# CSE/STAT 416

#### **Other Clustering Methods**

Pre-Class Videos

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May 17, 2024



Pre-Class Video 1

Clustering Recap

# Clustering









### SPORTS









### WORLD NEWS



# Define Clusters



In their simplest form, a **cluster** is defined by

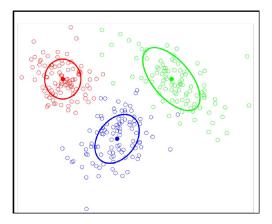
- The location of its center (centroid)
- Shape and size of its spread

**Clustering** is the process of finding these clusters and **assigning** each example to a particular cluster.

- $x_i$  gets assigned  $z_i \in [1, 2, ..., k]$
- Usually based on closest centroid

Will define some kind of score for a clustering that determines how good the assignments are

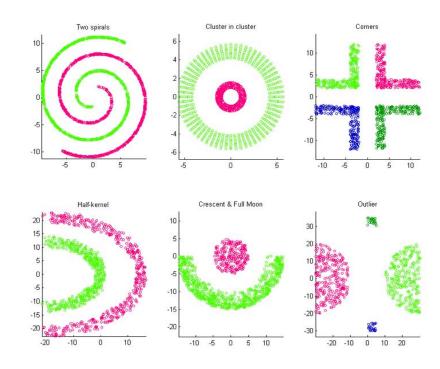
Based on distance of assigned



# Not Always Easy

There are many clusters that are harder to learn with this setup

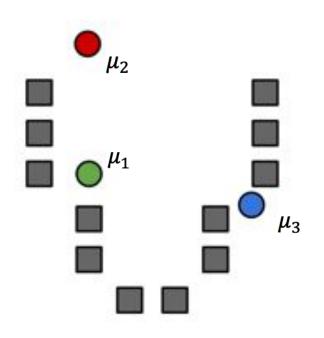
Distance does not determine clusters



### Step 0

Start by choosing the initial cluster centroids

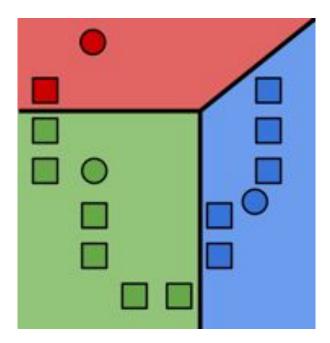
- A common default choice is to choose centroids at random
- Will see later that there are smarter ways of initializing



# Step 1

Assign each example to its closest cluster centroid

$$z_i \leftarrow \operatorname*{argmin}_{j \in [k]} \left| \left| \mu_j - x_i \right| \right|^2$$



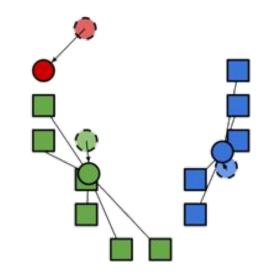


# Step 2

Update the centroids to be the mean of all the points assigned to that cluster.

$$u_j \leftarrow \frac{1}{n_j} \sum_{i: z_i = j} x_i$$

Computes center of mass for cluster!





### Smart Initializing w/ k-means++



Making sure the initialized centroids are "good" is critical to finding quality local optima. Our purely random approach was wasteful since it's very possible that initial centroids start close together.

Idea: Try to select a set of points farther away from each other.

**k-means++** does a slightly smarter random initialization

- 1. Choose first cluster  $\mu_1$  from the data uniformly at random
- 2. For the current set of centroids (starting with just  $\mu_1$ ), compute the distance between each datapoint and its closest centroid
- 3. Choose a new centroid from the remaining data points with probability of  $x_i$  being chosen proportional to  $d(x_i)^2$

# Problems with k-means



In real life, cluster assignments are not always clear cut

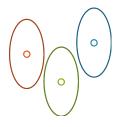
E.g. The moon landing: Science? World News? Conspiracy?

Because we minimize Euclidean distance, k-means assumes all the clusters are spherical



We can change this with weighted Euclidean distance

Still assumes every cluster is the same shape/orientation

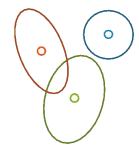


# Failure Modes of k-means

If we don't meet the assumption of spherical clusters, we will get unexpected results

disparate cluster sizes





different shaped/oriented clusters

### Mixture Models



A much more flexible approach is modeling with a mixture model

Model each cluster as a different probability distribution and learn their parameters

- E.g. Mixture of Gaussians
- Allows for different cluster shapes and sizes
- Typically learned using Expectation Maximization (EM) algorithm

Allows soft assignments to clusters

54% chance document is about world news, 45% science, 1% conspiracy theory, 0% other

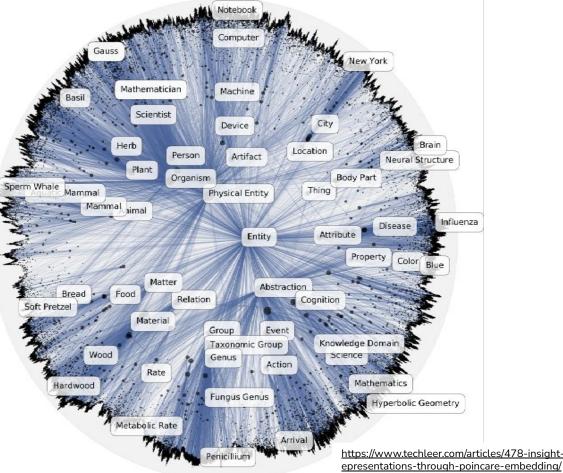
Pre-Class Video 2

Divisive Clustering

Hierarchical Clustering

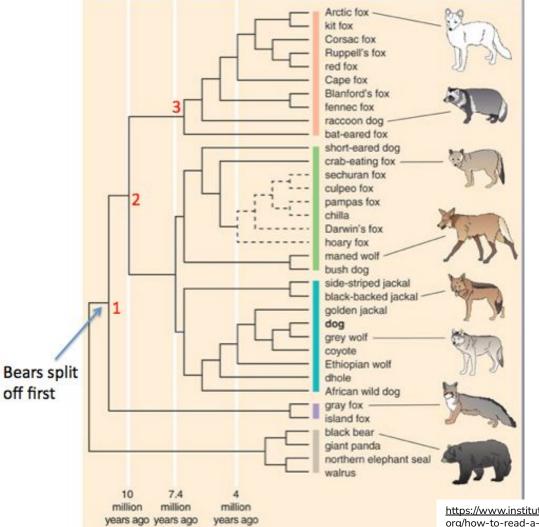
### Nouns





https://www.techleer.com/articles/478-insight-into-hierarchical-r

### Species



https://www.instituteofcaninebiology. org/how-to-read-a-dendrogram.html

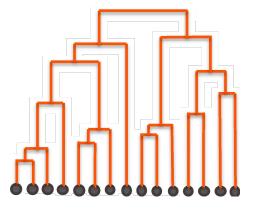
16

### Motivation



If we try to learn clusters in hierarchies, we can

- Avoid choosing the # of clusters beforehand
- Use dendrograms to help visualize different granularities of clusters
- Allow us to use any distance metric
  - K-means requires Euclidean distance
- Can often find more complex shapes than k-means



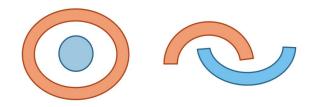
# Finding Shapes





Mixture Models





# Types of Algorithms

#### Divisive, a.k.a. top-down

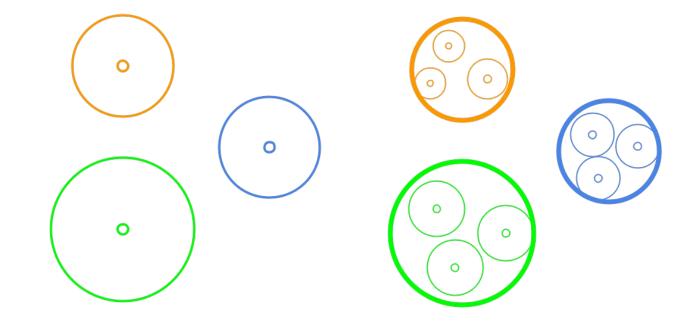
- Start with all the data in one big cluster and then recursively split the data into smaller clusters
  - Example: recursive k-means

#### Agglomerative, a.k.a. bottom-up:

- Start with each data point in its own cluster. Merge clusters until all points are in one big cluster.
  - Example: single linkage clustering

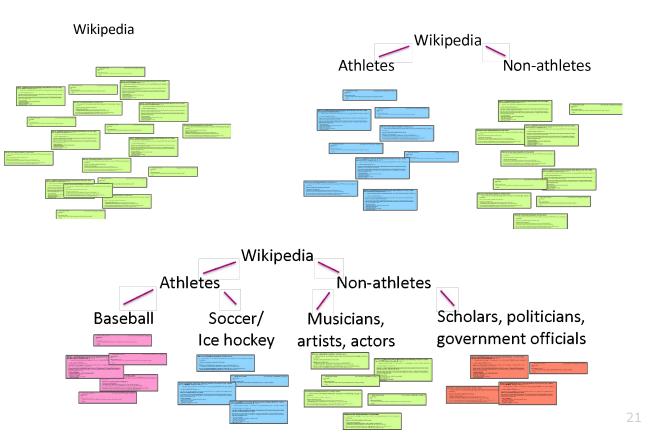
# Divisive Clustering

Start with all the data in one cluster, and then repeatedly run k-means to divide the data into smaller clusters. Repeatedly run k-means on each cluster to make sub-clusters.



### Example

Using Wikipedia



### Choices to Make

For divisive clustering, you need to make the following choices:

- Which algorithm to use (e.g., k-means)
- How many clusters per split
- When to split vs when to stop
  - Max cluster size
    - Number of points in cluster falls below threshold
  - Max cluster radius

distance to furthest point falls below threshold

Specified # of clusters
 split until pre-specified # of clusters is reached

# CSE/STAT 416

**Other Clustering Methods** 

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May 17, 2024

**?** Questions? Raise hand



# Define Clusters



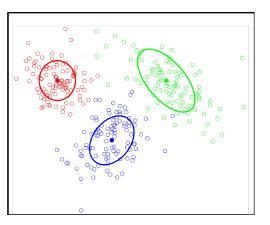
In their simplest form, a **cluster** is defined by

- The location of its center (centroid)
- Shape and size of its spread

**Clustering** is the process of finding these clusters and **assigning** each example to a particular cluster.

- $x_i$  gets assigned  $z_i \in [1, 2, ..., k]$
- Usually based on closest centroid

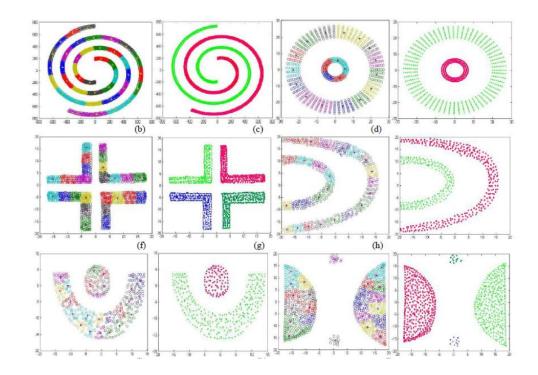
Will define some kind of objective function for a clustering that determines how good the assignments are



# Not Always Easy

There are many clusters that are harder to learn with this setup

Distance does not determine clusters



# Visualizing k-means

https://www.naftaliharris.com/blog/visualizing-k-means-clusterin g/



Smart Initializing w/ k-means++

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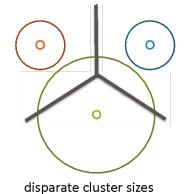
Idea: Try to select a set of points farther away from each other.

**k-means++** does a slightly smarter random initialization

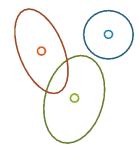
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- 2. For each datapoint  $x_i$ , compute the distance between  $x_i$  and the closest centroid from the current set of centroids (starting with just  $\mu_i$ ). Denote that distance  $d(x_i)$ .
- 3. Choose a new centroid from the remaining data points, where the probability of  $x_i$  being chosen is proportional to  $d(x_i)^2$ .
- 4. Repeat 2 and 3 until we have selected k centroids.

# Failure Modes of k-means

If we don't meet the assumption of spherical clusters, we will get unexpected results



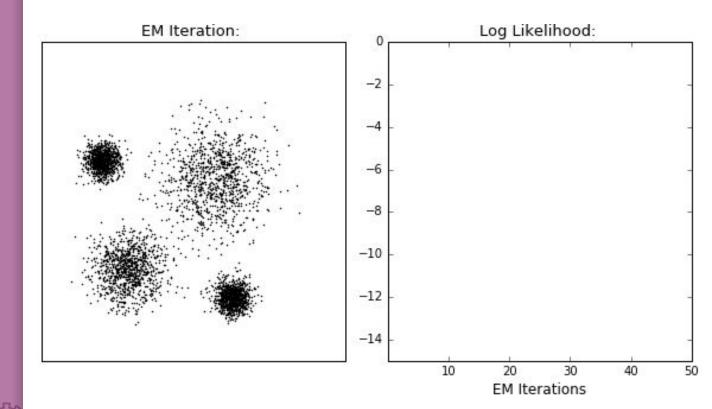




different shaped/oriented clusters

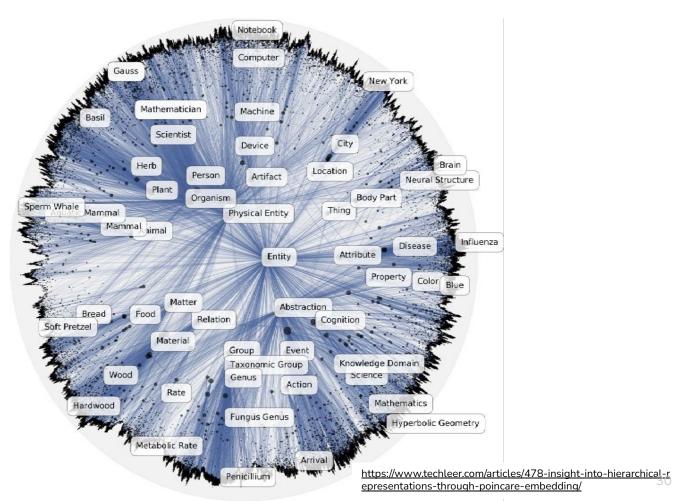


Visualizing Gaussian Mixture Models



### Nouns





# Types of Algorithms

#### Divisive, a.k.a. top-down

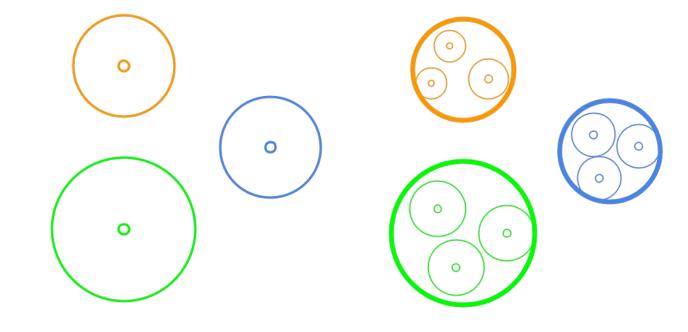
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- Start with each data point in its own cluster. Merge clusters until all points are in one big cluster.
  - Example: single linkage clustering

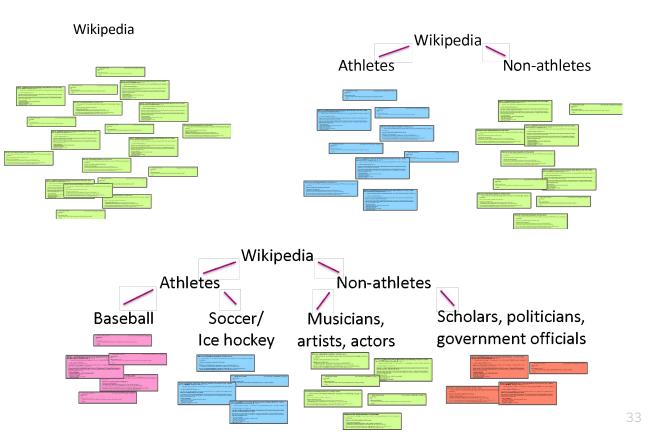
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Start with all the data in one cluster, and then repeatedly run k-means to divide the data into smaller clusters. Repeatedly run k-means on each cluster to make sub-clusters.



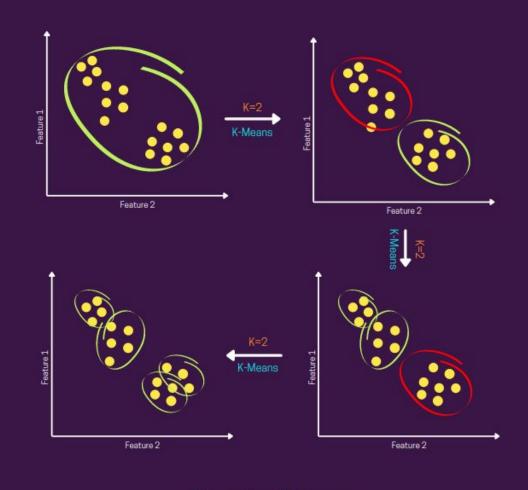
### Example

Using Wikipedia

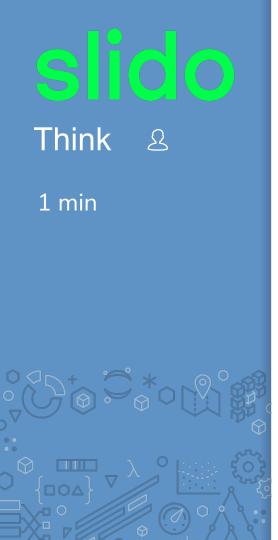


# Bisecting K-Means

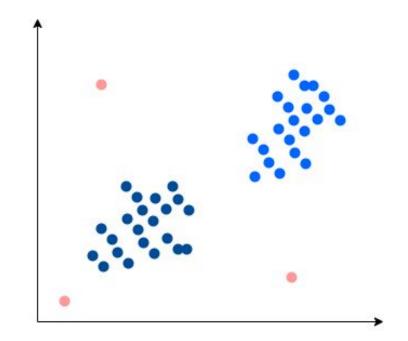


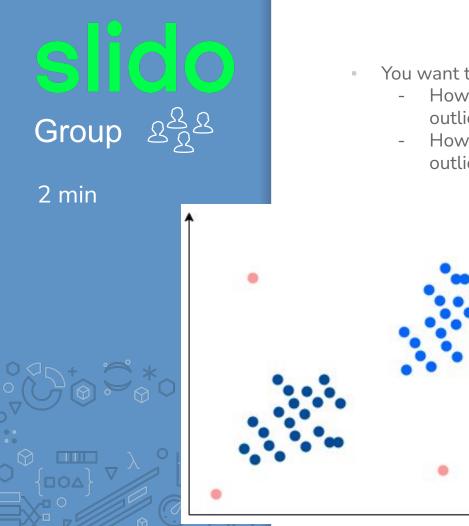


**Bisecting K-Means** 

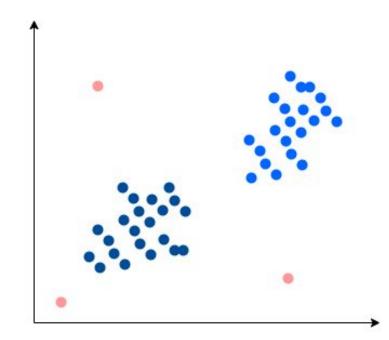


- You want to detect outliers in a dataset (shown below).
  - How would you use k-means clustering to detect outliers?
  - How would you use divisive clustering to detect outliers?

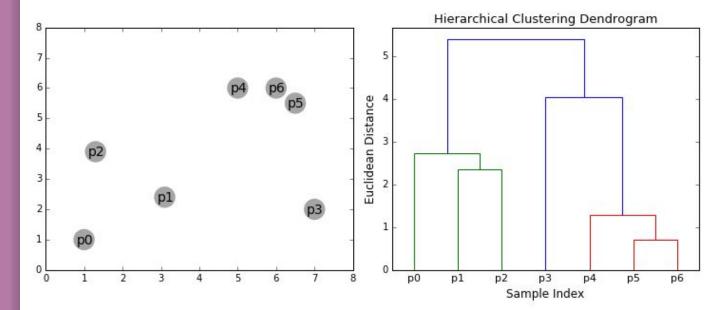




- You want to detect outliers in a dataset (shown below).
  - How would you use k-means clustering to detect outliers?
  - How would you use divisive clustering to detect outliers?



 $\nabla$ 



#### Algorithm at a glance

- 1. Initialize each point in its own cluster
- 2. Define a distance metric between clusters

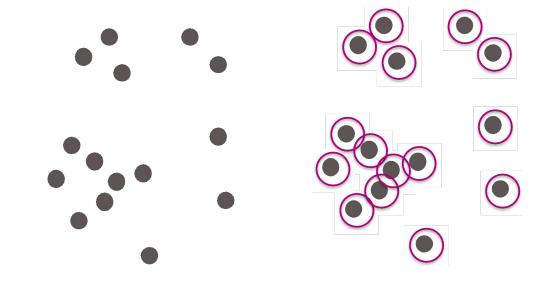
While there is more than one cluster

3. Merge the two closest clusters

## Step 1



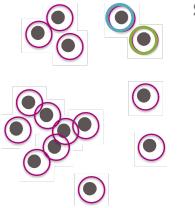
#### 1. Initialize each point to be its own cluster



## Step 2



#### 2. Define a distance metric between clusters



Single Linkage distance( $C_1$ ,  $C_2$ ) =  $\min_{x_i \in C_1, x_j \in C_2} d(x_i, x_j)$ 

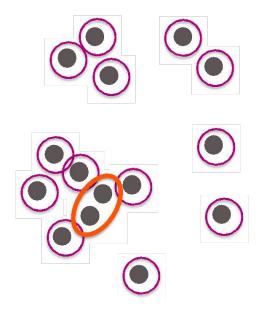
This formula means we are defining the distance between two clusters as the smallest distance between any pair of points between the clusters.



# Step 3

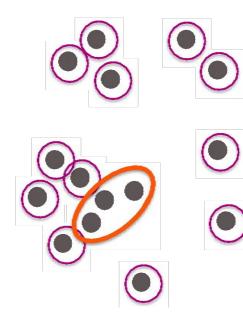


#### Merge closest pair of clusters



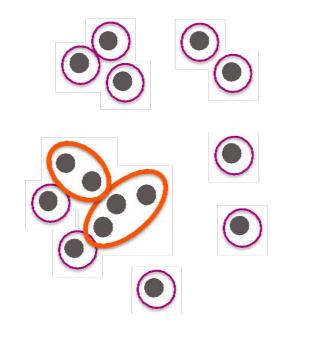
#### 



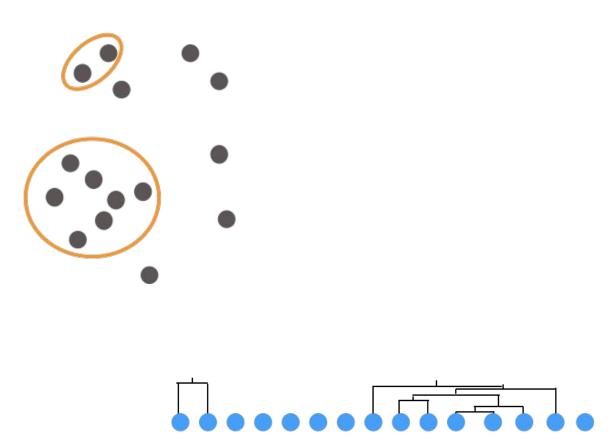




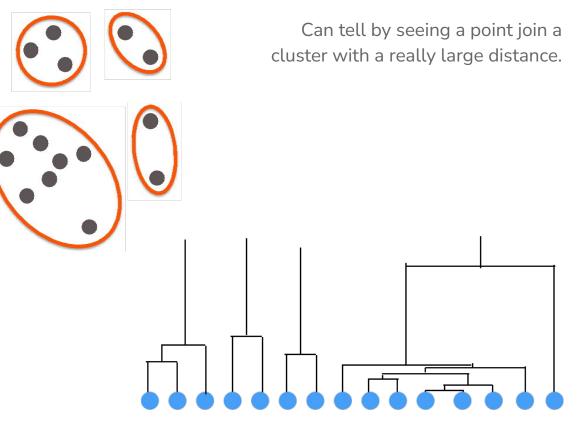
Notice that the height of the dendrogram is growing as we group points farther from each other



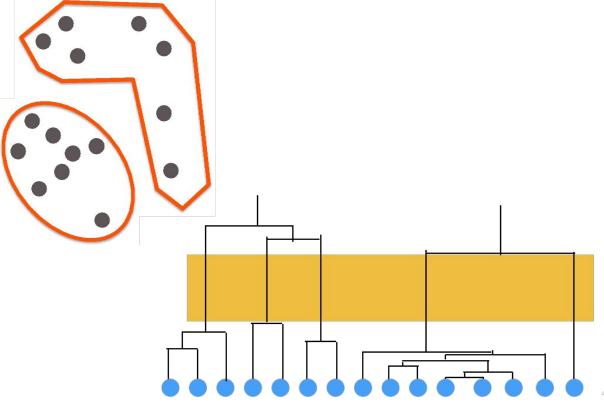




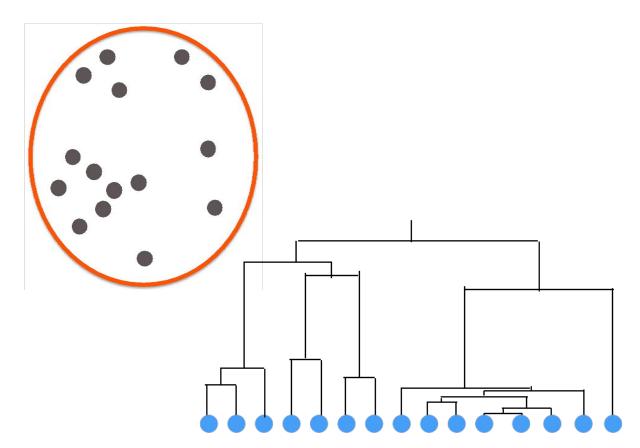
Looking at the dendrogram, we can see there is a bit of an outlier!



The tall links in the dendrogram show us we are merging clusters that are far away from each other

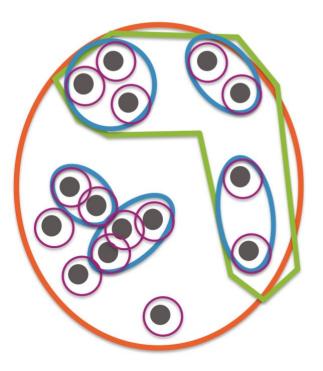


Final result after merging all clusters



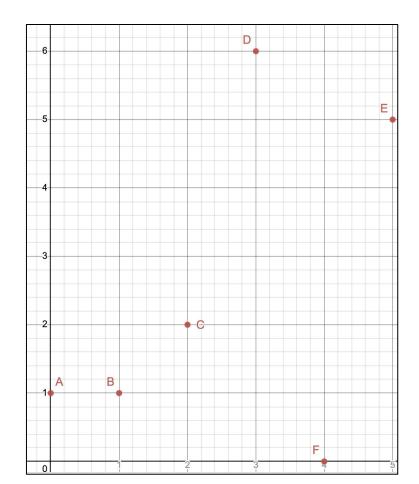
# Final Result

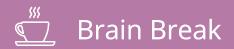






In what order will the following points get merged into clusters? Use L2 (Euclidean) distance, and the single linkage function.

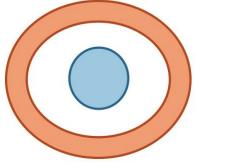






# Dendrograms

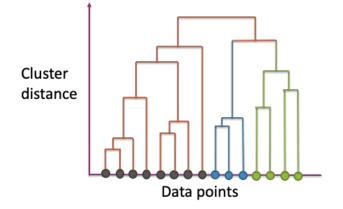
 With agglomerative clustering, we are now very able to learn weirder clusterings like





## Dendrogram

x-axis shows the datapoints (arranged in a very particular order) y-axis shows distance between merged clusters

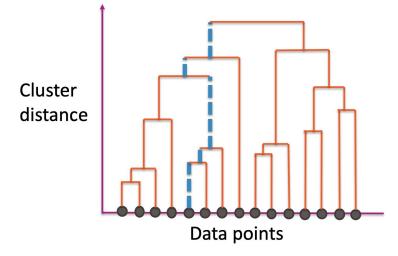






## Dendrogram

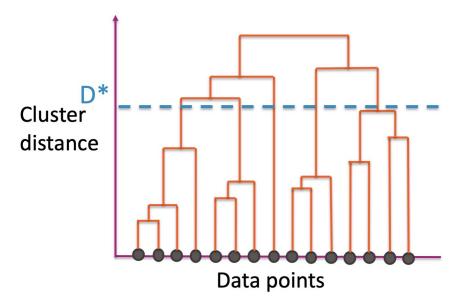
The path shows you all clusters that a single point belongs and the order in which its clusters merged

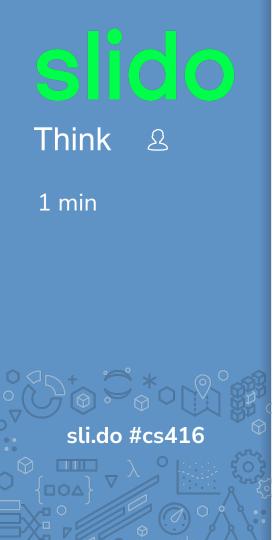


# Cut Dendrogram

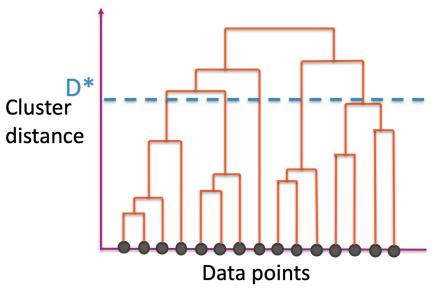
Choose a distance  $D^*$  to "cut" the dendrogram

- Use the largest clusters with distance < D\*</p>
- Usually ignore the idea of the nested clusters after cutting





How many clusters would we have if we use this threshold?

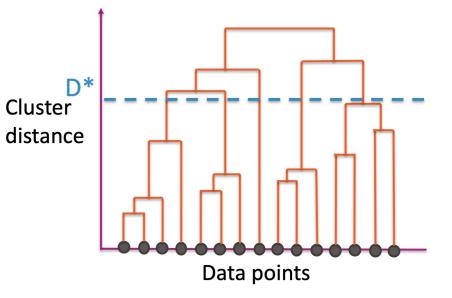




2 min

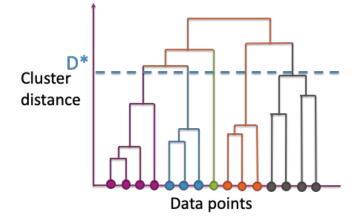


How many clusters would we have if we use this threshold?



# Cut Dendrogram

Every branch that crosses  $D^*$  becomes its own cluster





# Choices to Make

 For agglomerative clustering, you need to make the following choices:

- Distance metric  $d(x_i, x_j)$
- Linkage function
  - Single Linkage:

$$D(C_1, C_2) = \min_{x_i \in C_1, x_j \in C_2} d(x_i, x_j)$$

Complete Linkage:

$$D(C_1, C_2) = \max_{x_i \in C_1, x_j \in C_2} d(x_i, x_j)$$

Centroid Linkage

$$\mathsf{D}(\mathcal{C}_1,\mathcal{C}_2) = d(\mu_1,\mu_2)$$

- Others

Cluster distance

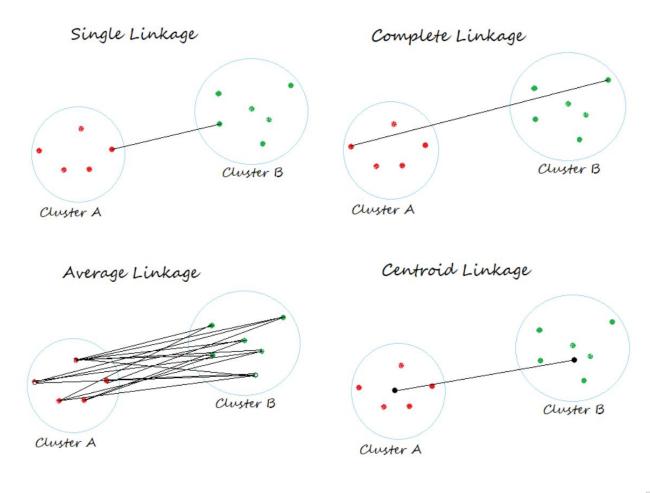
D

Where and how to cut dendrogram

Data points

# Linkage Functions





# Practical Notes

For visualization, generally a smaller # of clusters is better

For tasks like outlier detection, cut based on:

- Distance threshold
- Or some other metric that tries to measure how big the distance increased after a merge

No matter what metric or what threshold you use, no method is "incorrect". Some are just more useful than others.



Computational Cost of Agglomerative Clustering

Computing all pairs of distances is pretty expensive!

• A simple implementation takes  $O(n^2 \log(n))$ 

Can be much implemented more cleverly by taking advantage of the **triangle inequality** 

"Any side of a triangle must be less than the sum of its sides"

Best known algorithm is  $\mathcal{O}(n^2)$ 

k-means vs. Agglomerative Clustering

- K-means is more efficient on big data than hierarchical clustering.
- Initialization changes results in k-means, not in agglomerative clustering has reproducible results.
- K-means works well only for hyper-spherical clusters, agglomerative clustering can handle more complex cluster shapes.
- K-means requires selecting a number of clusters beforehand.
  In agglomerative clustering, you can decide on the number of clusters afterwards using the dendrogram.

# Concept Inventory

This week we want to practice recalling vocabulary. Spend 10 minutes trying to write down all the terms for concepts we have learned in this class and try to bucket them into the following categories.

Regression

Classification

**Deep Learning** 

**Document Retrieval** 

Misc – For things that fit in multiple places or none of the above

You don't need to define/explain the terms for this exercise, but you should know what they are!

Try to do this for at least 5 minutes from recall before looking at your notes!

# Recap

- Problems with k-means
- Mixture Models
- Hierarchical clustering
- Divisive Clustering
- Agglomerative Clustering
- Dendrograms

