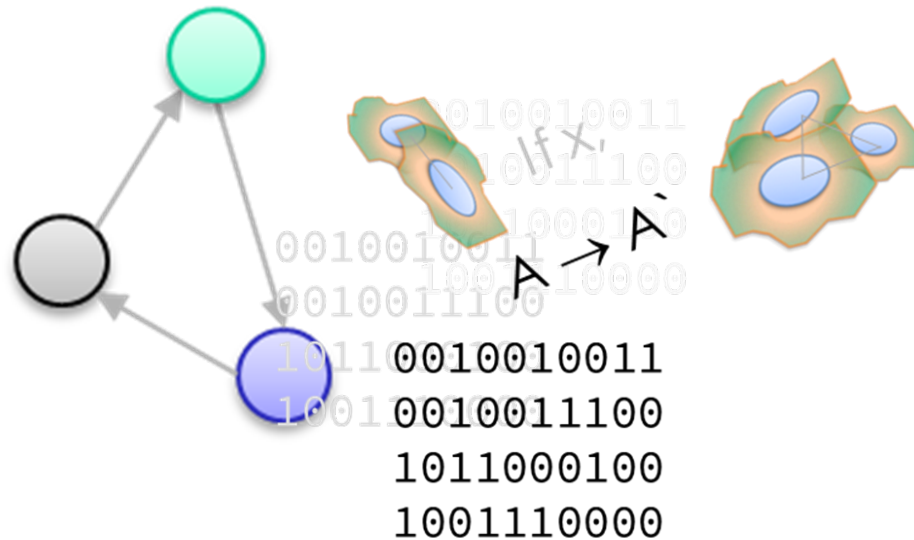


Computational Bioengineering

BME 4803 | BME 6313



Clustering Introduction

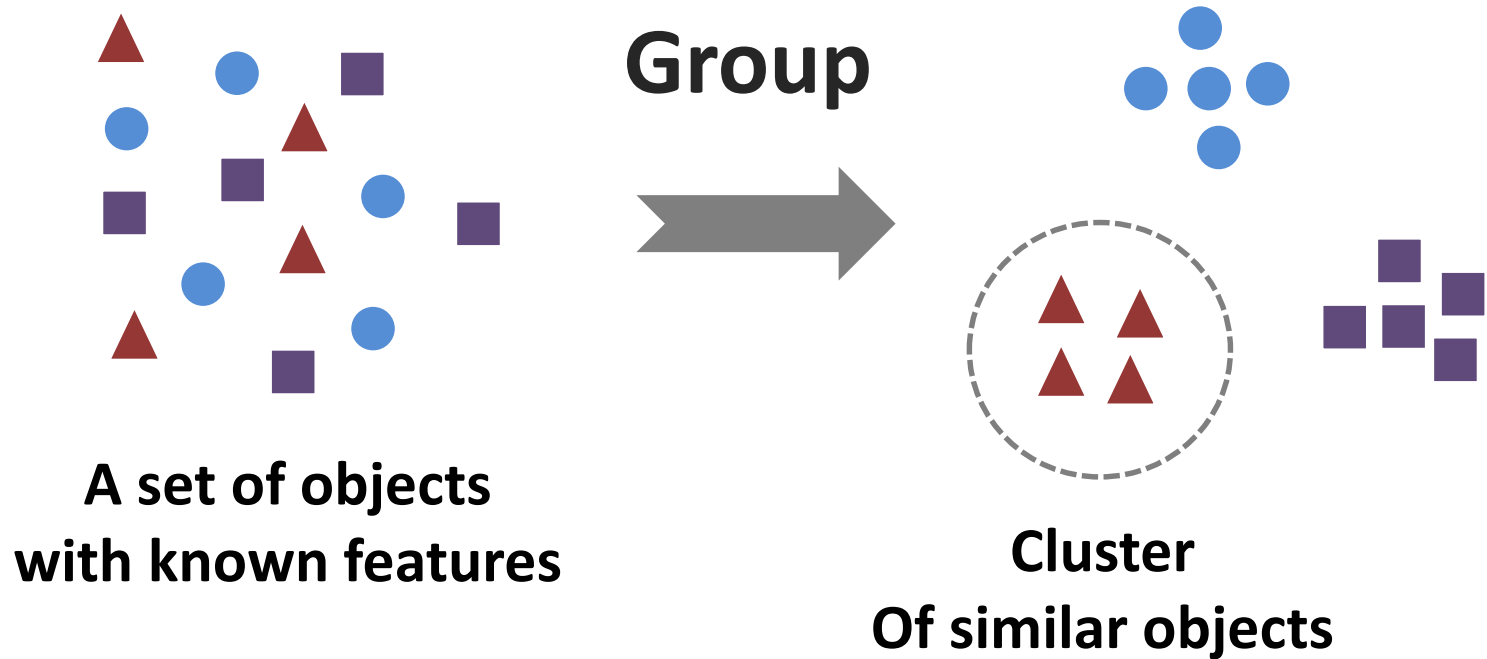
Slide Credits:

Dr. Chenyue Wendy Hu

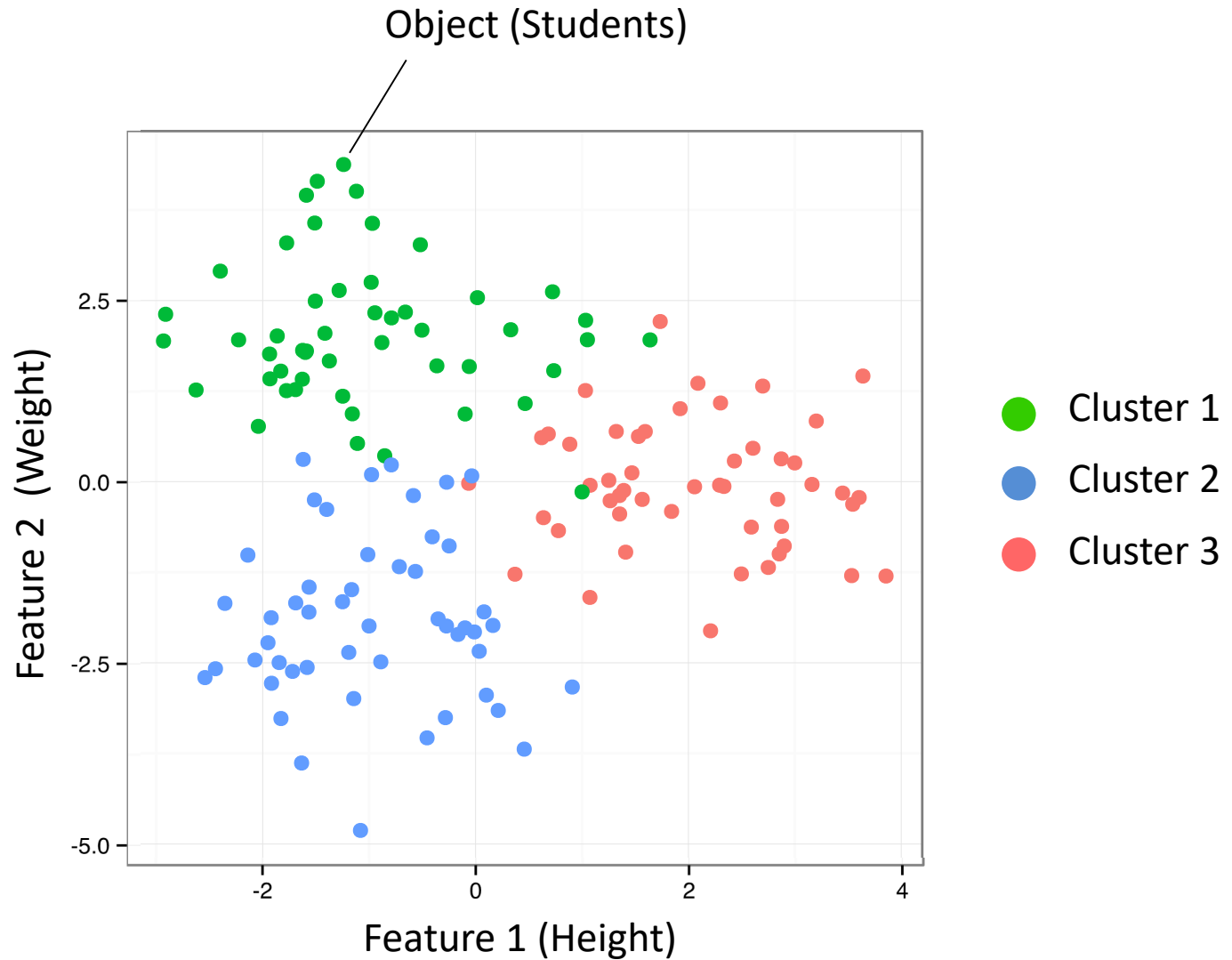
Outline

- What is cluster analysis?
 - concept, examples, application scenarios
- How does cluster analysis work?
 - Similarity (Euclidean distance)
 - K-means Clustering
 - Hierarchical Clustering
- How to pick the cluster number?
 - Elbow Method
 - Silhouette
- Best practices & review

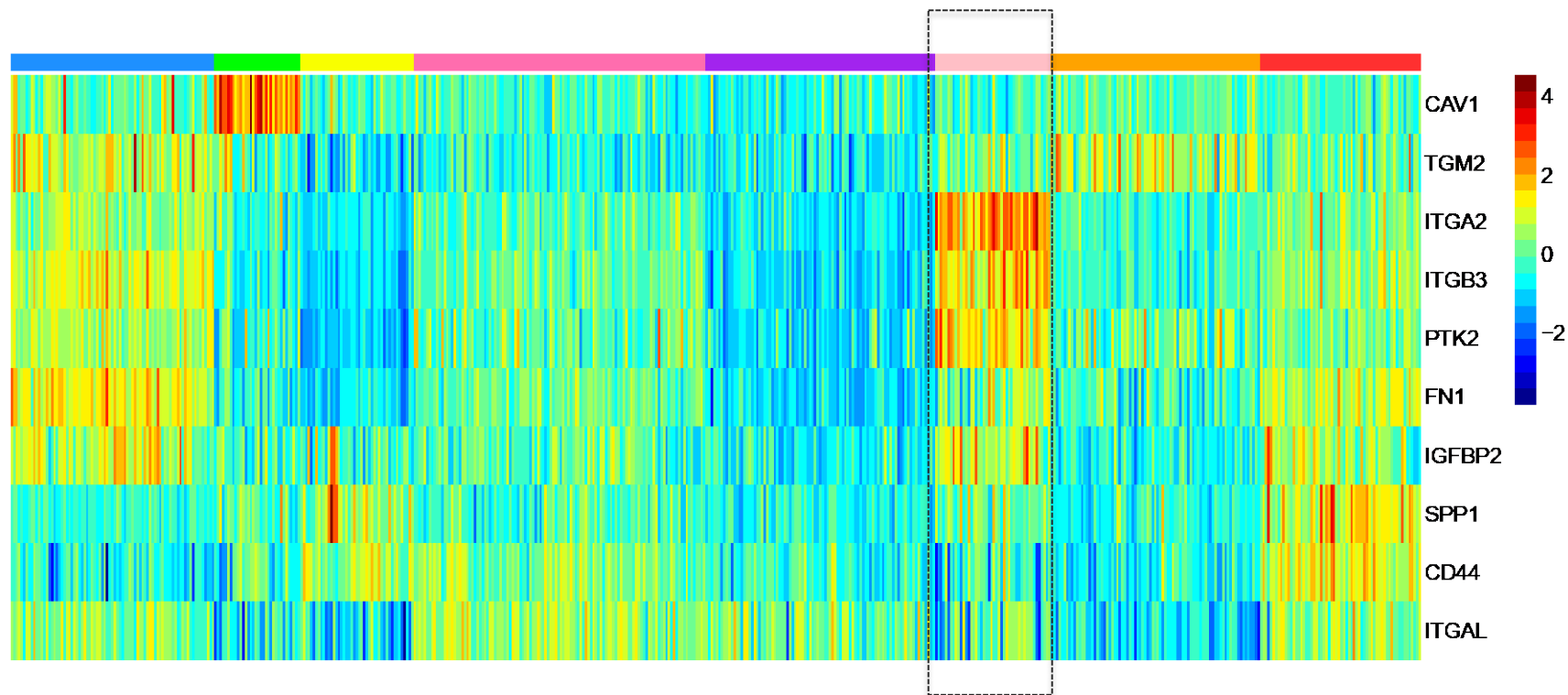
Concept



Example 1



Example 2



Cluster

Application

Application scenarios

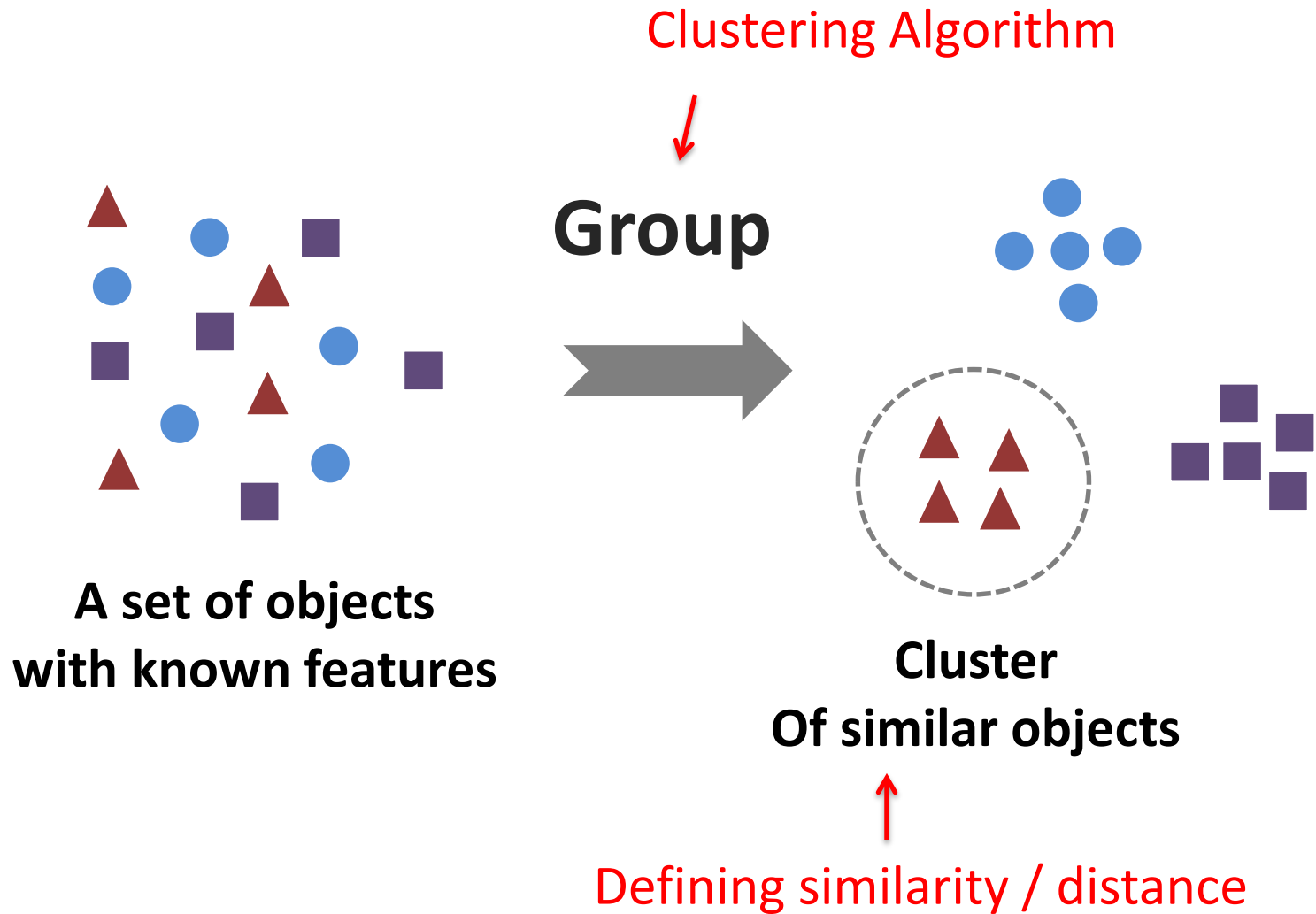
- You have multiple features (> 1)
 - matrix or data table
- You believe the population is heterogeneous
- You don't know the true grouping of objects



Application goals

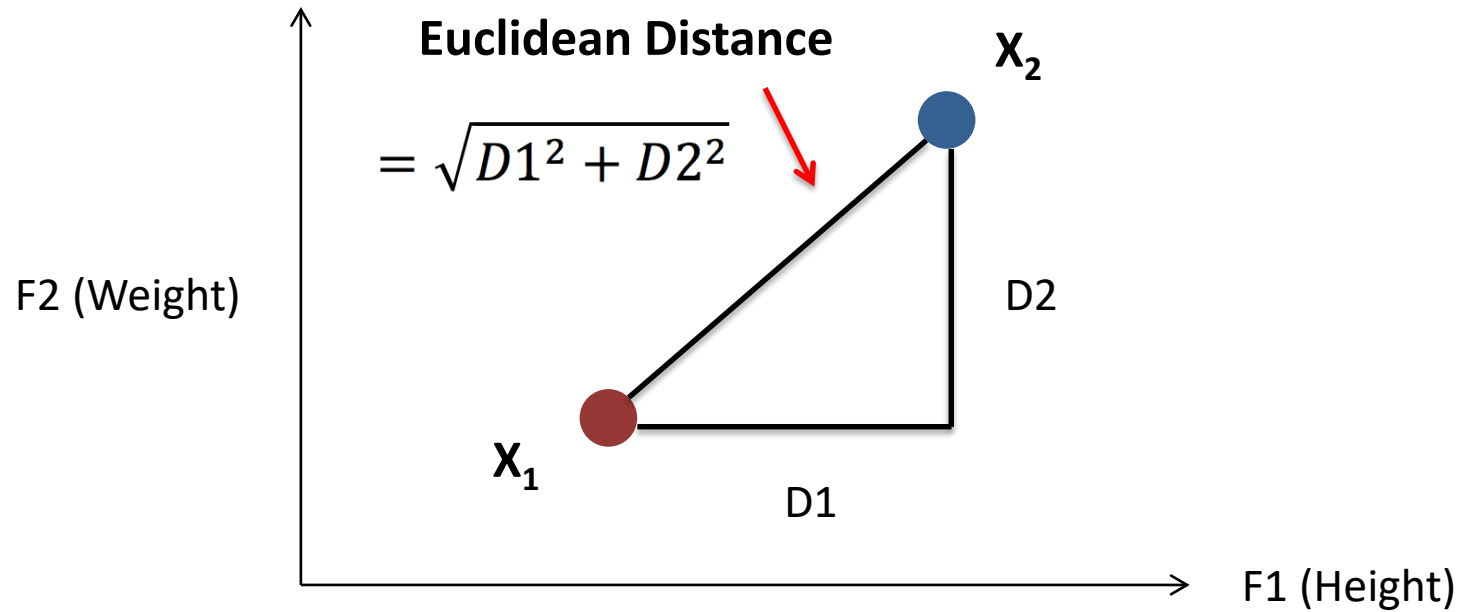
- ✓ End-step: Define classes or categories
- ✓ Pre-step: Segment samples for subsequent analyses

Concept



Similarity

Distance (Dissimilarity)



Generalized Euclidean Distance
Between X_i and X_j
In n dimensional space

$$D_{ij} = \sqrt{\sum_{k=1}^n (D_{ik} - D_{jk})^2}$$

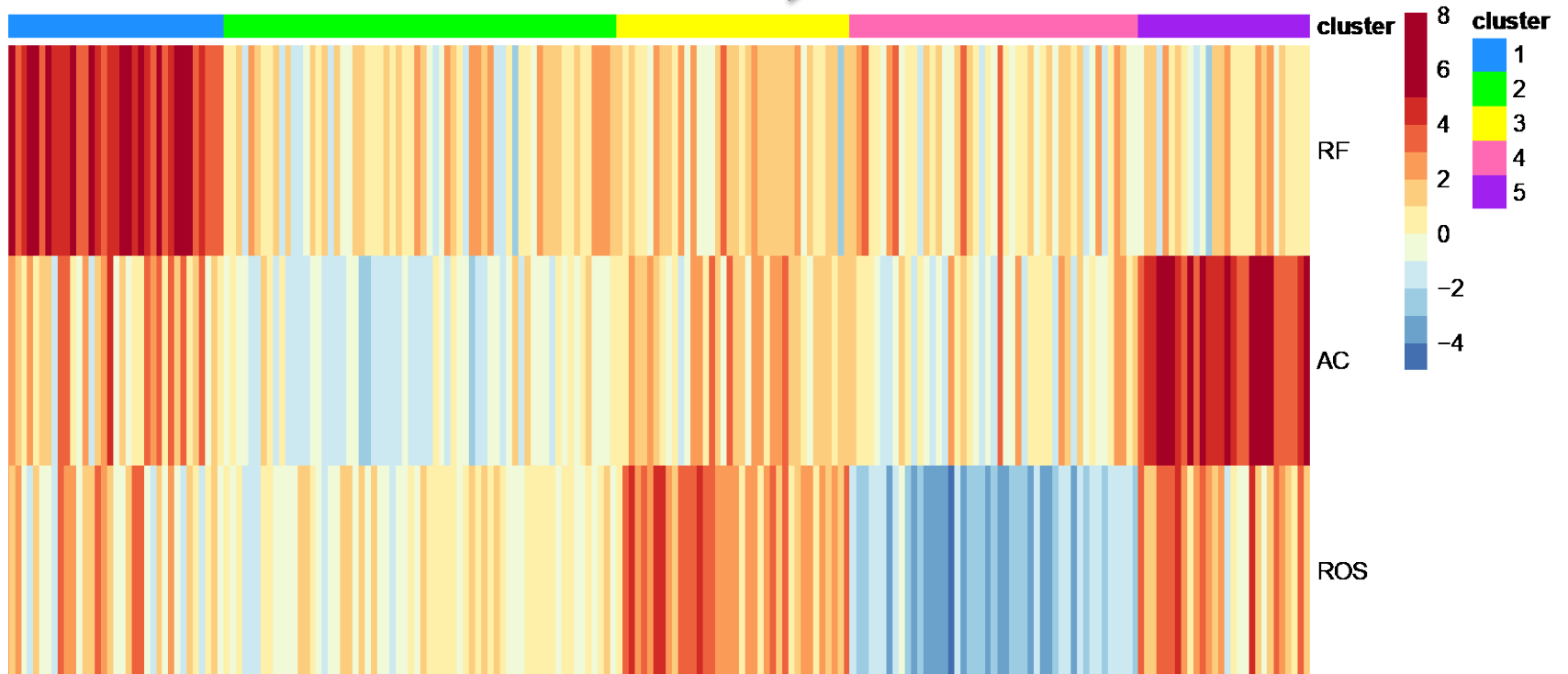
Algorithms

Common Clustering Algorithms

- **Hierarchical Clustering**
- **K-means**
- Affinity Propagation
- Spectral Clustering
- DBSCAN

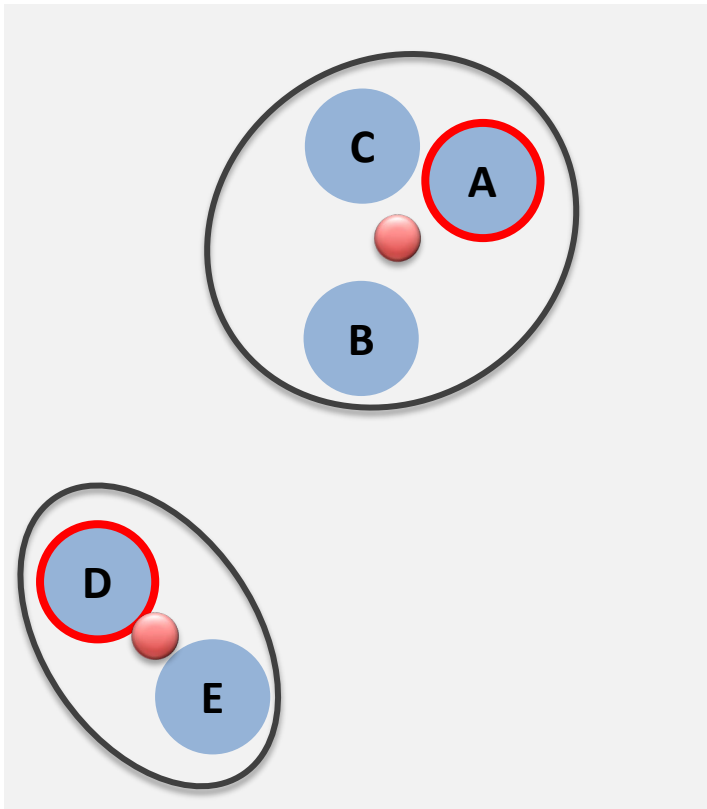
K-means

No dendrogram



K-means

- Start with K initial random cluster centers
- Repeat the following steps:
 - Assign objects to their closest cluster centers
 - Recompute cluster centers
- End when the solution is optimized (nothing to do)



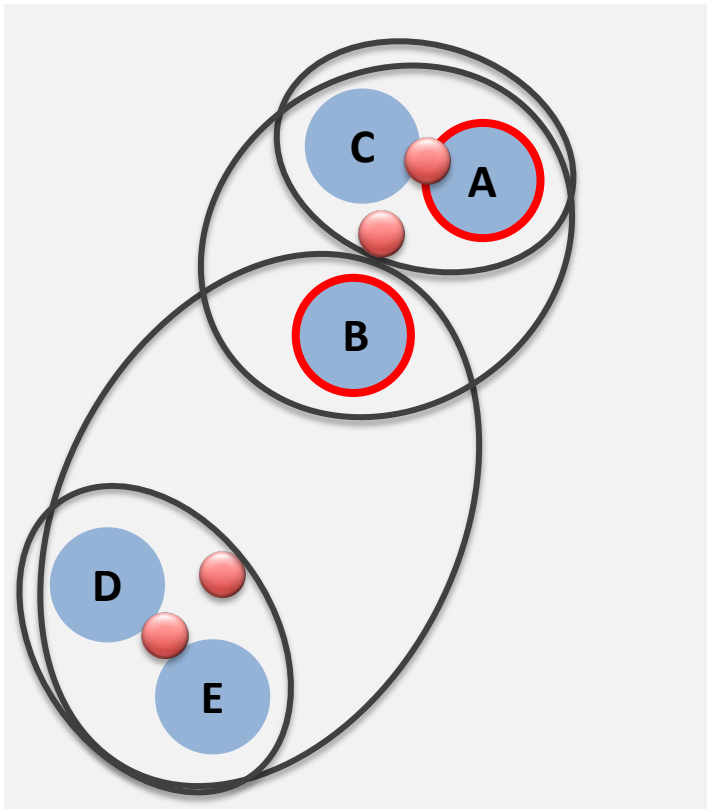
Initialization: A, D

Cluster: {A, B, C}, {D, E}

Cluster: {A, B, C}, {D, E}

K-means

- Start with K initial random cluster centers
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Initialization: A, D

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Initialization: A, B

Cluster: {A, C}, {B, D, E}

Cluster: {A, B, C}, {D, E}

Cluster: {A, B, C}, {D, E}

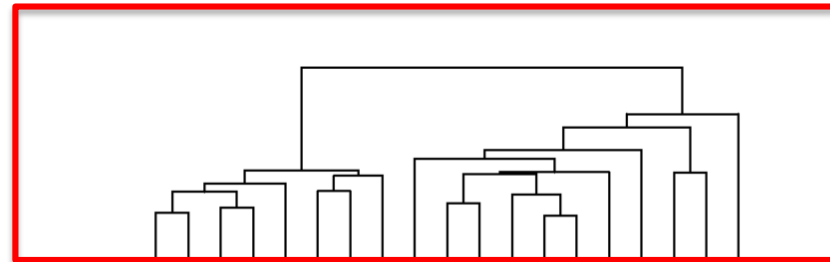
K-means

Summary

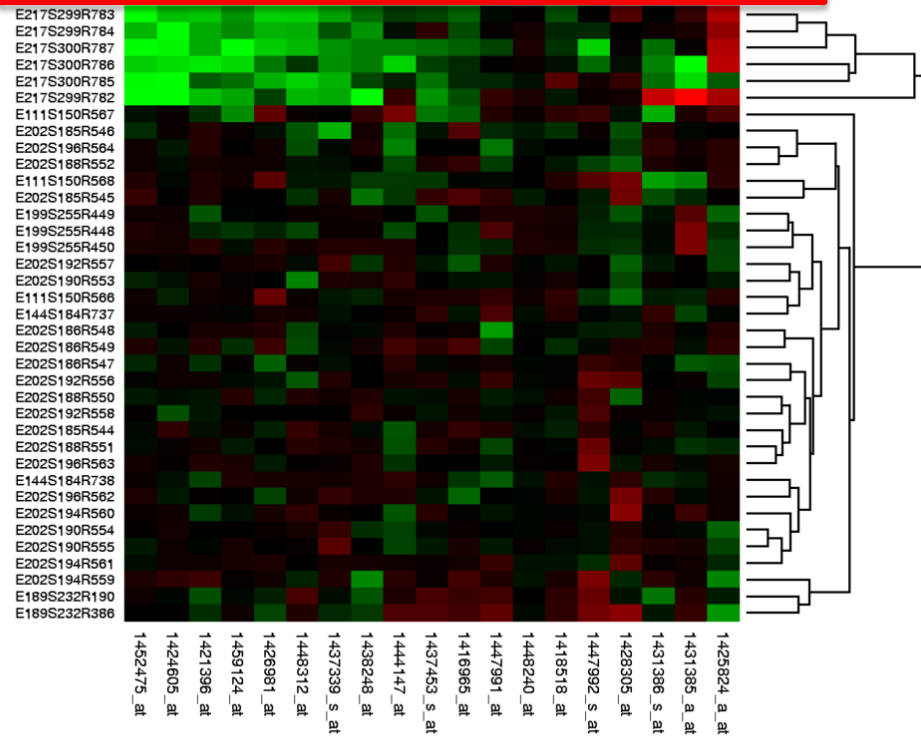
- Need to provide K (number of clusters)
 - Iterative assignment and center update
 - Optimize within-cluster distances
 - Converge to local optimum
- Not necessarily the global optimum
 - Different random initiations -> different solutions
 - Make sure to run the algorithm multiple times (≥ 100)

Slower than Hierarchical, but better clustering quality

Hierarchical

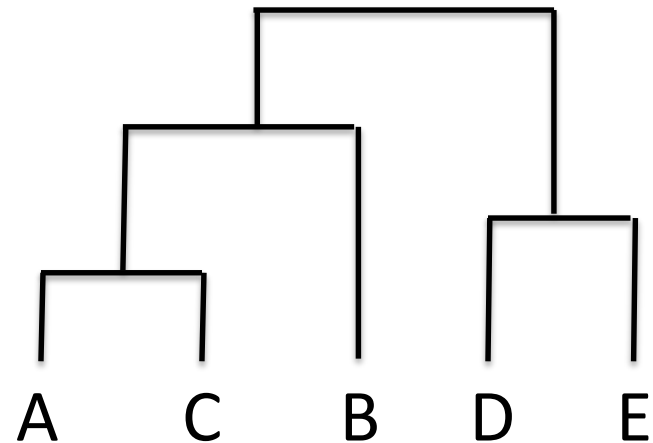
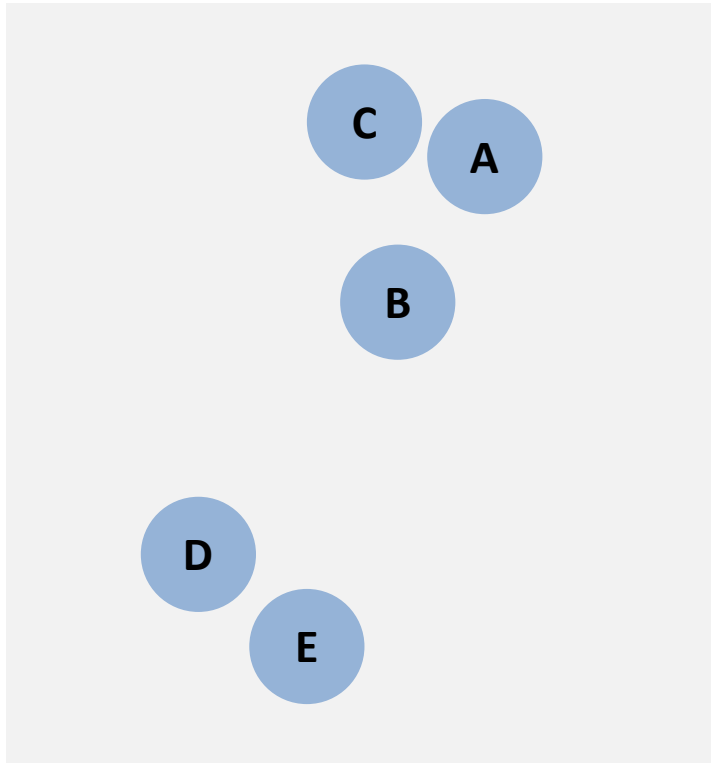


Dendrogram



Hierarchical

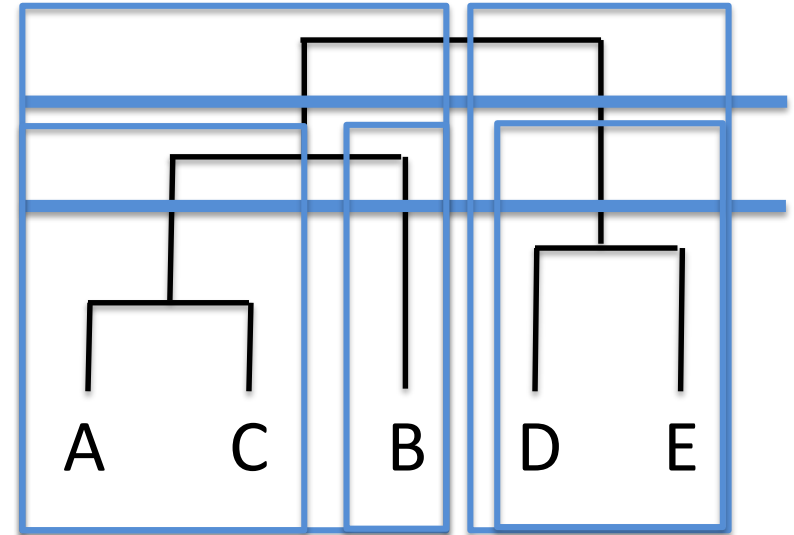
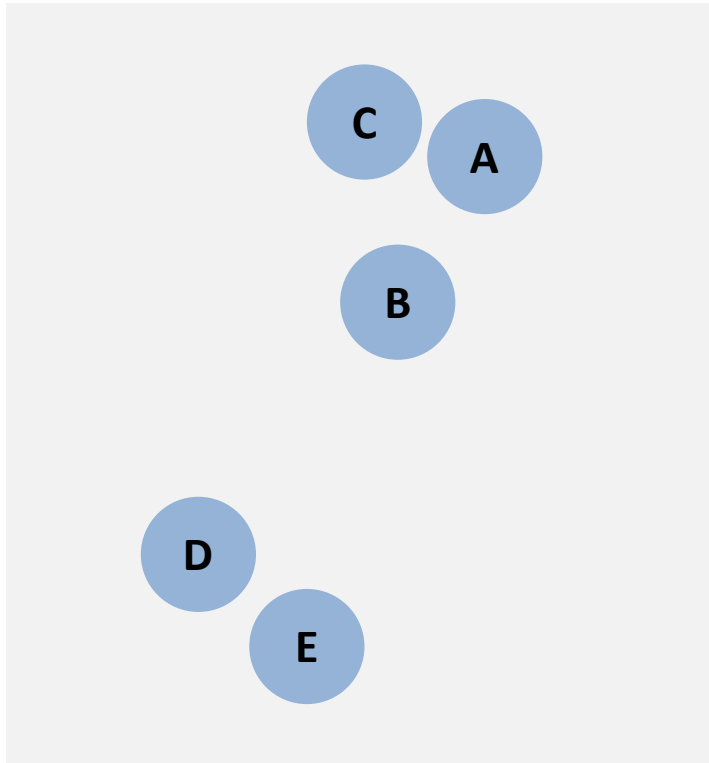
- Start with each object as a cluster
- Repeatedly connect the closest two clusters
- End when there is only one cluster left



Minimum **Linkage**: the smallest distance between elements of each cluster

Hierarchical

Cut tree to obtain a clustering solution

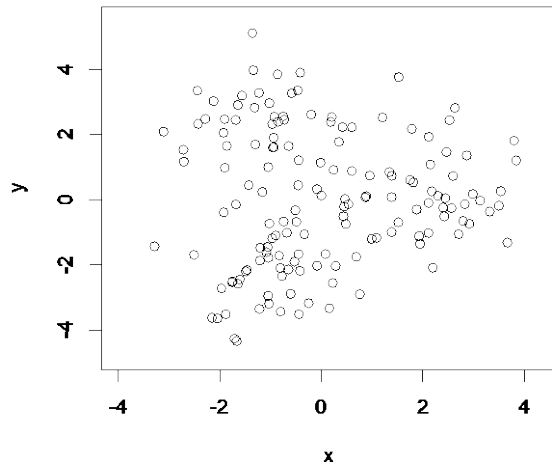


Hierarchical

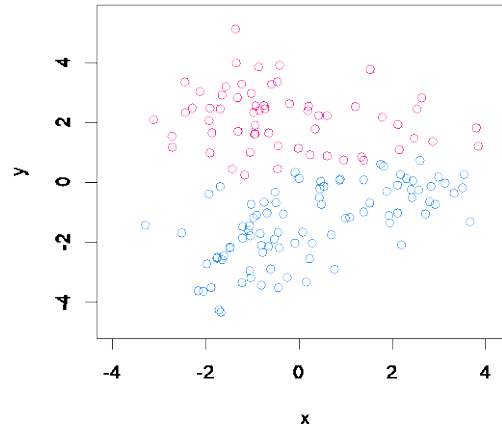
Summary

- Dendrogram (tree structure)
 - Bottom-up or agglomerative approach
 - Linkage Method
 - Cut-tree
-
- Linkage Method Choice: Maximum (Complete), Minimum (Single), Average, Ward
 - Pro: Good visualization, Fast Computing
 - Con: Biased towards outliers

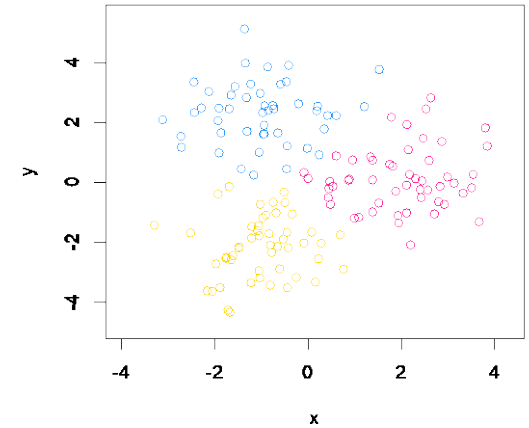
Cluster Number



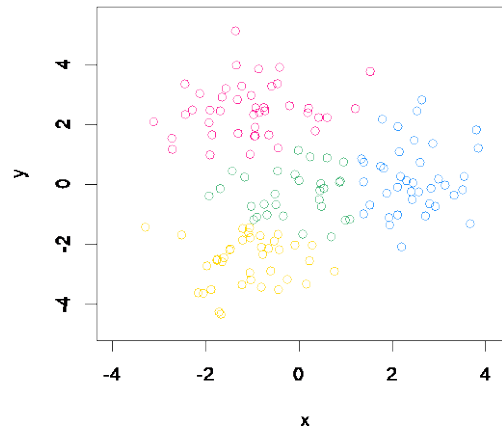
K=2



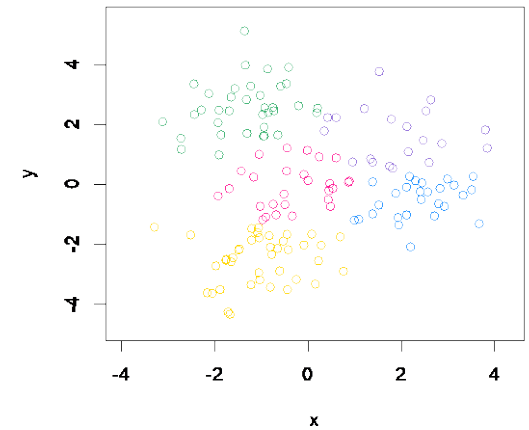
K=3



K=4



K=5



Cluster Number

Computational Methods

- Elbow Method
- Silhouette
- Gap Statistic
- Progeny Clustering*

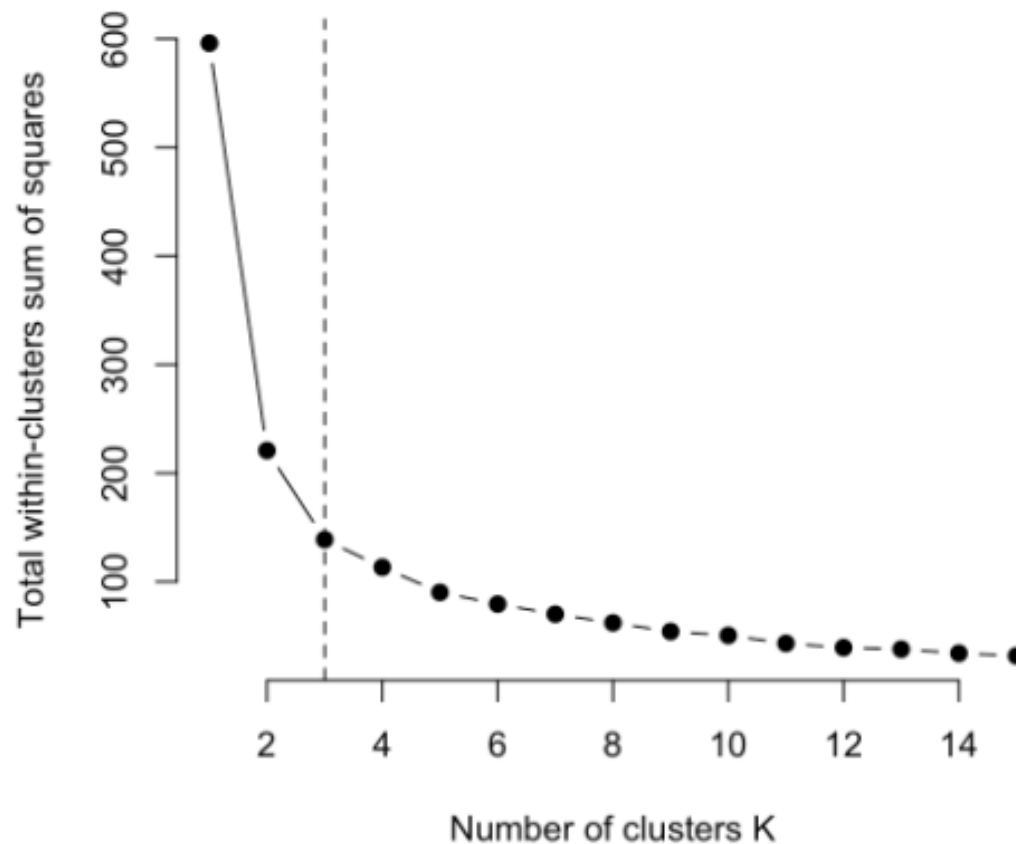
Procedure

1. Cluster data repeated with varying cluster numbers ($K=2, \dots, 10$)
2. Pick the number with best clustering quality

*Progeny Clustering (Hu et al.): <https://www.nature.com/articles/srep12894>

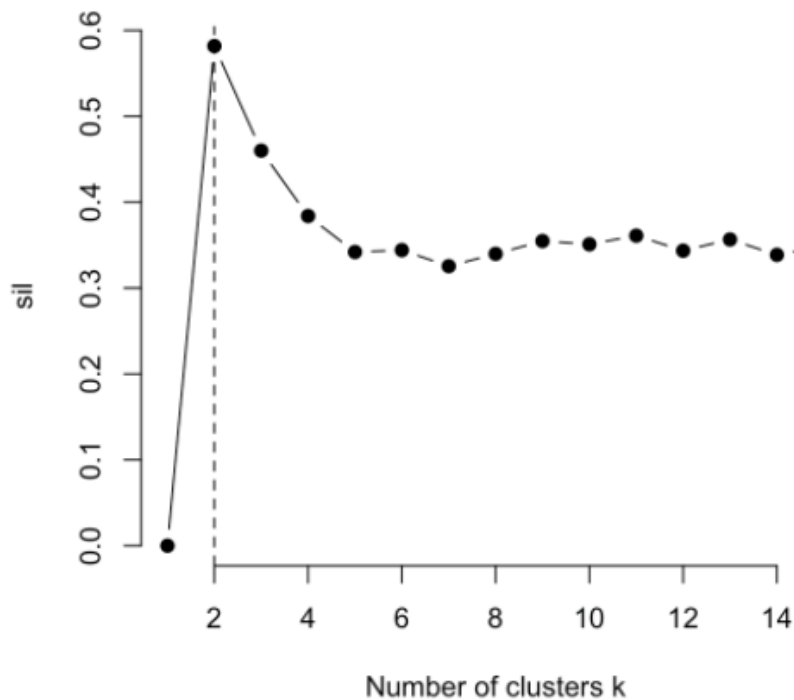
Cluster Number

Elbow Method



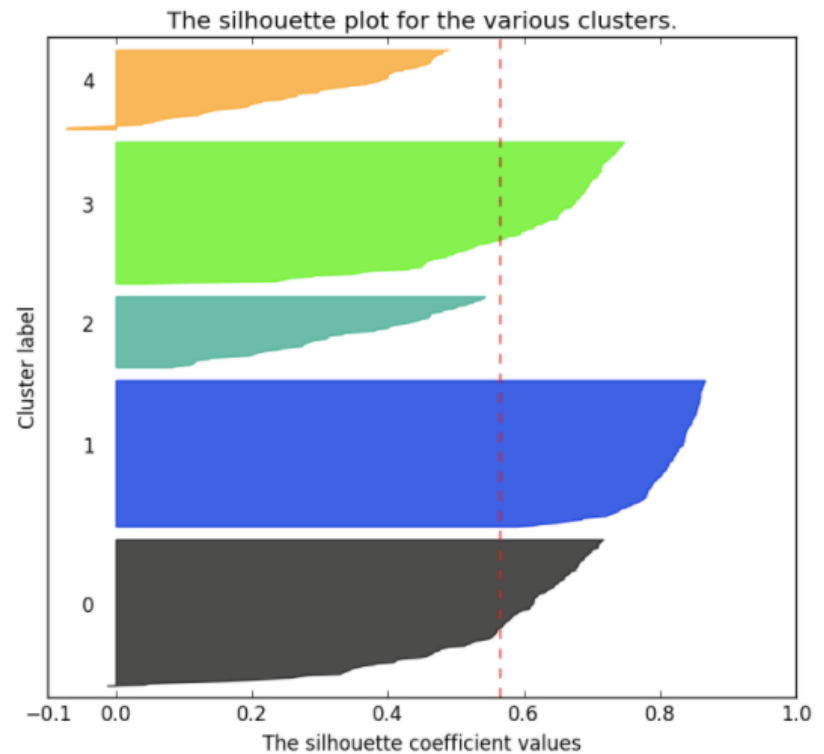
Cluster Number

$$s(i) = \frac{b(i) - a(i)}{\max\{a(i), b(i)\}}$$



Silhouette

$$-1 \leq s(i) \leq 1$$



Best Practice

- Data Normalization
 - Scaling -> Equal Variance
- Clustering Algorithm Choice
 - K-means with multiple runs
- Cluster Number Choice
 - Silhouette or your expert knowledge
- Visualization
 - Heatmaps with dendrograms or annotation bars
 - Scatter Plot with Principal Components (PCA)

Outline

- **Concept**
 - **Application**
 - **Best Practices**
 - **Circular Heatmap**
 - **Tutorial**
- ✓ what is a heatmap?
 - ✓ when do I need a heatmap?
 - ✓ how should I choose the color?
 - ✓ what is a circular heatmap?
 - ✓ I can draw heatmaps!