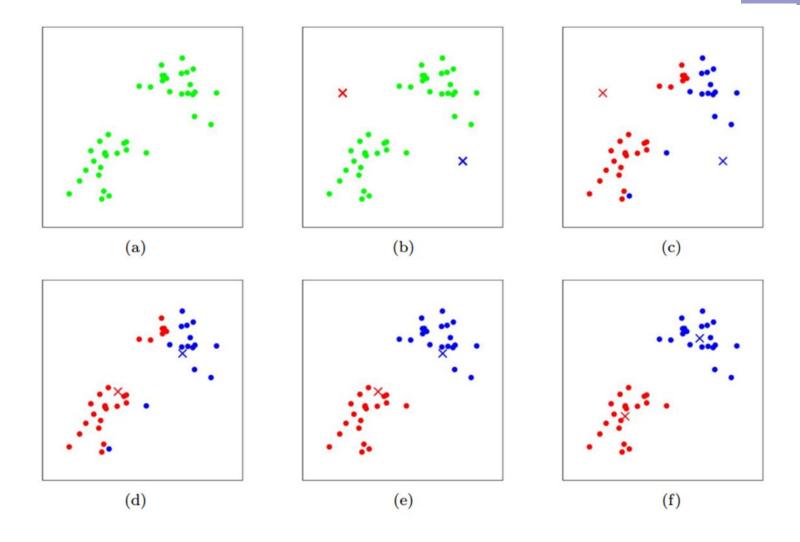




■ Iterate until convergence (the cluster centroids will not change).









```
输入: 样本集D = \{x_1, x_2, \dots, x_m\};
       聚类簇数k.
过程:
 1: 从D中随机选择k个样本作为初始均值向量{\mu_1, \mu_2
 2: repeat
    \diamondsuit C_i = \emptyset \ (1 \le i \le k)
      for j = 1, \ldots, m do
    计算样本x_j与各均值向量\mu_i (1 \le i \le k)的足根据距离最近的均值向量确定x_j的簇标记:
      将样本x_j划入相应的簇: C_{\lambda_j} = C_{\lambda_j} \bigcup \{x_j\};
      end for
      for i = 1, \ldots, k do
         计算新均值向量: \mu'_i = \frac{1}{|C_i|} \sum_{\boldsymbol{x} \in C_i} \boldsymbol{x};
10:
     if \mu_i' \neq \mu_i then
      将当前均值向量\mu_i更新为\mu'_i
12:
     else
13:
       保持当前均值向量不变
14:
         end if
15:
      end for
16:
17: until 当前均值向量均未更新
18: return 簇划分结果
```

**输出:** 簇划分 $\mathcal{C} = \{C_1, C_2, \ldots, C_k\}$ 

- Are the results with different random initialization same?
- How to choose K?

移动 聚类中心



- We need all possible initializations and get the best result.
- The measure to find the best result is minimizing square error E(SSE, sum of the Squared Error).

$$E = \sum_{i=1}^{k} \sum_{x \in C_i} ||x - u_i||_2^2 \qquad E = \sum_{i=1}^{m} ||x_i - u_{\lambda_i}||_2^2$$

$$u_i = \frac{1}{|C_i|} \sum_{x \in C_i} x$$

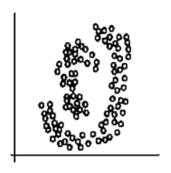


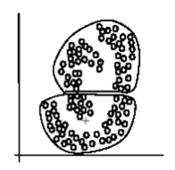
#### **□** How to initialize

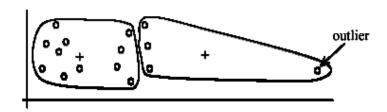
- ➤ It is NP-Hard to minimizing E.
- ➤ K-means uses an iterative optimal algorithm. Each step of every iteration is the process of optimizing E.
- We can choose multiple initializations to get the best result(Attention: Whether this measure is effective depends on k).



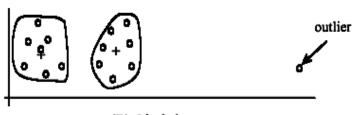
■ K-means is not always suitable.







(A): Undesirable clusters



(B): Ideal clusters



Gaussian mixture distribution

$$p(x) = \sum_{k=1}^{K} p(k)p(x|k)$$
$$= \sum_{k=1}^{K} \pi_k \mathcal{N}(x|\mu_k, \Sigma_k)$$

- $\blacksquare \pi_k$ : corresponding mixture coefficient



Objective function:

$$\sum_{i=1}^{N} \log \left\{ \sum_{k=1}^{K} \pi_k \mathcal{N}(x_i | \mu_k, \Sigma_k) 
ight\}$$

■ Then, we use MLE(Maximum likelihood estimate) to optimize function.



■ 估计数据由每个 Component 生成的概率 (并不是每个 Component 被选中的概率) : 对于每个数据 x\_i 来说, 它由第 k 个 Component 生成的概率为

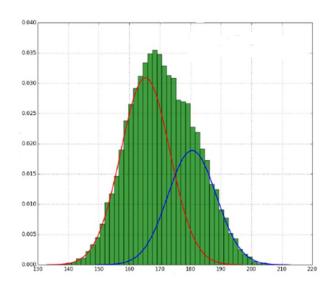
$$\gamma(i,k) = rac{\pi_k \mathcal{N}(x_i | \mu_k, \Sigma_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(x_i | \mu_j, \Sigma_j)}$$

■ 估计每个 Component 的参数

$$egin{aligned} \mu_k &= rac{1}{N_k} \sum_{i=1}^N \gamma(i,k) x_i \ \Sigma_k &= rac{1}{N_k} \sum_{i=1}^N \gamma(i,k) (x_i - \mu_k) (x_i - \mu_k)^T \end{aligned}$$



- Suppose we have 100 students and the only data we can get is their height. Try to model the distribution of male and female.
- The height obeys Gaussian distribution.





先假定男生服从参数为~N(180,10)的高斯分布,女生服从参数为~N(160,8)的高斯分布(Assume a determined Gaussian model for boys and girls)

- 对每个样本计算出分别属于男生和女生的概率(Compute the probability that clustering each sample to boys and girls)
- 认定:每个样本分属于男生和女生的部分(即概率,用 $\gamma(i,k)$ 表示,即第i个样本属于第k个类别的概率)同样服从高斯分布,且具有更好的拟合属性。(如一个样本身高175,我们可以通过设定的参数计算出他有80%的概率为男生,20%的概率为女生,那可以把这个样本看作由80%的男生和20%的女生组成,并将这个样本看作是一个80%的男生样本和一个20%的女生样本。)(Divide each sample into two components according to the probability in step 2)
- 根据每个样本的男生组分和女生组分拟合出新的高斯模型。(Update the parameters according to the components in step 3)
- 迭代直到收敛。(Iterating these steps until convergence)



```
输入: 样本集D = \{x_1, x_2, \dots, x_m\};
           高斯混合成分个数k.
 过程:
 1: 初始化高斯混合分布的模型参数\{(\alpha_i, \mu_i, \Sigma_i) \mid 1 \le i \le k\}
 2: repeat
         for j = 1, ..., m do
             根据(9.30)计算x_i由各混合成分生成的后验概率,即
            \gamma_{ii} = p_{\mathcal{M}}(z_i = i \mid \boldsymbol{x}_i) \ (1 \leq i \leq k)
         end for
         for i = 1, \ldots, k do
        计算新均值向量: \mu_i' = \frac{\sum_{j=1}^m \gamma_{ji} \boldsymbol{x}_j}{\sum_{j=1}^m \gamma_{ji}};
            计算新协方差矩阵: \Sigma_i' = \frac{\sum_{j=1}^m \gamma_{ji} (\boldsymbol{x}_j - \boldsymbol{\mu}_i') (\boldsymbol{x}_j - \boldsymbol{\mu}_i')^\top}{\sum_{j=1}^m \gamma_{ji}};
            计算新混合系数: \alpha_i' = \frac{\sum_{j=1}^m \gamma_{ji}}{m};
         end for
10:
         将模型参数\{(\alpha_i, \boldsymbol{\mu}_i, \boldsymbol{\Sigma}_i) \mid 1 \leq i \leq k\}更新为\{(\alpha'_i, \boldsymbol{\mu}'_i, \boldsymbol{\Sigma}'_i) \mid 1 \leq i \leq k\}
12: until 满足停止条件
13: C_i = \emptyset \ (1 < i < k)
14: for j = 1, ..., m do
         根据(9.31)确定x_i的簇标记\lambda_i;
         将x_i划入相应的簇: C_{\lambda_i} = C_{\lambda_i} \bigcup \{x_i\}
17: end for
18: return 簇划分结果
 输出: 簇划分\mathcal{C} = \{C_1, C_2, \dots, C_k\}
```

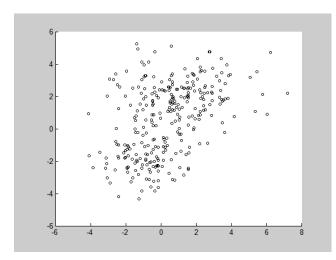
固定模型参数

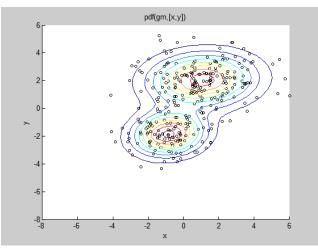
更新后验概率

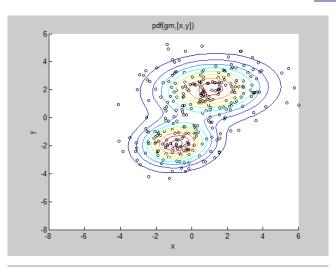
固定后验概率

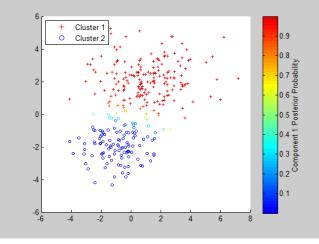
更新模型参数









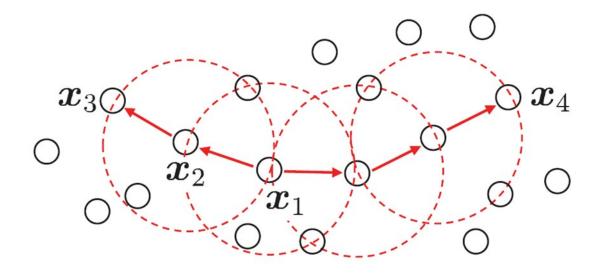




- > It is a famous density-based-clustering.
- $\succ$   $\epsilon$ -neighborhood: $N_{\epsilon}(x_j) = \{x_i \in D | dist(x_i, x_j) \le \epsilon\}$  of the point  $x_j$
- > core object :a point with a  $|N_{\epsilon}(x_j)| \ge MinPts$
- $\triangleright$  Directly density-reached:  $x_j$  is directly density-reachable from a core object  $x_i$  if  $x_j$  is in  $N_{\epsilon}(x_i)$
- Density-reached:  $x_j$  is density-reachable from a core object  $x_i$  if a sequence of core objects  $p_1, p_2, ..., p_n$  between  $x_i$  and  $x_j$  exists and  $p_{i+1}$  is directly density-reached from  $p_i$ .
- $\triangleright$  Density-connected:  $x_i$  and  $x_j$  are density-connected if they are density-reachable from a common core object  $x_k$ .

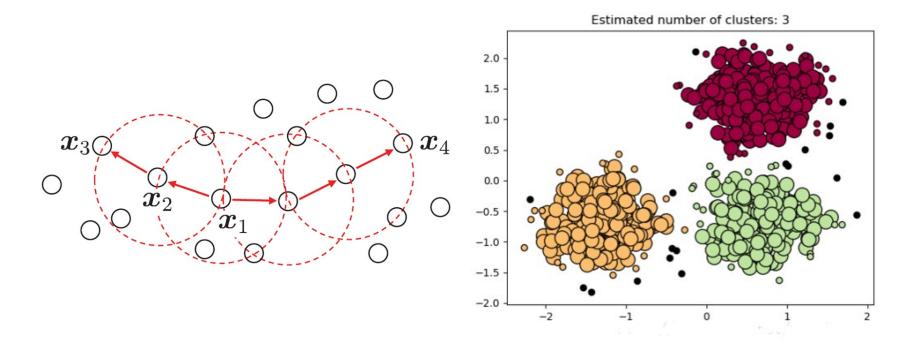


- □ Directly density-reached: X2 from X1
- □ Density-reached: X3 from X1
- □ Density-connected: X4 from X3





■ DBSCAN defines cluster as such sample set which is most density-connected.





找出所有核心 对象

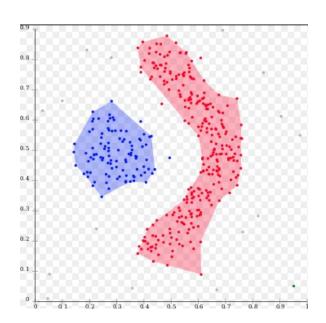
随机选一个 核心对象生长 出一个簇,并 在核心对象集 合里删去该核 心对象

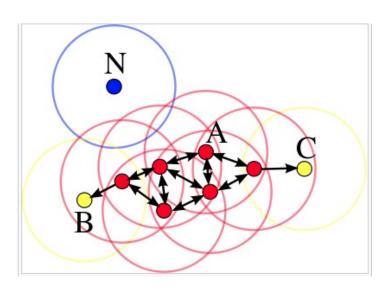
```
输入: 样本集D = \{x_1, x_2, \dots, x_m\};
        邻域参数(\epsilon, MinPts).
过程:
 1: 初始化核心对象集合: \Omega = \emptyset
 2: for j = 1, ..., m do
 3: 确定样本x_i的\epsilon-邻域N_{\epsilon}(x_i);
 4: if |N_{\epsilon}(\boldsymbol{x}_{j})| \geq MinPts then
     将样本x_i加入核心对象集合: \Omega = \Omega \cup \{x_i\}
       end if
 7: end for
 8: 初始化聚类簇数: k=0
 9: 初始化未访问样本集合: \Gamma = D
10: while \Omega \neq \emptyset do
     记录当前未访问样本集合: \Gamma_{\text{old}} = \Gamma;
     随机选取一个核心对象o \in \Omega, 初始化队列 Q = \langle o \rangle;
      \Gamma = \Gamma \setminus \{o\};
13:
      while Q \neq \emptyset do
          取出队列Q中的首个样本q;
15:
         if |N_{\epsilon}(q)| \geq MinPts then
16:
         \diamondsuit \Delta = N_{\epsilon}(\mathbf{q}) \cap \Gamma;
17:
       将\Delta中的样本加入队列Q;
18:
            \Gamma = \Gamma \setminus \Delta;
19:
     end if
20:
     end while
     k = k + 1, 生成聚类簇C_k = \Gamma_{\text{old}} \setminus \Gamma;
      \Omega = \Omega \setminus C_k
23:
24: end while
25: return 簇划分结果
输出: 簇划分\mathcal{C} = \{C_1, C_2, \dots, C_k\}
```



#### Strengths

- > There is no K.
- > It can discover any shape of spatial clustering.
- > It can discard the remote point.



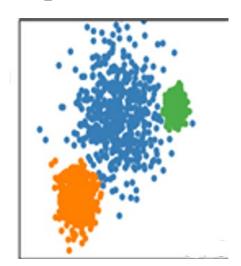




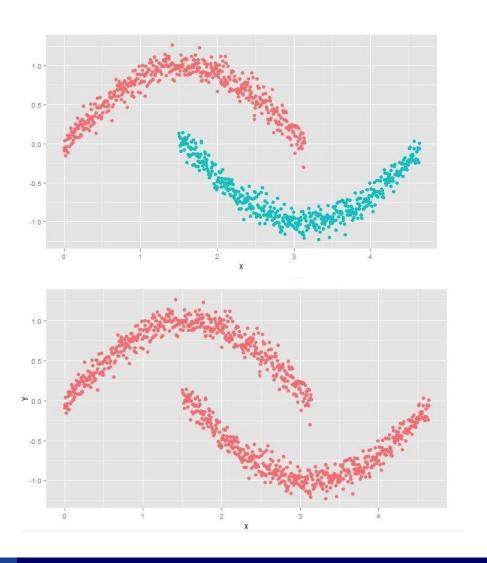
#### Weaknesses

- ➤ It is not suitable when the cluster spacing difference is very different.
- > Parameters adjustment are more complicated.



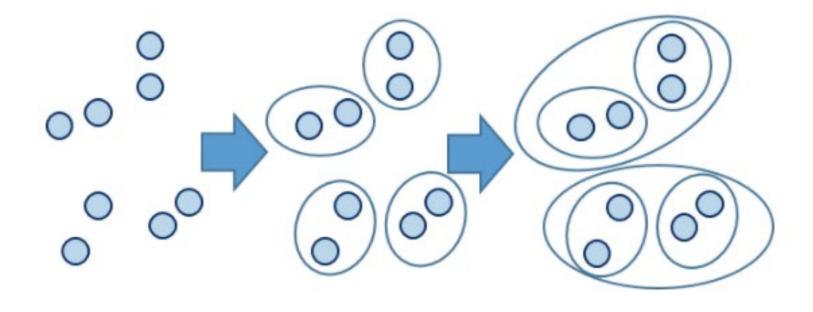




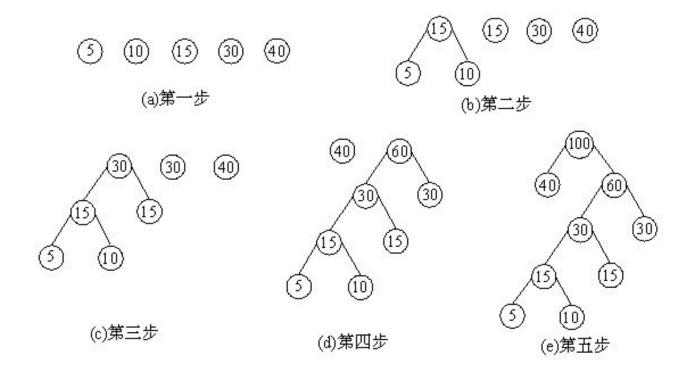




AGNES is a kind of Hierarchical clustering









- Given cluster  $C_i$ ,  $C_j$ , Usually the distance between two clusters is one of the following
  - Maximum distance(also called complete-linkage clustering)  $dist_{max}(C_i, C_j) = \max_{x \in C_i, z \in C_j} dist(x, z)$
  - Minimum distance(also called single-linkage clustering)  $dist_{\min}(C_i, C_j) = \min_{x \in C_i, z \in C_j} dist(x, z)$
  - Average distance (also called average-linkage clustering)

$$dist_{avg}(C_i, C_j) = \frac{1}{|C_i||C_j|} \sum_{x \in C_i} \sum_{z \in C_j} dist(x, z)$$

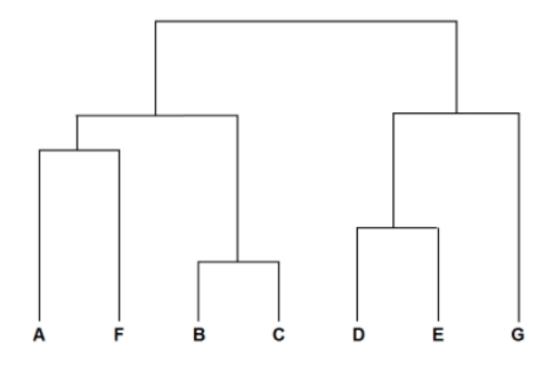


```
输入: 样本集D = \{x_1, x_2, \dots, x_m\};
       聚类簇距离度量函数d \in \{d_{\min}, d_{\max}, d_{\text{avg}}\};
       聚类簇数k.
过程:
1: for j = 1, ..., m do
2: C_j = \{x_j\}
3: end for
4: for i = 1, ..., m do
5: for j = i, ..., m do
    M(i,j) = d(C_i, C_j);
7: M(j,i) = M(i,j)
8: end for
9: end for
10: 设置当前聚类簇个数: q=m
11: while q > k do
     找出距离最近的两个聚类簇(C_{i^*}, C_{i^*});
     合并(C_{i^*}, C_{j^*}): C_{i^*} = C_{i^*} \bigcup C_{j^*};
13:
    for j = j^* + 1, ..., q do
14:
     将聚类簇C_i重编号为C_{i-1}
15:
16:
    end for
     删除距离矩阵M的第j*行与第j*列;
17:
   for j = 1, ..., q - 1 do
18:
    M(i^*, j) = d(C_{i^*}, C_j);
19:
     M(j, i^*) = M(i^*, j)
20:
   end for
21:
     q = q - 1
23: end while
24: return 簇划分结果
输出: 簇划分\mathcal{C} = \{C_1, C_2, \dots, C_k\}
```

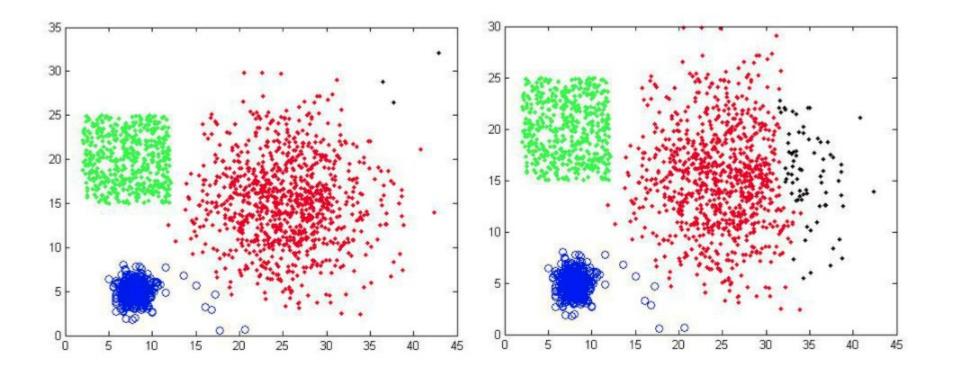
计算距离矩阵

每次循环合并 两个簇并更新 距离矩阵











- Partitioning Methods
- k-means, k-medoids, CLARANS, FCM
- Hierarchical Methods
- AGNES, Birch, Cure, Rock, CHEMALOEN
- Density-based Methods
- DBSCAN,OPTICS
- Grid-based Methods
- Model-Based Methods
- Transitive closure, Boolean matrix, direct clustering, correlation analysis clustering, clustering method based on statistics.....

## **Performance Measurement**



Good clustering should be:

Inter-cluster similarity implication minimized

#### Intra distance & Inter distance



Intra
distance
$$distance = \frac{2}{|C|(|C|-1)} \sum_{1 \le i \le j \le |C|} dist(x_i, x_j)$$

$$distance = \max_{1 \le i \le j \le |C|} dist(x_i, x_j)$$
Inter
distance
$$d_{min}(C_i, C_j) = \min_{x_i \in C_i, x_j \in C_j} dist(x_i, x_j)$$

$$d_{cen}(C_i, C_j) = dis(\mu_i, \mu_j)$$

## **Performance Measurement**



- Internal Index
- Evaluate clustering results directly without using reference model.

- External Index
- Compare clustering results with reference model, for example, partitioning results given by domain expert.

#### **Internal Index**



Davies-Boukdin Index:

#### The smaller the better

$$DBI = \frac{1}{k} \sum_{i=1}^{k} \max_{j \neq i} \left( \frac{avg(C_i) + avg(C_j)}{d_{cen}(C_i, C_j)} \right)$$

Dunn Index:

#### The bigger the better

$$DI = \min_{1 \le i \le k} \left\{ \min_{j \ne i} \frac{d_{min}(C_i, C_j)}{\max_{1 \le l \le k} diam(C_l)} \right\}$$

#### **External Index**



Assume our cluster partition is

$$C = \{C_1, C_2, ... C_k\}$$

- The partition given by reference model is
- $C^* = \{C_1^*, C_2^*, ... C_s^*\}$
- $\triangleright$  let  $\lambda$  and  $\lambda^*$  be clustering label vectors corresponding to
- C and  $C^*$ . Consider  $C_m^2$  sample pairs

$$a = |SS|, SS = \{(x_i, x_j) | \lambda_i = \lambda_j, \lambda_i^* = \lambda_j^*, i < j\}$$

$$b = |SD|, SS = \{(x_i, x_j) | \lambda_i = \lambda_j, \lambda_i^* \neq \lambda_j^*, i < j\}$$

$$c = |DS|, SS = \{(x_i, x_j) | \lambda_i \neq \lambda_j, \lambda_i^* = \lambda_j^*, i < j\}$$

$$d = |DD|, SS = \{(x_i, x_j) | \lambda_i \neq \lambda_j, \lambda_i^* \neq \lambda_j^*, i < j\}$$

$$a + b + c + d = C_m^2 = m(m - 1)/2$$

#### **External Index**



■ JC: Jaccard Coefficient

$$JC = \frac{a}{a+b+c}$$

■ FMI: Fowlkes and Mallows Index

$$FMI = \sqrt{\frac{a}{a+b} \cdot \frac{a}{a+c}}$$

■ RI: Rand Index

$$RI = \frac{2(a+d)}{m(m-1)}$$

[0,1] interval the bigger the better

# **Applications**

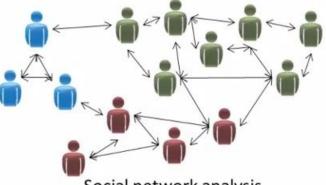




Organize computing clusters



Market segmentation



Social network analysis



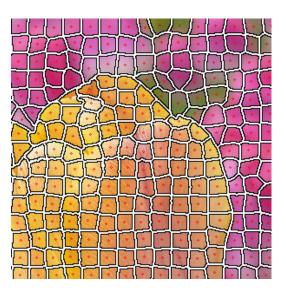
Astronomical data analysis

## An Example



Superpixel segmentation uses the similarity of features between pixels to group pixels, and replaces a large number of pixels with a small number of superpixels to express image features.





# MIMA Group

# Thank You!

**Any Question?**