# 人工智能 无监督学习 +Cluster Analysis

中山大学 计算机学院



### **Unsupervised Learning**

• Why?

• How?

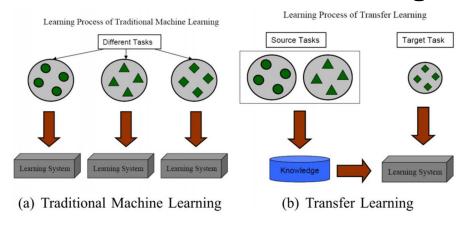
#### Pre-trained Models

- For DNN, one of challenges: data hungry
  - overfit
  - poor generalization ability

- One solution: construct high-quality datasets
  - expensive
  - > time-consuming

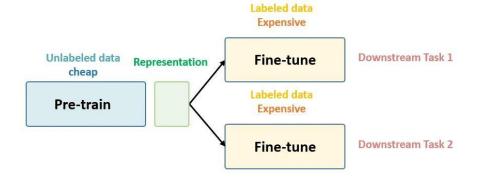
Key: train effective deep neural models with limited human-annotated data

#### An Intuitive idea : Transfer learning



#### • Two stage:

- > 1. Pre-training on source tasks
- 2. Fine-tuning on target tasks



#### The history of pre-trained models

1. Vision models(based on CNNs)

AlexNet, VGG, ResNet

2. Language models(based on Transformer)

GPT, BERT, UNILM

3. Vision models(based on Vision Transformer)

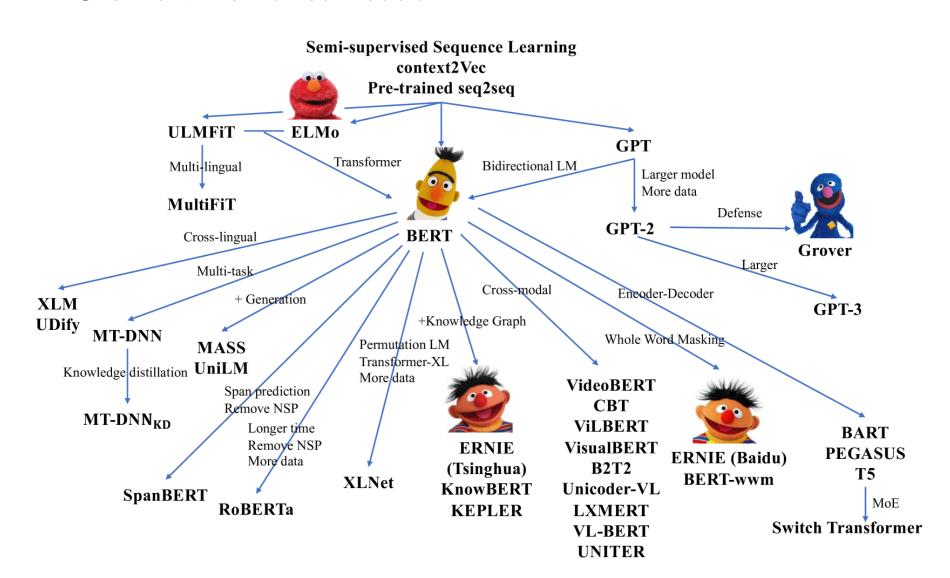
MAE, MoCo, BEiT, simMIM

4. Multimodal models

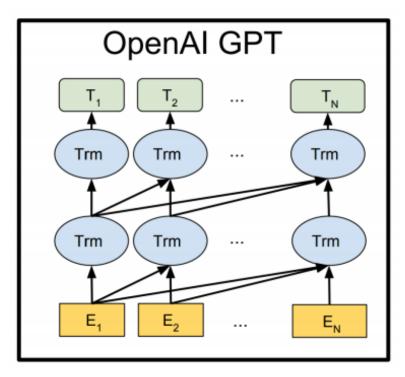
CLIP, BLIP

#### Pre-trained Models

Overview: Pre-trained models



- First SOTA pre-trained language model
- From OpenAl
- Later: GPT-2, GPT-3, ......



#### Unsupervised pre-training

Unsupervised pre-training, maximizing the log-likelihood

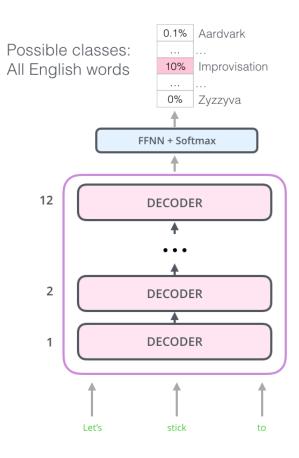
$$L_1(\mathcal{U}) = \sum_i \log P(u_i \mid u_{i-k}, \dots, u_{i-1}; \Theta)$$

where  $\mathcal{U} = \{u_1, \dots, u_n\}$  s an unsupervised corpus of tokens, k is the size of context window, P is modelled as a neural network with parameters  $\Theta$ .

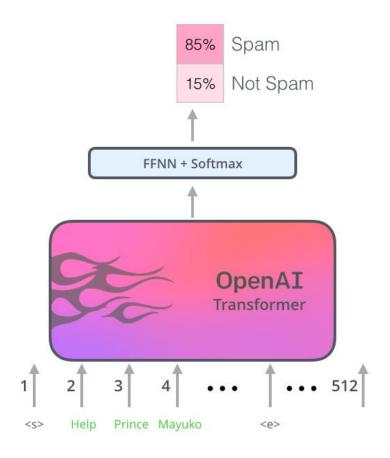
$$egin{aligned} h_0 &= UW_e + W_p \ h_l &= ext{transformer}\, b \operatorname{bock}(h_{l-1}) orall i \in [1,n] \ P(u) &= \operatorname{softmax}ig(h_n W_e^Tig) \end{aligned}$$

where U is one-hot representation of tokens in the window, n is the total number of transformer layers, transformer\_block() denotes the decoder of the Transformer model (multi-headed self-attention and position-wise feedfoward layers).

- Pre-training dataset:
  - > 7000+ books



• An example: Transfer Learning to Downstream Tasks

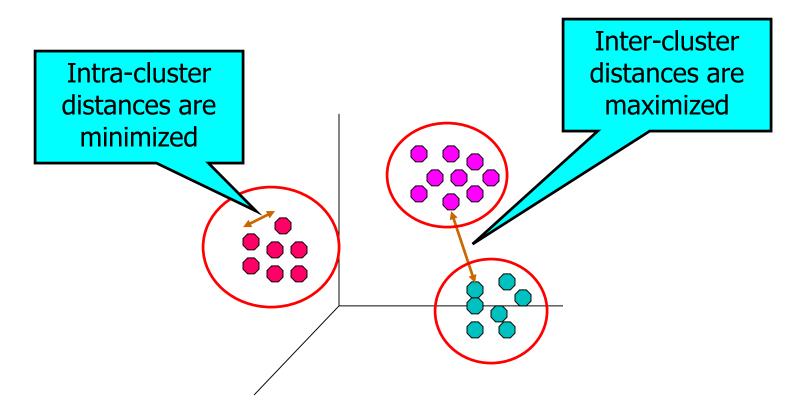


## **Cluster Analysis**

### What is Cluster Analysis?



 Finding groups of objects such that the objects in a group will be similar (or related) to one another and different from (or unrelated to) the objects in other groups



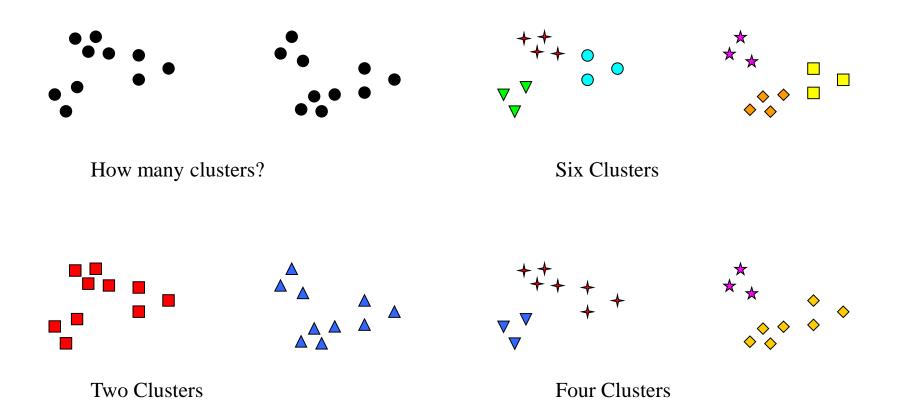
### **Clustering in Social Network**





### Notion of a Cluster can be Ambiguous

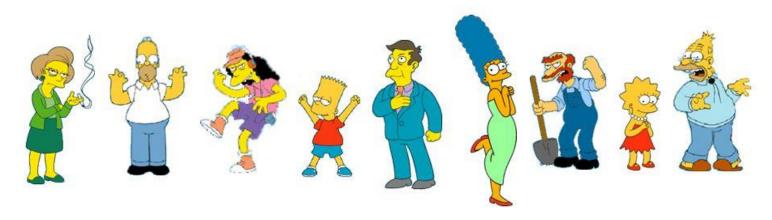




### **Notion of a Cluster can be Ambiguous**



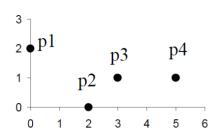
What is a natural grouping among these objects?



### Types of Clusters: Objective Function ...



- Map the clustering problem to a different domain and solve a related problem in that domain
  - Proximity (亲近度) matrix defines a weighted graph, where the nodes are the data points, and the weighted edges represent the proximities (similarity or dissimilarity) between data points



point	X	у
p1	0	2
p2	2	0
p3	3	1
p4	5	1

	p1	p2	p3	p4
p1	0.000	2.828	3.162	5.099
p2	2.828	0.000	1.414	3.162
р3	3.162	1.414	0.000	2.000
p4	5.099	3.162	2.000	0.000

**Figure** Four points and their corresponding data and proximity (distance) matrices.

Similarity? Dissimilarity?

Euclidean Distance / Manhattan Distance / Chebyshev Distance/
Cosine Similarity /..... Similarity=1-Dissimilarity

#### **Bonus: Some Distance Metrics**



• 
$$L_p(x_1, x_2) = \left(\sum_{l=1}^n \left| x_i^{(l)} - x_j^{(l)} \right|^p \right)^{\frac{1}{p}}$$

• 
$$L_1(x_1, x_2) = \sum_{l=1}^{n} \left| x_i^{(l)} - x_j^{(l)} \right|$$

Manhattan Distance

• 
$$L_2(x_1, x_2) = \sum_{l=1}^n \sqrt{\left(x_i^{(l)} - x_j^{(l)}\right)^2}$$

**Euclidean Distance** 

• 
$$L_{\infty}(x_1, x_2) = \sqrt[\infty]{\sum_{l=1}^{n} |x_i^{(l)} - x_j^{(l)}|^{\infty}} = max(|x_i - x_j|)$$

Chebyshev Distance

• 
$$L_{-\infty}(x_1, x_2) = \sqrt[-\infty]{\sum_{l=1}^{n} |x_i^{(l)} - x_j^{(l)}|^{-\infty}} = min(|x_i - x_j|)$$

• Cosine Distance: 
$$D(x_1, x_2) = 1 - \cos(x_1, x_2) = 1 - \frac{x_1 \cdot x_2}{||x_1||_2 ||x_2||_2}$$

### **K-means Clustering**



- Partitional clustering approach
- Each cluster is associated with a centroid (center point)
- Each point is assigned to the cluster with the closest centroid
- Number of clusters, K, must be specified
- The basic algorithm is very simple

- 1: Select K points as the initial centroids.
- 2: repeat
- 3: Form K clusters by assigning all points to the closest centroid.
- 4: Recompute the centroid of each cluster.
- 5: **until** The centroids don't change

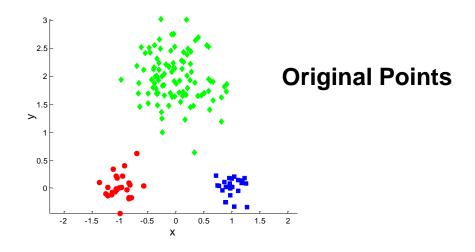
### K-means Clustering — Details

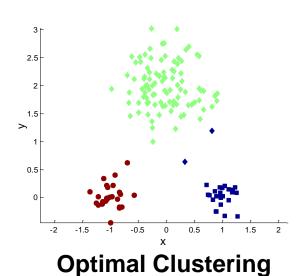


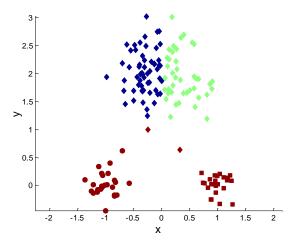
- Initial centroids are often chosen randomly.
  - Clusters produced vary from one run to another.
- The centroid is (typically) the mean of the points in the cluster.
- 'Closeness' is measured by Euclidean distance, cosine similarity, correlation, etc.
- K-means will converge for common similarity measures mentioned above.
- Most of the convergence happens in the first few iterations.
  - Often the stopping condition is changed to 'Until relatively few points change clusters'
- Complexity is O( n \* K \* I \* d )
  - n = number of points, K = number of clusters,
     I = number of iterations, d = number of attributes

### **Two different K-means Clusterings**





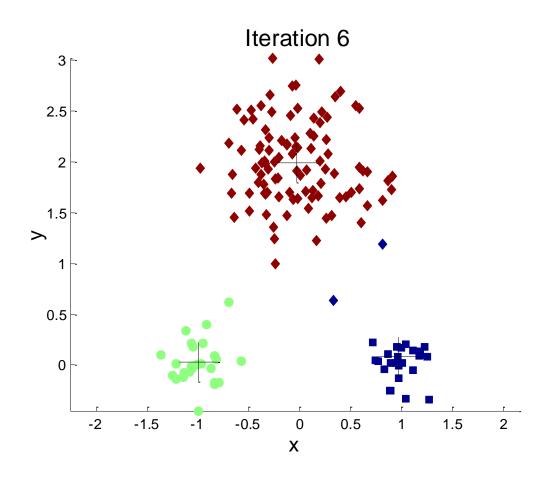




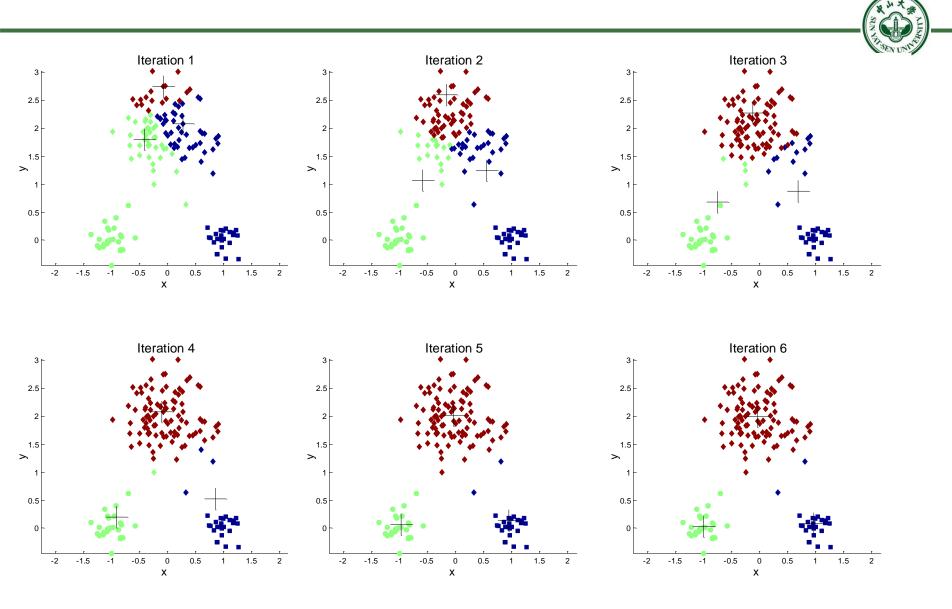
**Sub-optimal Clustering** 

### **Importance of Choosing Initial Centroids**





### **Importance of Choosing Initial Centroids**



### **Evaluating K-means Clusters**

- Most common measure is Sum of Squared Error (SSE)
  - For each point, the error is the distance to the nearest cluster
  - To get SSE, we square these errors and sum them.

$$SSE = \sum_{i=1}^{K} \sum_{\mathbf{x} \in C_i} dist^2(m_i, \mathbf{x})$$

- x is a data point in cluster  $C_i$  and  $m_i$  is the representative point for cluster  $C_i$ 
  - ◆ can show that m<sub>i</sub> corresponds to the center (mean) of the cluster
- Given two clusters, we can choose the one with the smallest error
- One easy way to reduce SSE is to increase K, the number of clusters
  - ◆ A good clustering with smaller K can have a lower SSE than a poor clustering with higher K

#### **Limitations**



- K-means has problems when clusters are of differing
  - Sizes
  - Densities
  - Non-globular shapes

 K-means also has problems when the data contains outliers.

#### **Discussion**

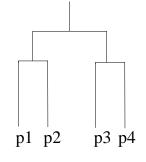


对象?



### **Method 1: Hierarchical Clustering**

将全宇宙的生命按照基因相似度构造Dendrogram,确定聚类数目后,在每两个相邻的聚类簇中随机挑选一个进行消灭,以保证宇宙生物多样性。



#### **Discussion**



### **Method 2: K-means Clustering**

宇宙包含"九大国度",为防止某个国度灭绝,在每个国度选取一个聚类中心,通过K-means得到聚类簇,再在每个聚类中随机选取一半生物进行消灭,宇宙生物多样性得到进一步保证。

