

Pre-Class Video 1

Clustering Reca

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**

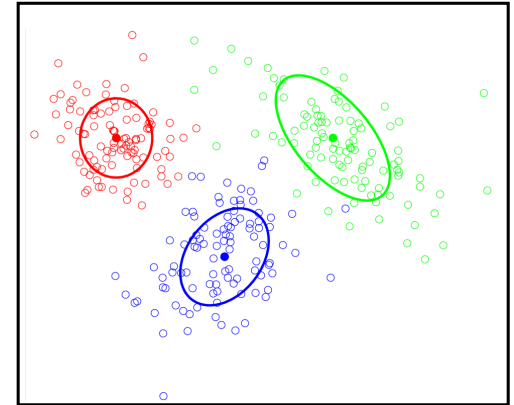
Clustering is the process of finding these clusters and **assigning** each example to a particular cluster.

x_i gets assigned $z_i \in [1, 2, \dots, k]$

Usually based on closest centroid

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

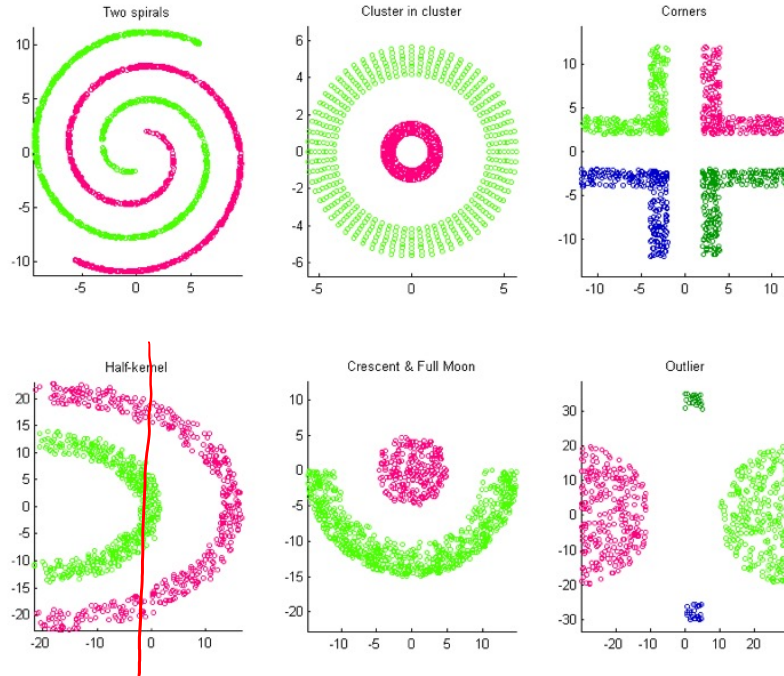
Based on distance of assigned examples to each cluster



Not Always Easy

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

Distance does not determine clusters

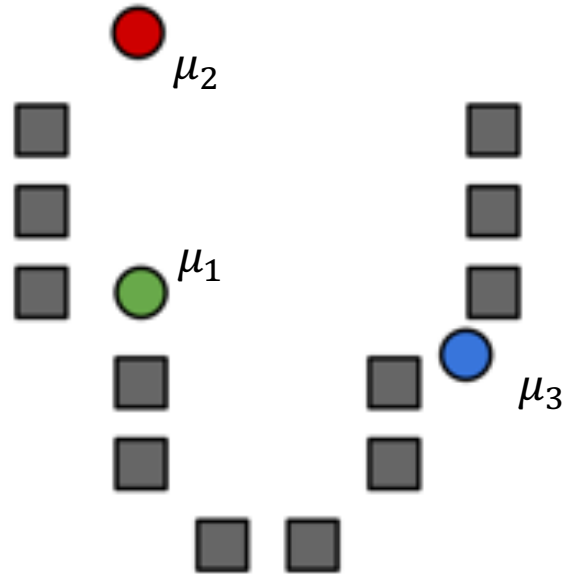


Step 0

Start by choosing the initial cluster centroids

A common default choice is to choose centroids at random

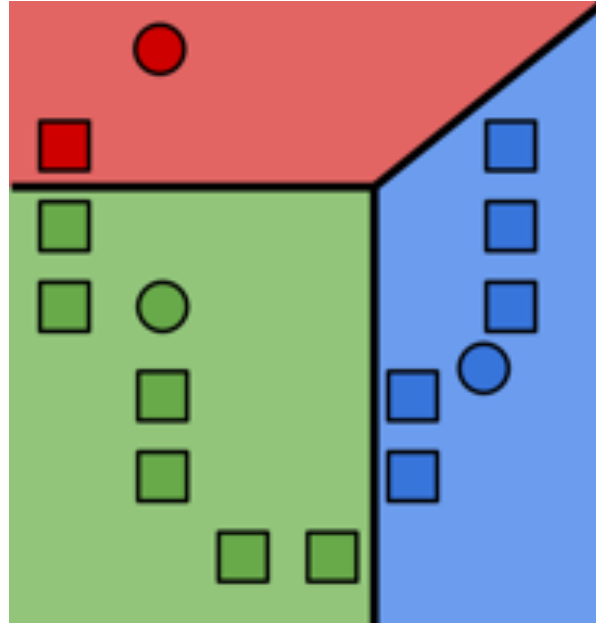
Will see later that there are smarter ways of initializing



Step 1

Assign each example to its closest cluster centroid

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

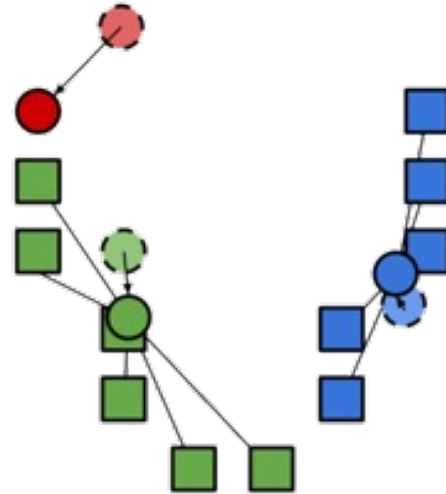


Step 2

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

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

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

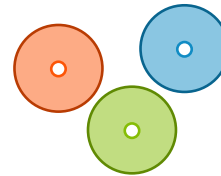
1. Choose first cluster μ_1 from the data uniformly at random
2. For the current set of centroids (starting with just μ_1), compute the distance between each datapoint and its closest centroid
3. Choose a new centroid from the remaining data points with probability of x_i being chosen proportional to $d(x_i)^2$
4. Repeat 2 and 3 until we have selected k centroids

Problems with k-means

In real life, cluster assignments are not always clear cut

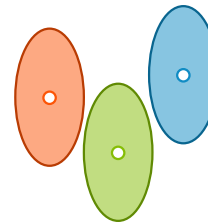
E.g. The moon landing: Science? World News? Conspiracy?

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



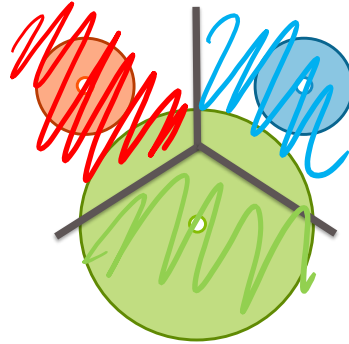
We can change this with weighted Euclidean distance

Still assumes every cluster is the same shape/orientation

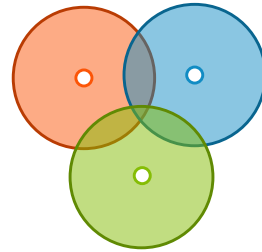


Failure Modes of k-means

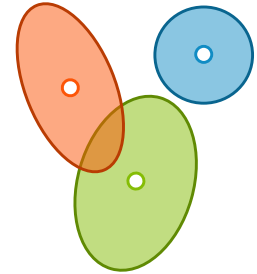
If we don't meet the assumption of spherical clusters, we will get unexpected results



disparate cluster sizes



overlapping clusters



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



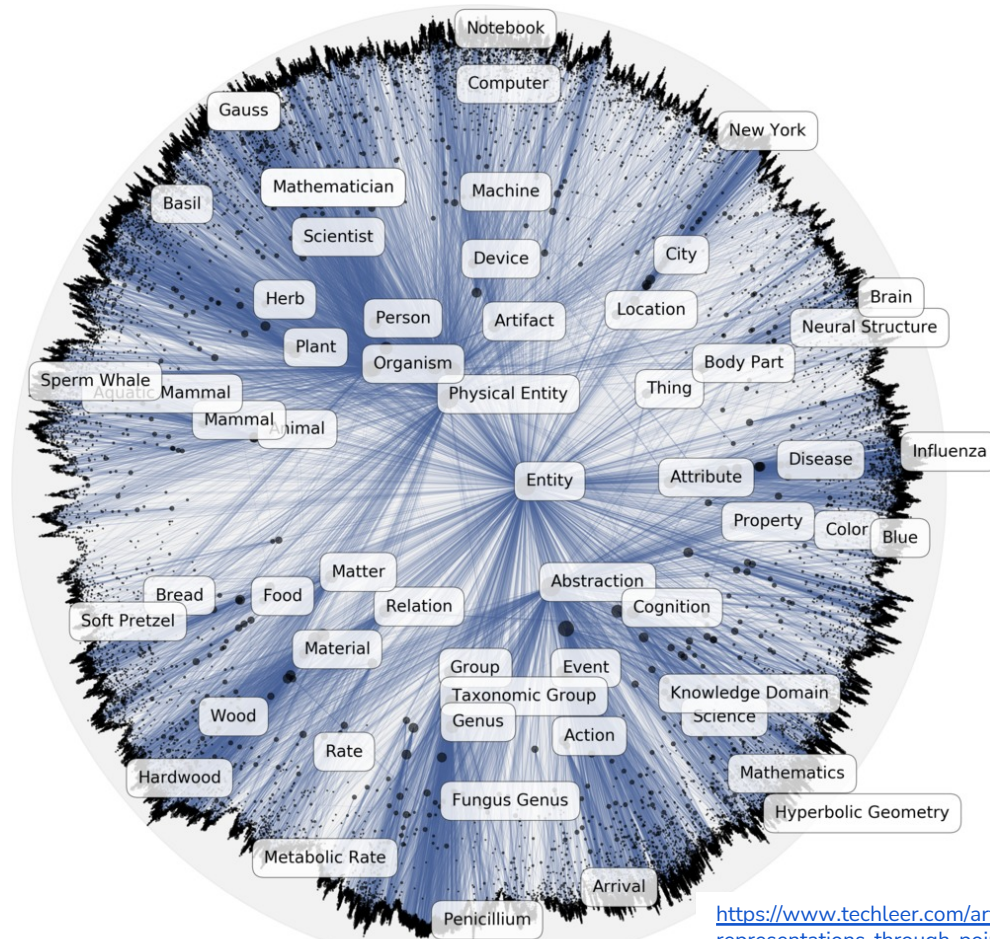
Pre-Class Video 2

Divisive Clustering

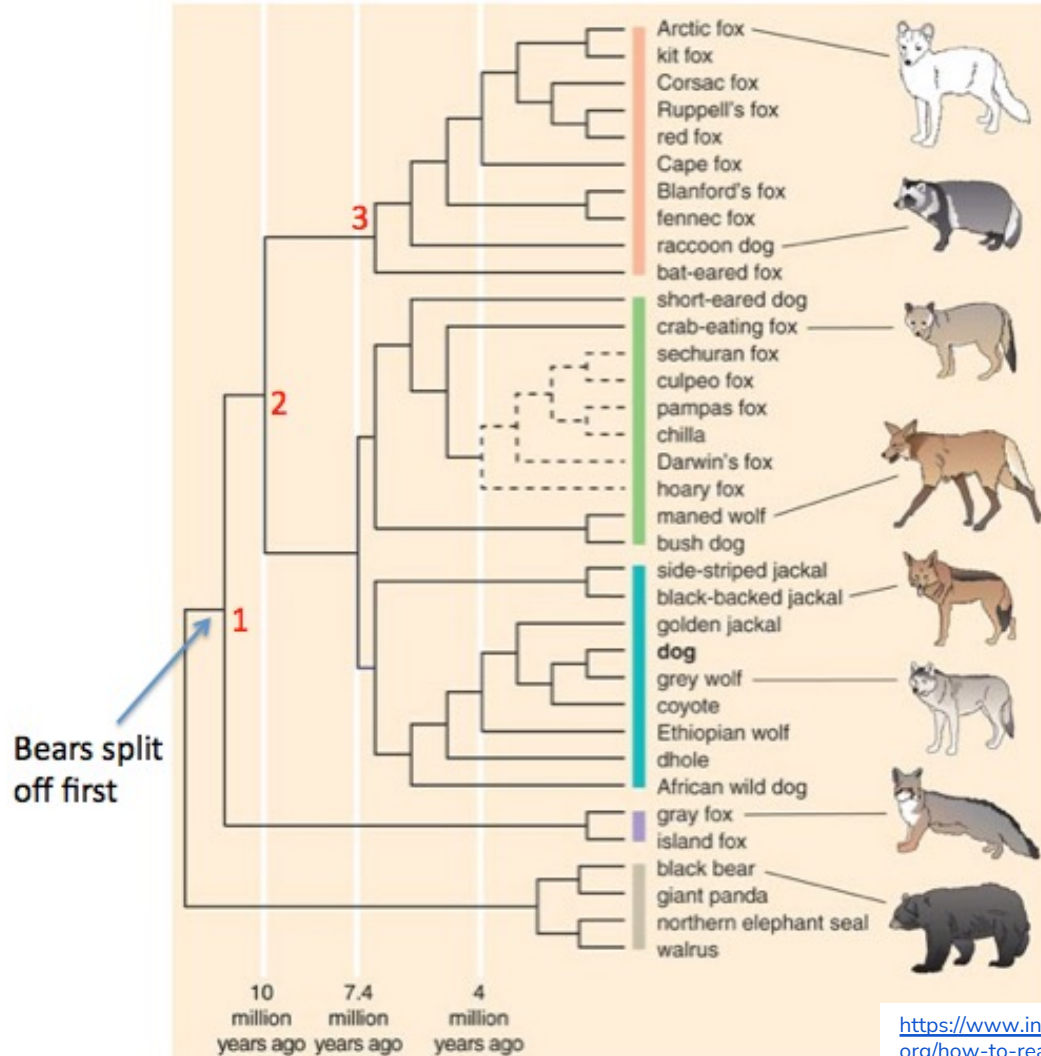
Hierarchical Clustering

Nouns

Lots of data is hierarchical by nature



Species



Motivation

If we try to learn clusters in hierarchies, we can

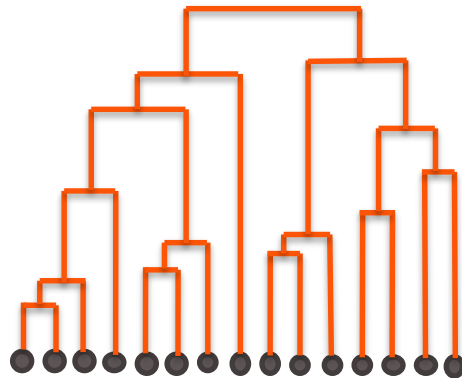
Avoid choosing the # of clusters beforehand

Use dendrograms to help visualize different granularities of clusters

Allow us to use any distance metric

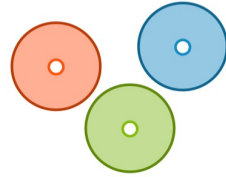
- K-means requires Euclidean distance

Can often find more complex shapes than k-means

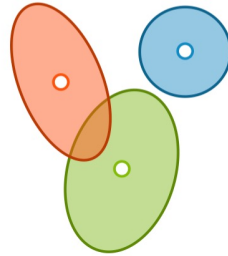


Finding Shapes

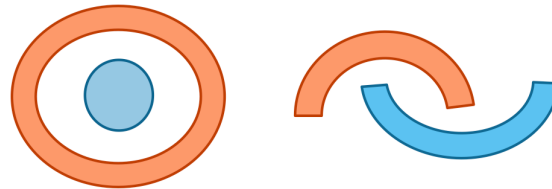
k-means



Mixture Models



Hierarchical Clustering



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*:

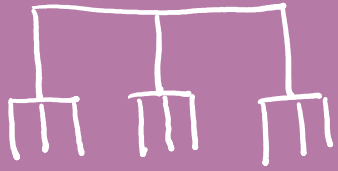
Start with each data point in its own cluster. Merge clusters until all points are in one big cluster.

- Example: single linkage clustering

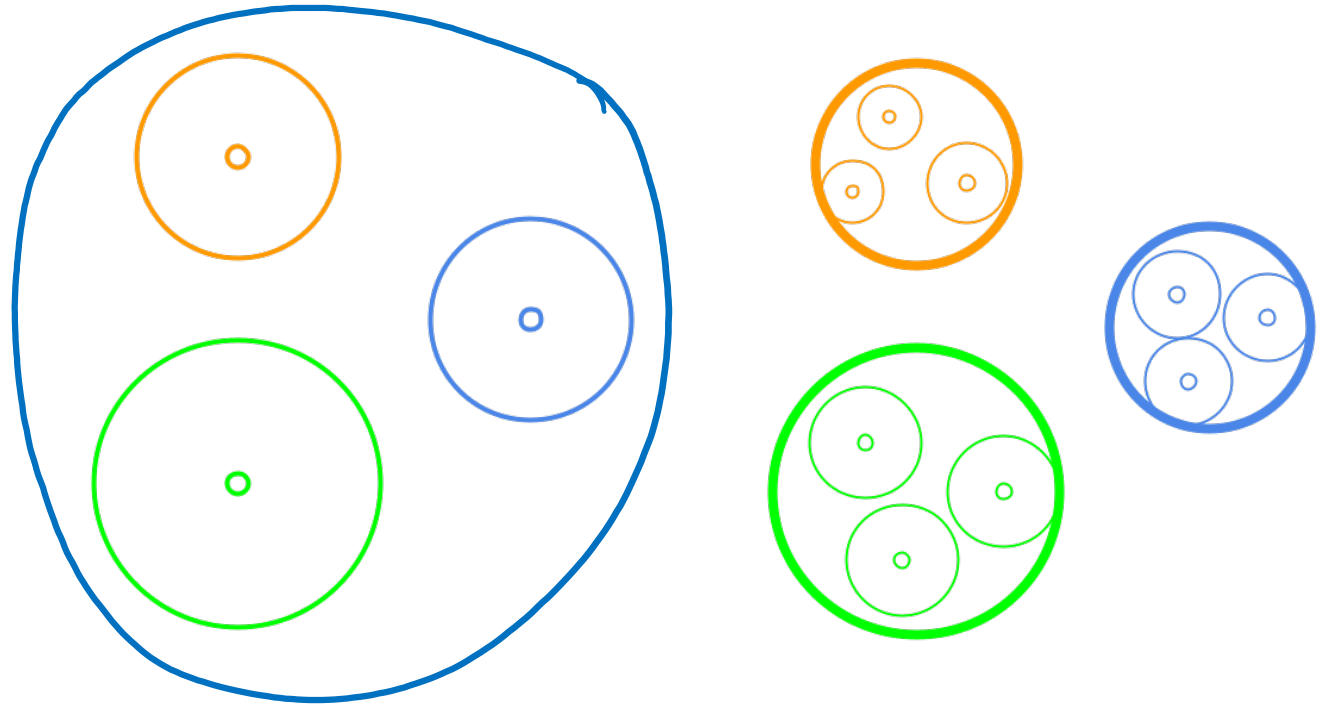
→ how we measure distance between clusters

Divisive Clustering

$k=3$



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.



Using Wikipedia

Example

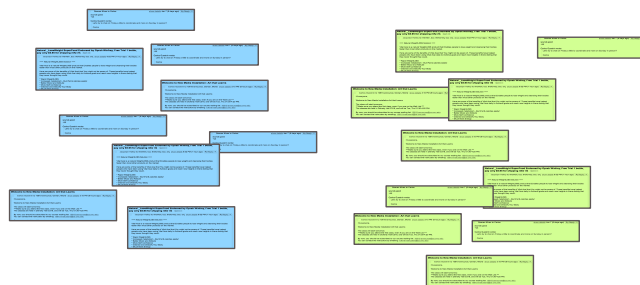
$K=2$

Bisecting
K-means

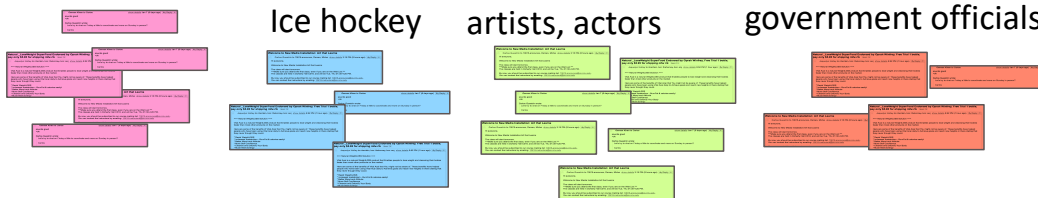
Wikipedia



Wikipedia
Athletes Non-athletes



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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

Hunter Schafer
Paul G. Allen School of Computer Science & Engineering
University of Washington

May 17, 2023

? Questions? Raise hand or sli.do #cs416
🎵 Listening to: Still Woozy



Define Clusters

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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.

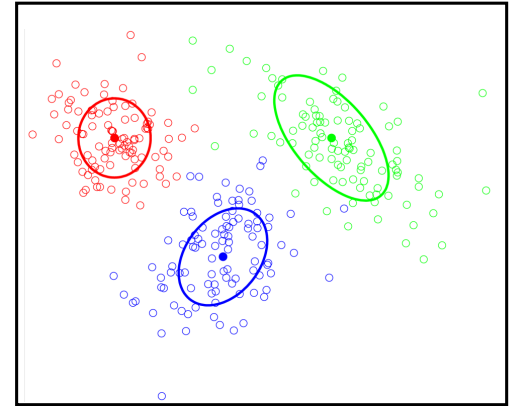
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Usually based on closest centroid

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

Based on distance of assigned examples to each cluster.

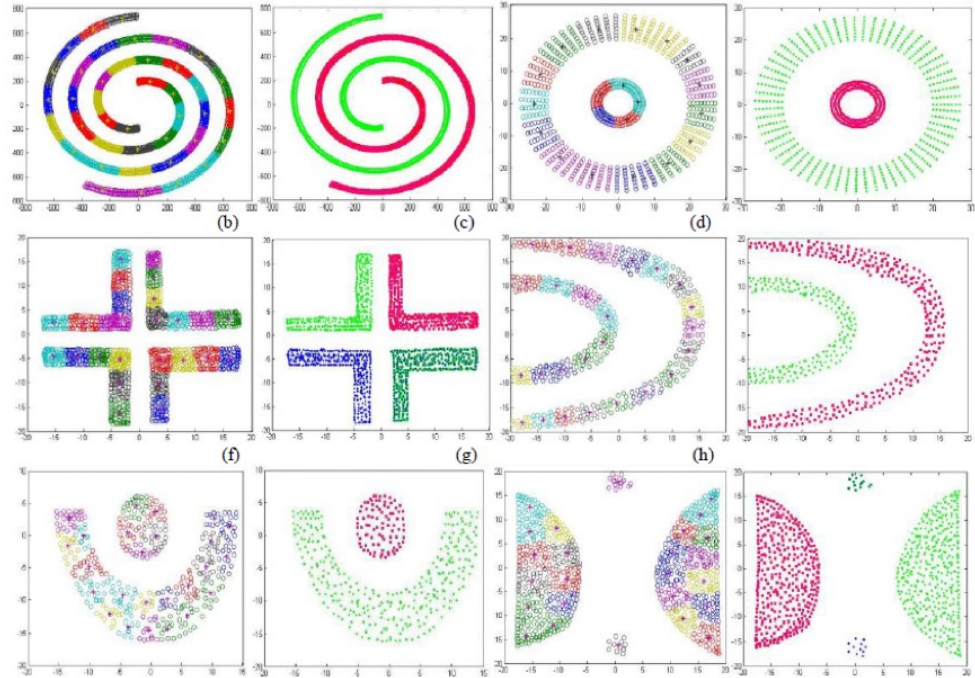
Close distance reflects strong similarity between datapoints.



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-clustering/>



Smart Initializing w/ k-means++

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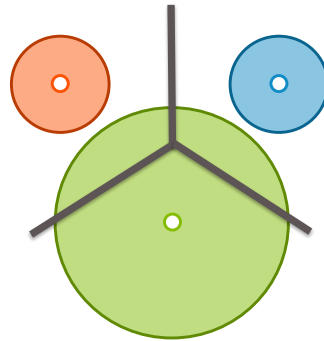
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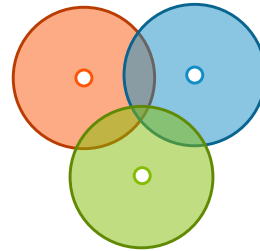
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Failure Modes of k-means

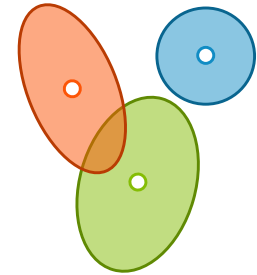
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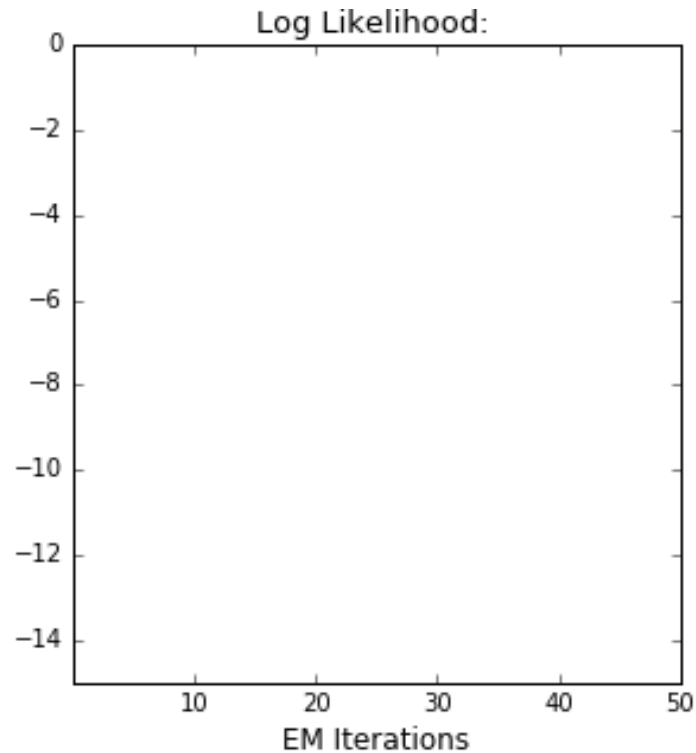
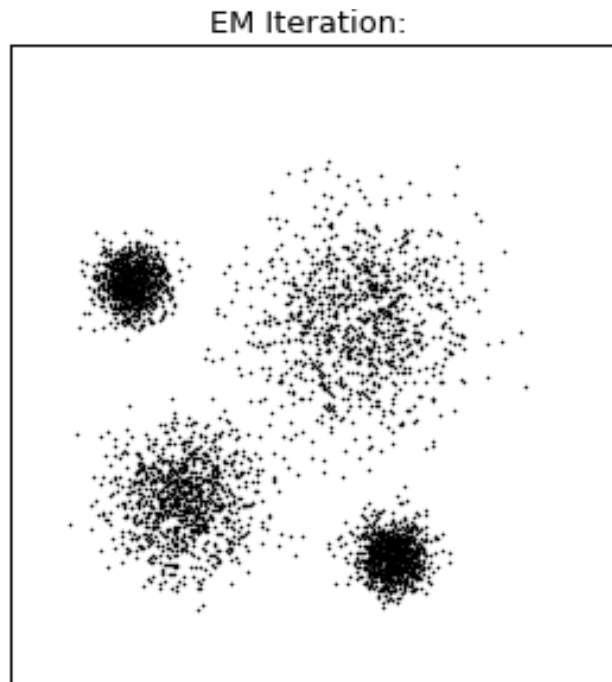
overlapping clusters



different shaped/oriented clusters

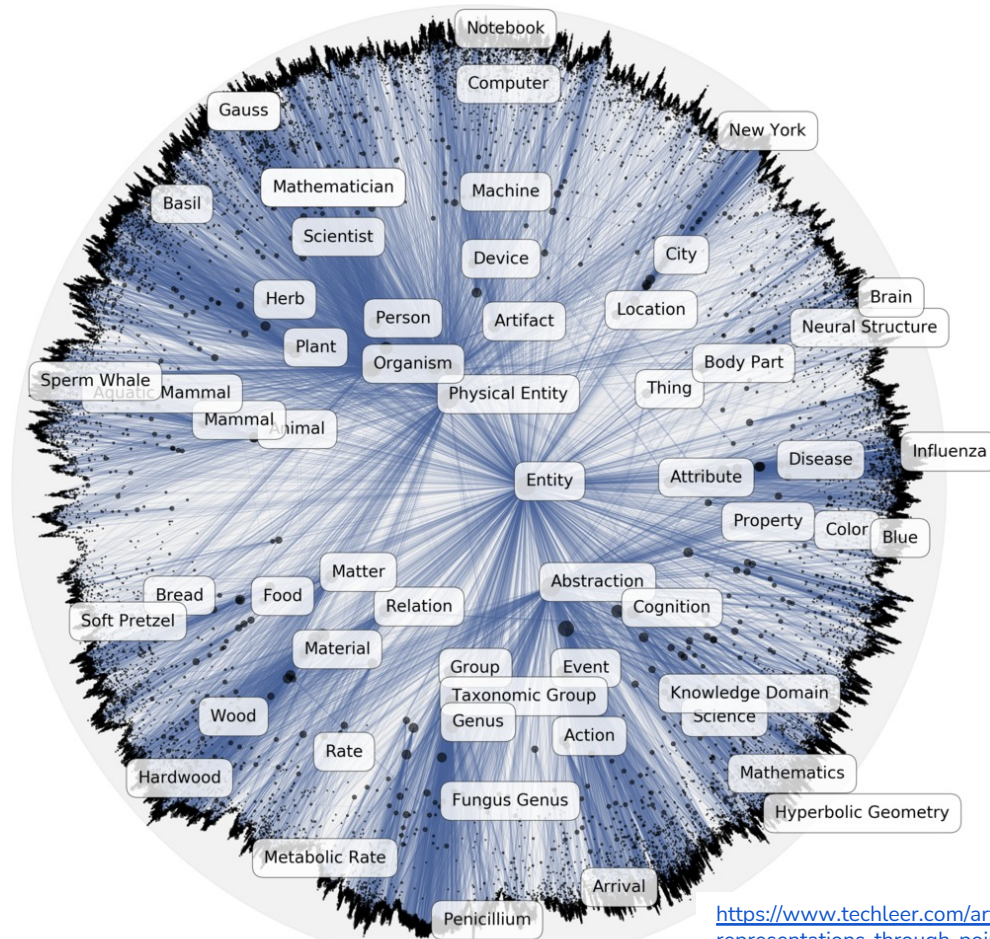


Visualizing Gaussian Mixture Models



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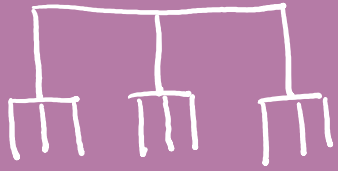
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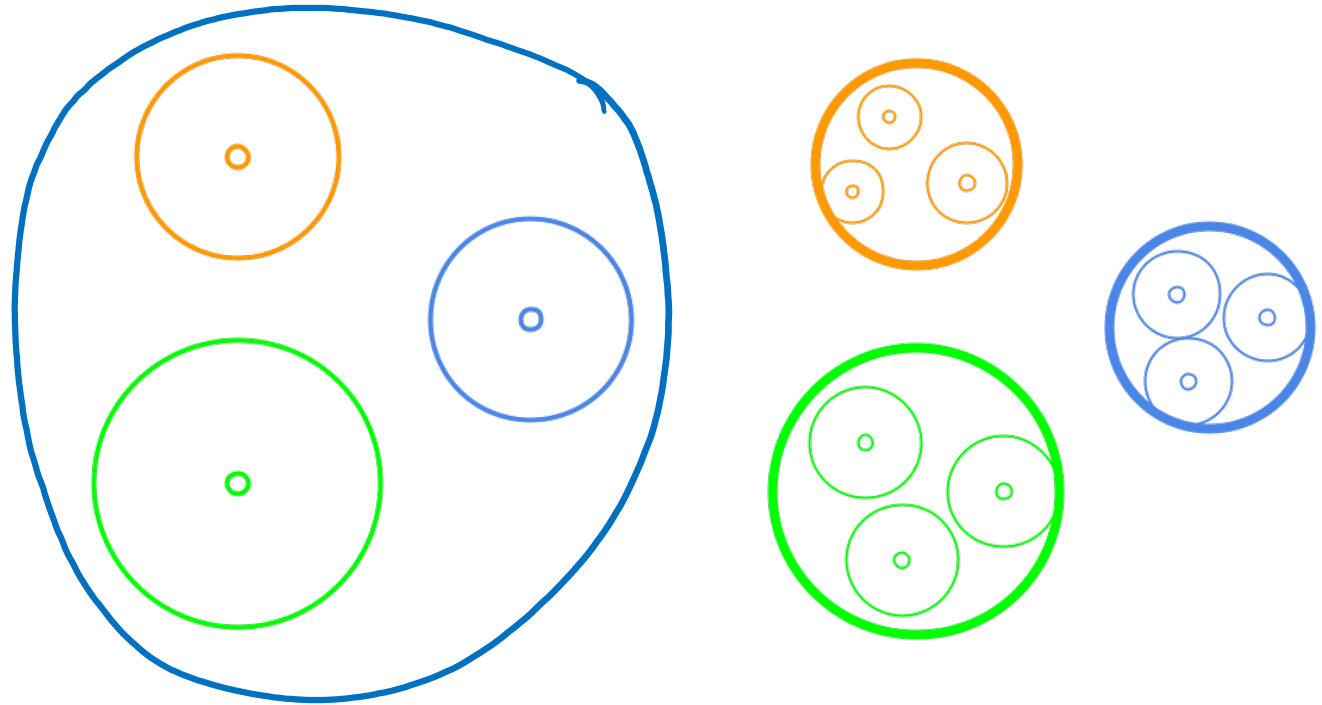
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Divisive Clustering

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Example

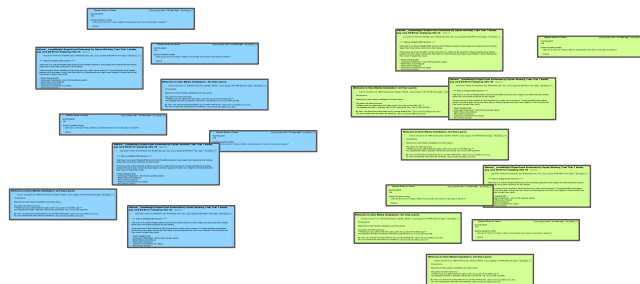
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Bisecting
K-means

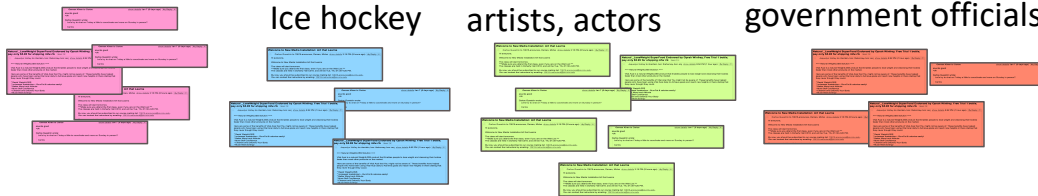
Wikipedia



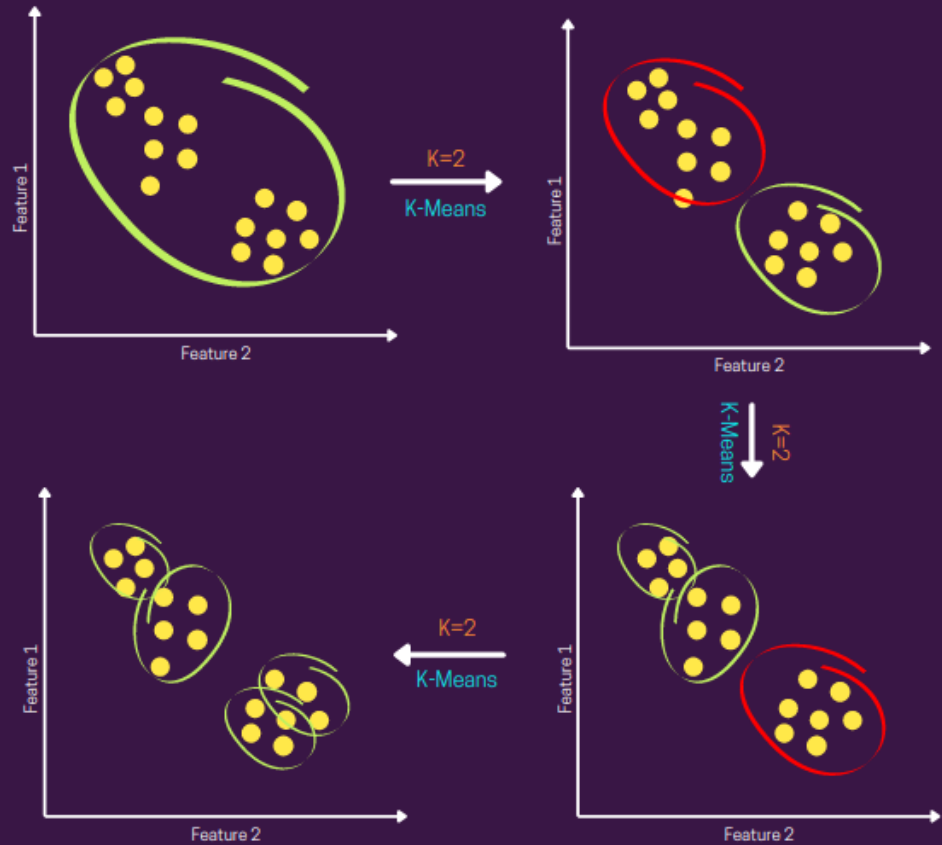
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Bisecting K-Means



Bisecting K-Means

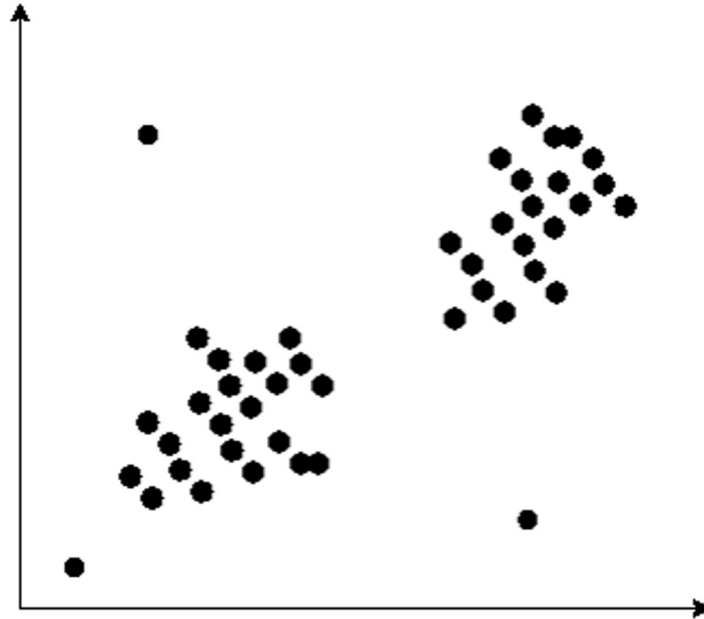
slido

Think 

1 min

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?



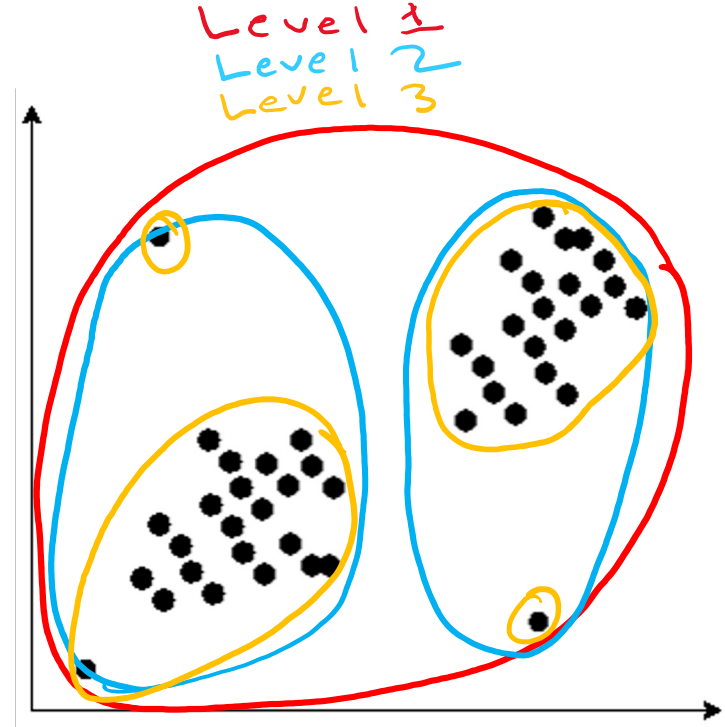
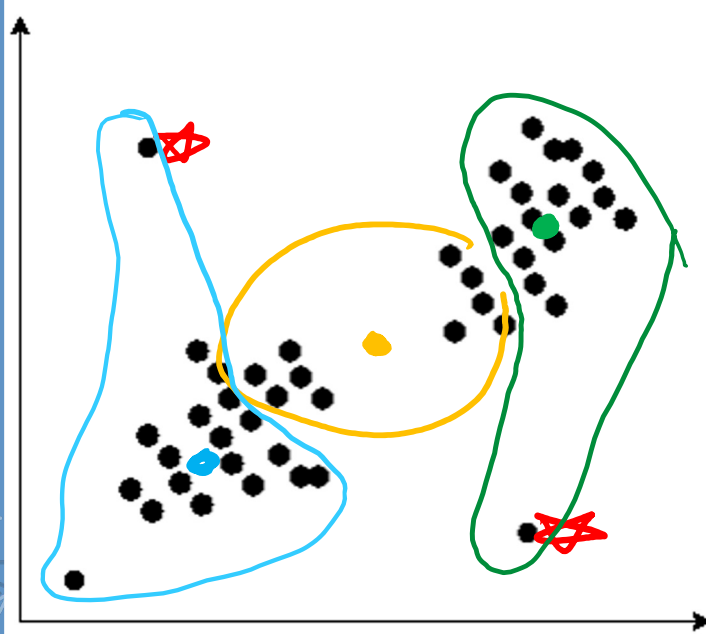
slido

Group 

2 min

You want to detect outliers in a dataset (shown below).

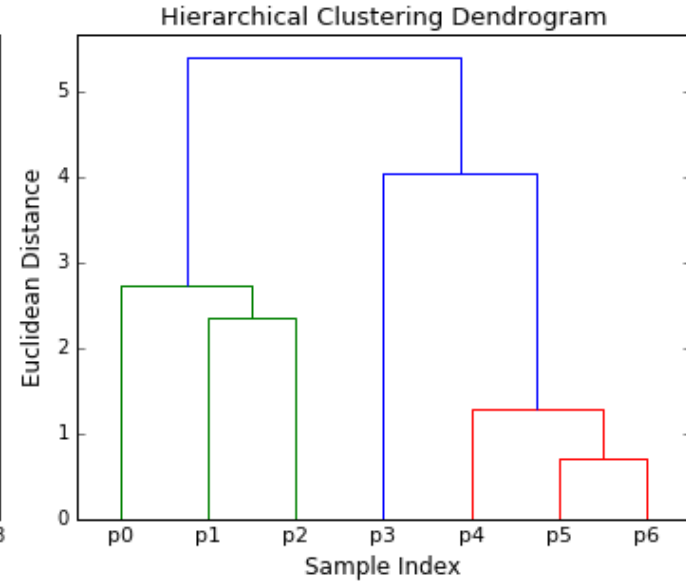
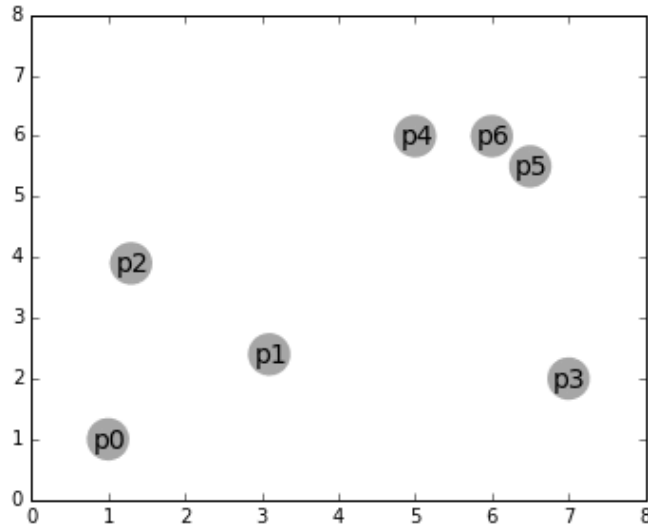
- How would you use k-means clustering to detect outliers?
- How would you use divisive clustering to detect outliers?



Agglomerative Clustering

Merge closest pair of clusters

Agglomerative Clustering



Agglomerative Clustering

Algorithm at a glance

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

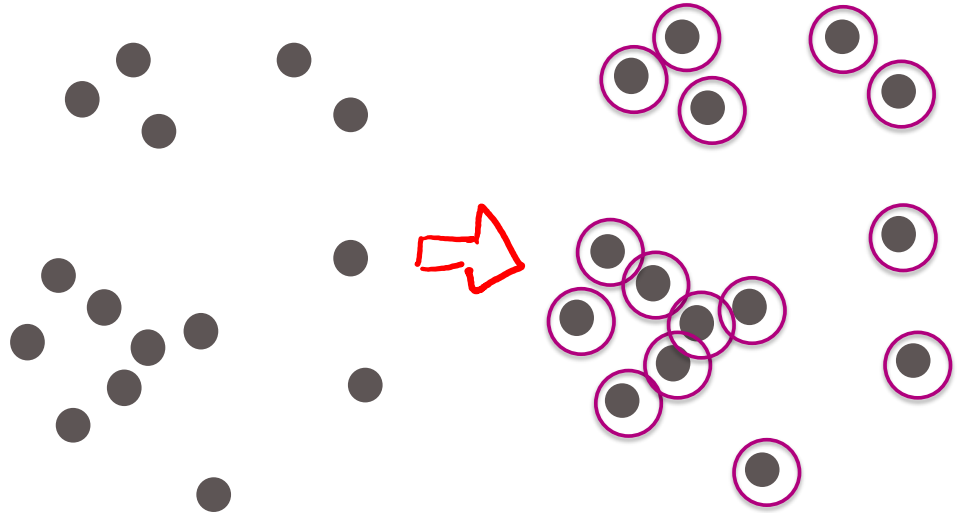
While there is more than one cluster

3. Merge the two closest clusters (and add it to dendrogram)



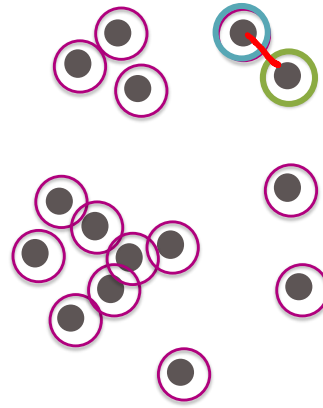
Step 1

1. Initialize each point to be its own cluster



Step 2

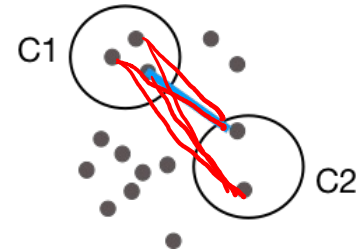
2. Define a distance metric between clusters



Single Linkage

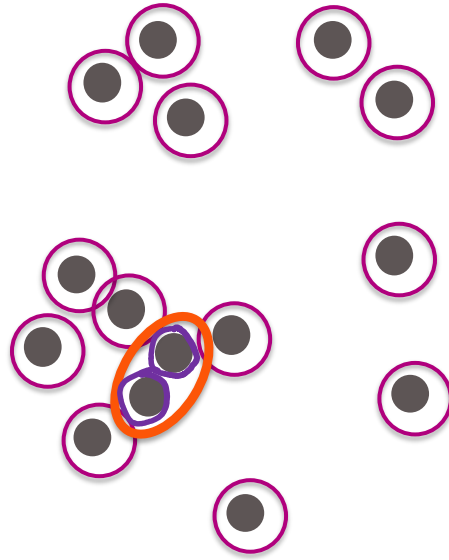
$$distance(C_1, C_2) = \min_{x_i \in C_1, x_j \in C_2} d(x_i, x_j)$$

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

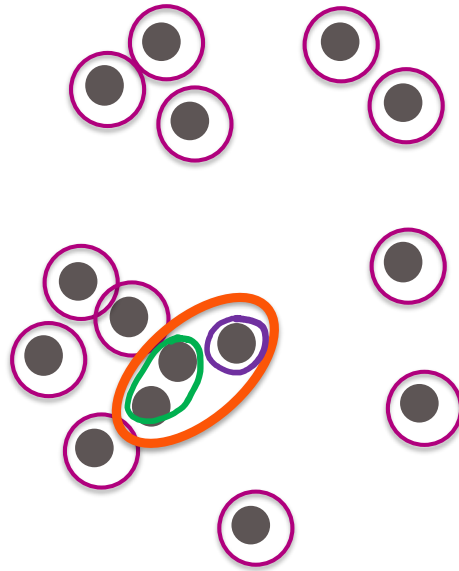


Step 3

Merge closest pair of clusters

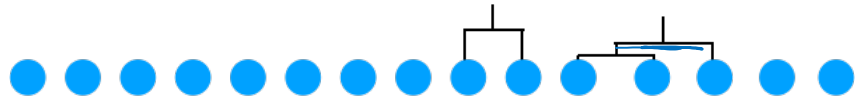
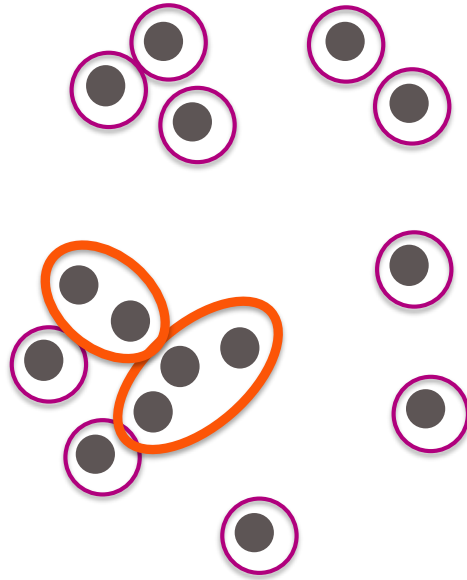


Repeat

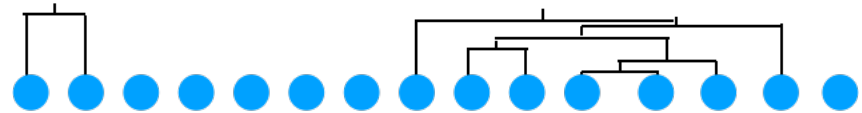
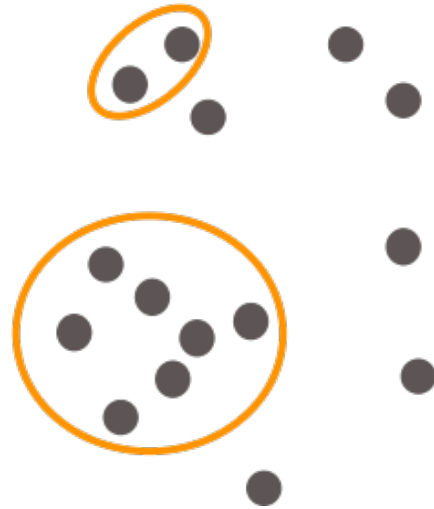


Repeat

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

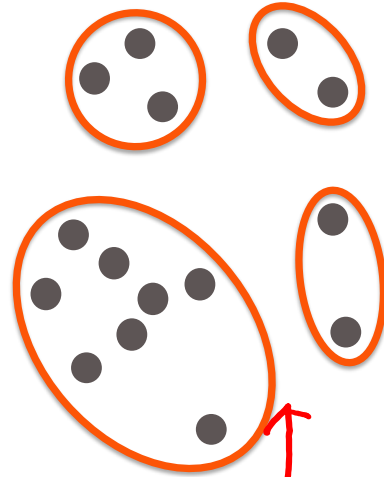


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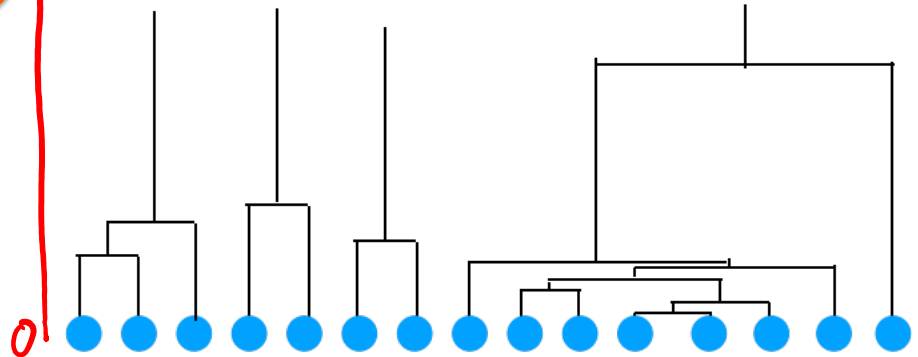
Repeat

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



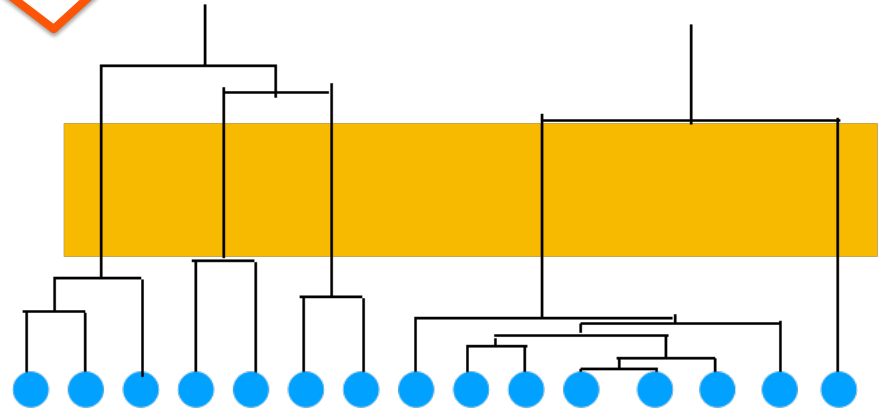
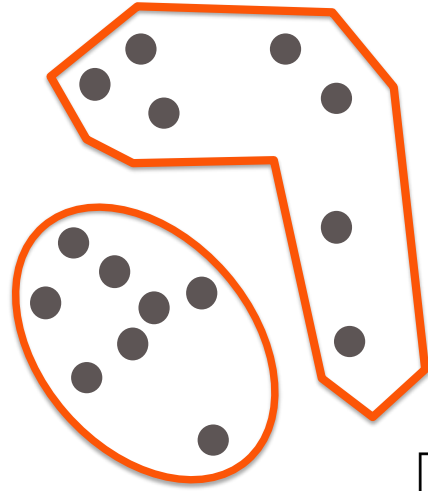
Can tell by seeing a point join a cluster with a really large distance.

Distance
between
the two
merged
clusters



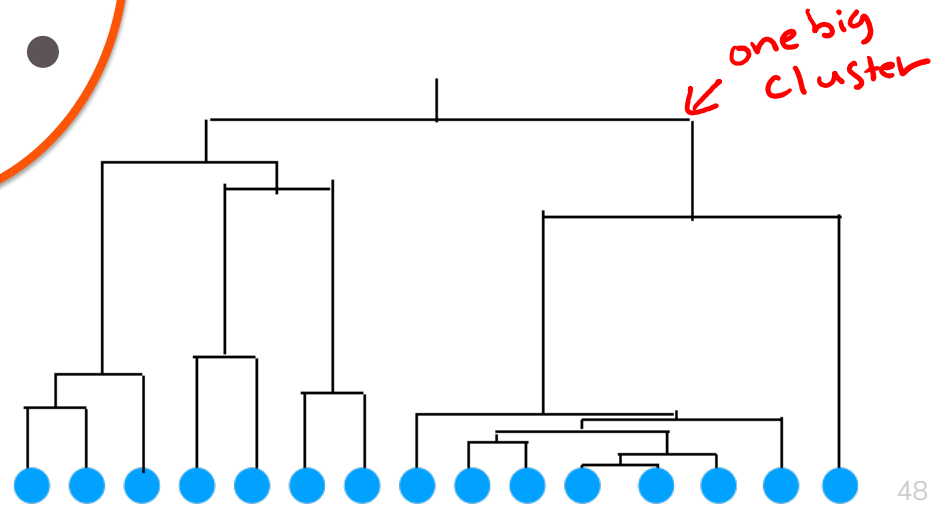
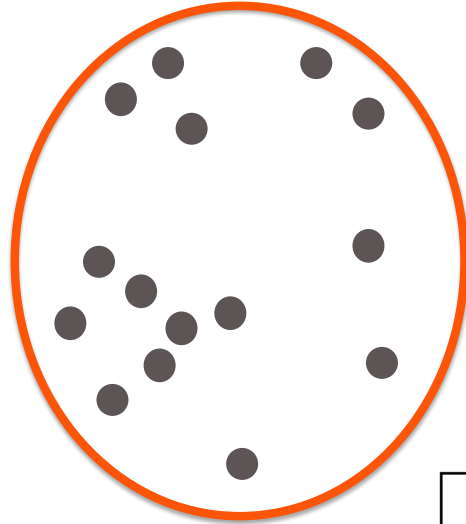
Repeat

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

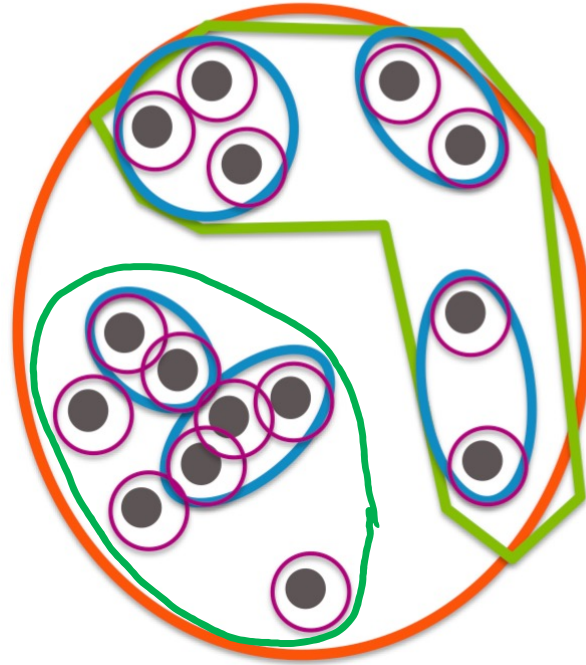


Repeat

Final result after merging all clusters



Final Result

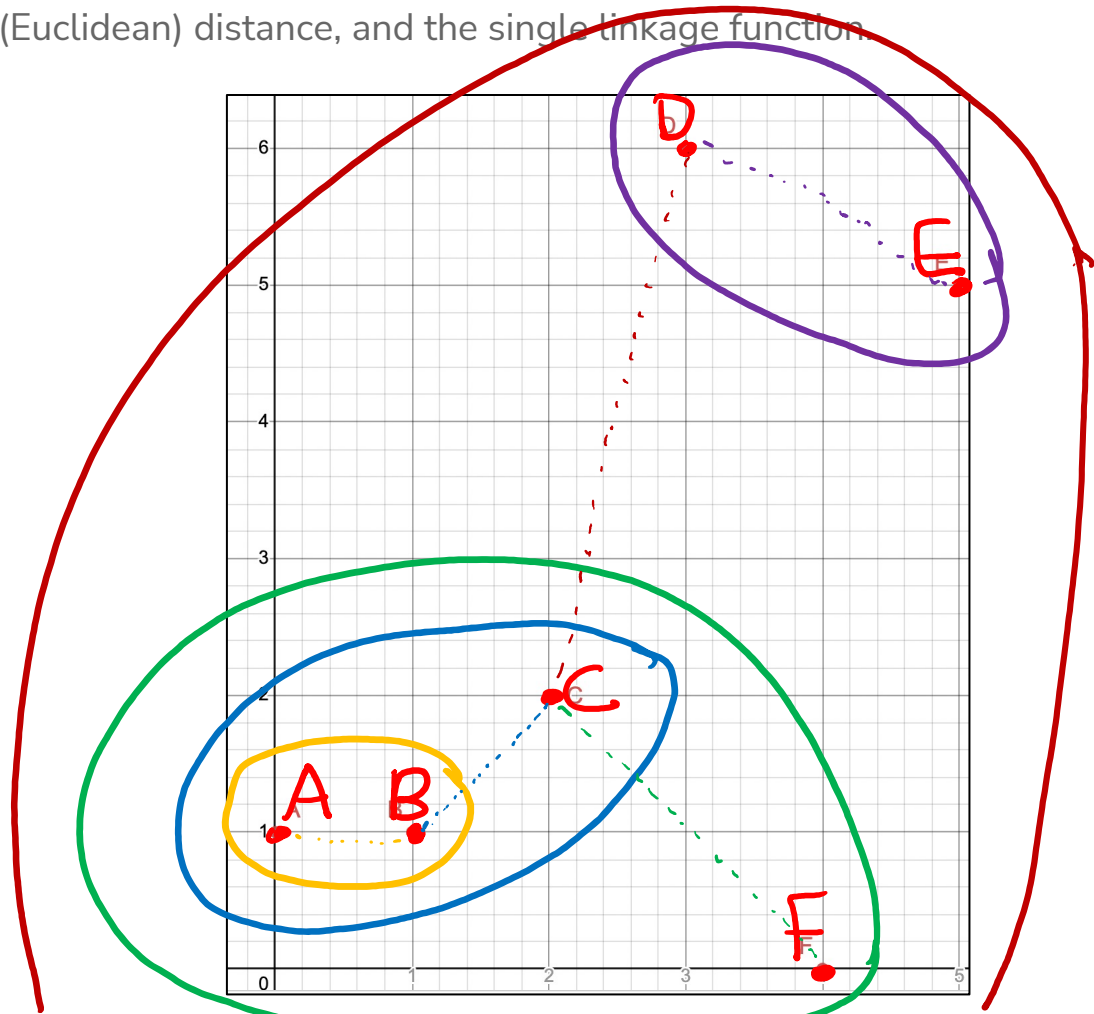


slido

Group 

1 min

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



3:12



Brain Break

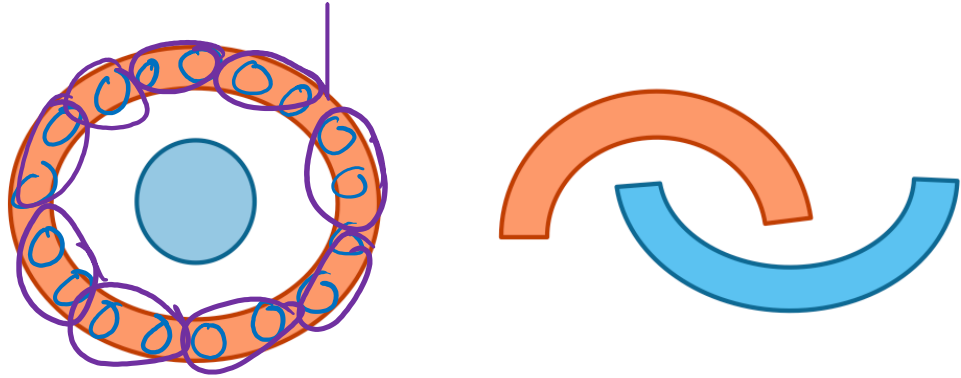


Dendrograms

Agglomerative Clustering

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

Single Linkage: $\min_{x_i \in C_1, x_j \in C_2} \text{dist}(x_i, x_j)$

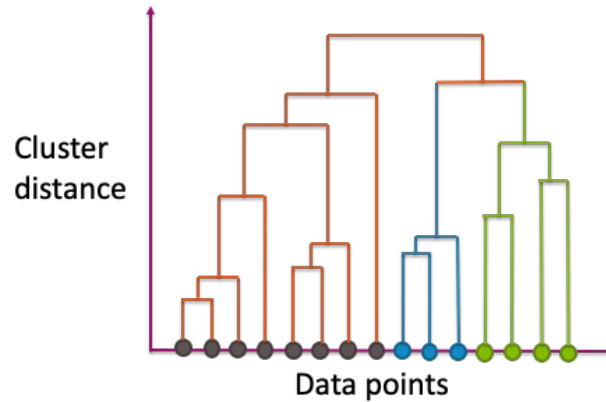


Single Linkage can merge long chains

Dendrogram

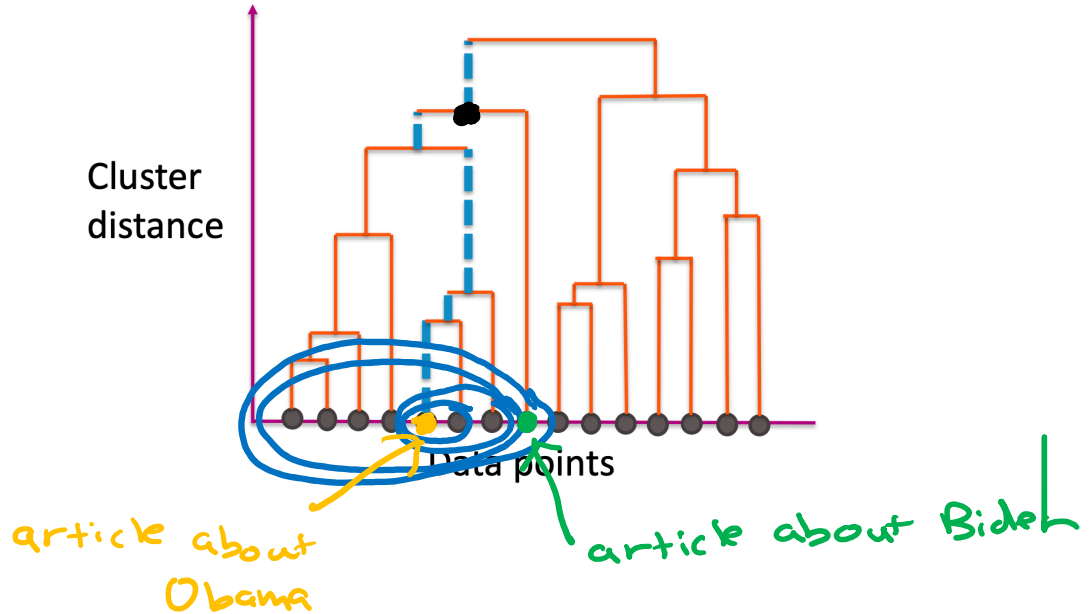
x-axis shows the datapoints (arranged in a very particular order)

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

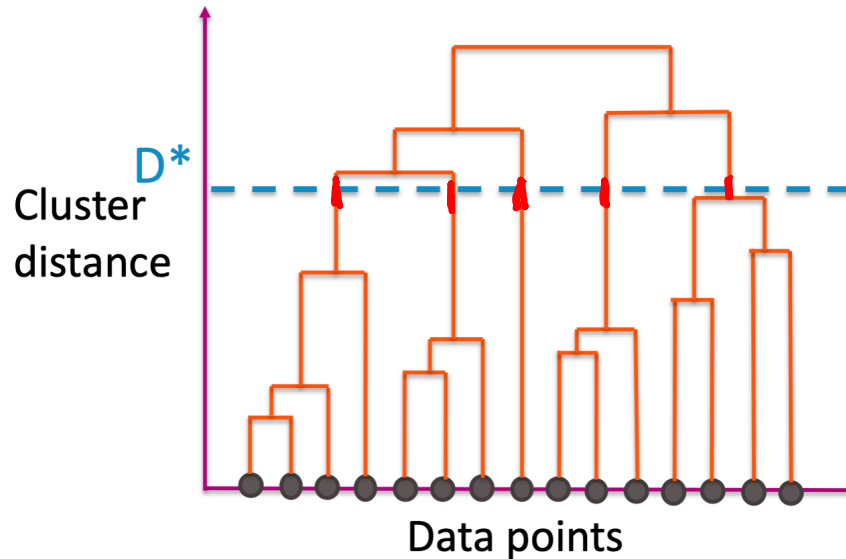


Cut Dendrogram

Choose a distance D^* to “cut” the dendrogram

Use the largest clusters with distance $< D^*$

Usually ignore the idea of the nested clusters after cutting



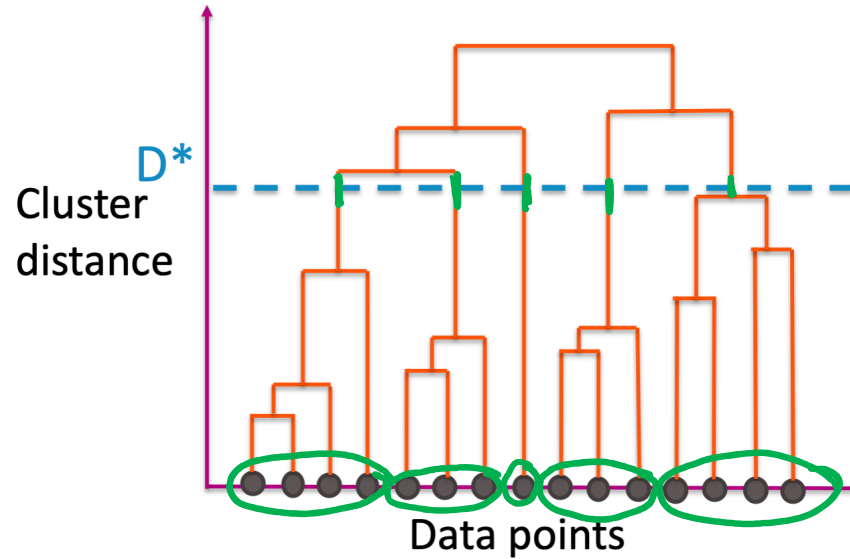
slido

Think 

1 min

slido #cs416

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



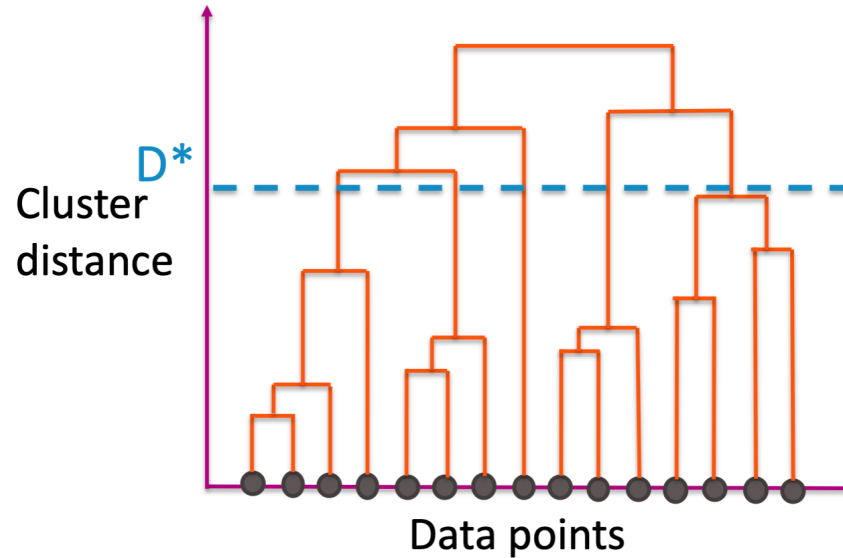
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Group 

2 min

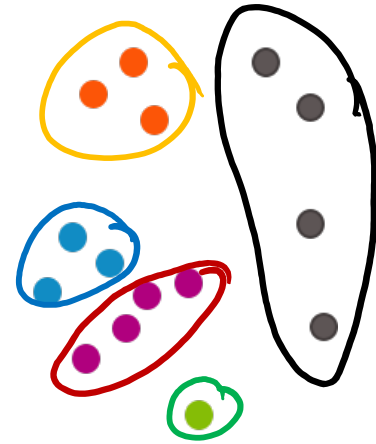
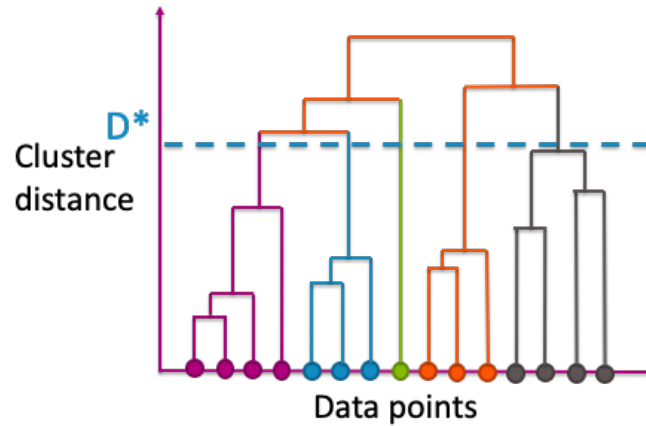
slido #cs416

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:

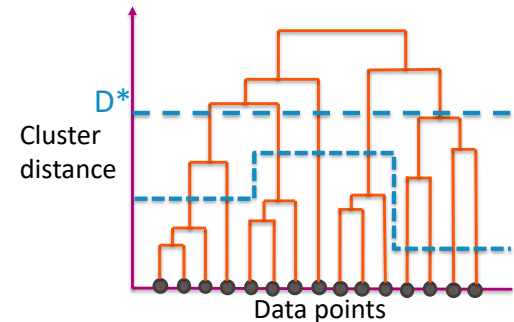
$$D(C_1, C_2) = \max_{x_i \in C_1, x_j \in C_2} d(x_i, x_j)$$

- Centroid Linkage

$$D(C_1, C_2) = d(\mu_1, \mu_2)$$

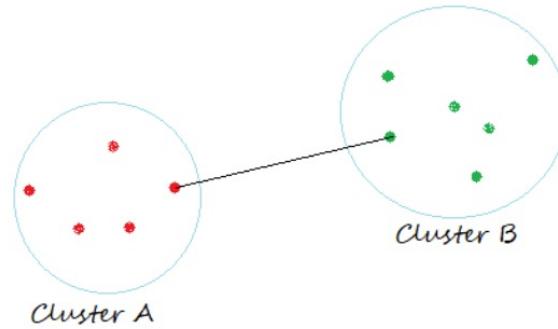
- Others

Where and how to cut dendrogram

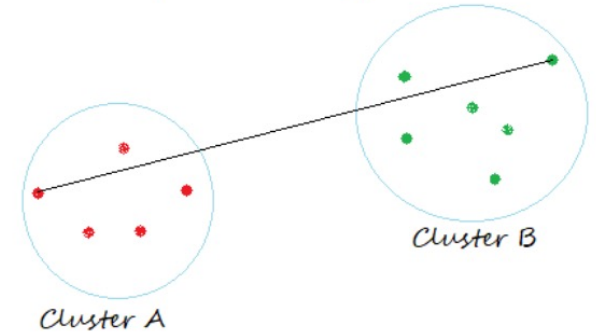


Linkage Functions

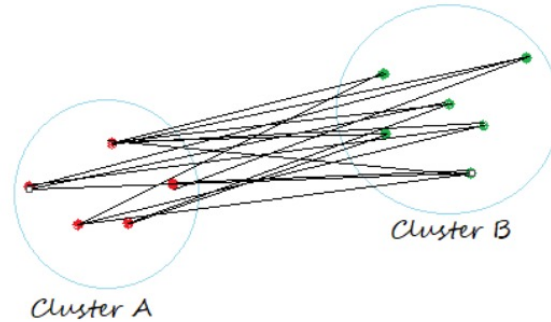
Single Linkage



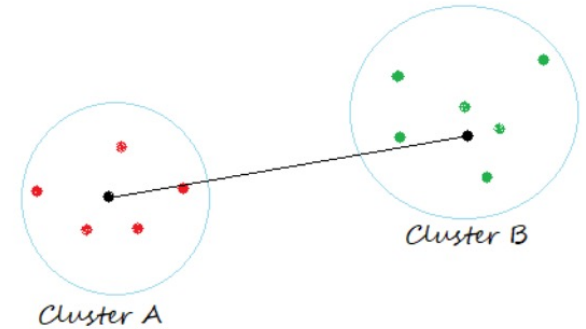
Complete Linkage



Average Linkage



Centroid Linkage



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

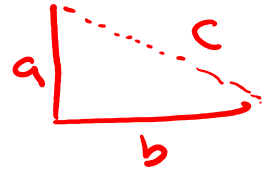
Computing all pairs of distances is pretty expensive!

A simple implementation takes $\mathcal{O}(n^2 \log(n))$

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 $\mathcal{O}(n^2)$



$$c \leq a + b$$



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

