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# Clustering methods

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# DataSets

**Features(Attribute/variable)**

**Data records (samples)**

ID	Address	# Bed	#Bath	...	School Score	Year Build	Crime Rate
				...			
				...			

**# Features = Dimension of dataset**

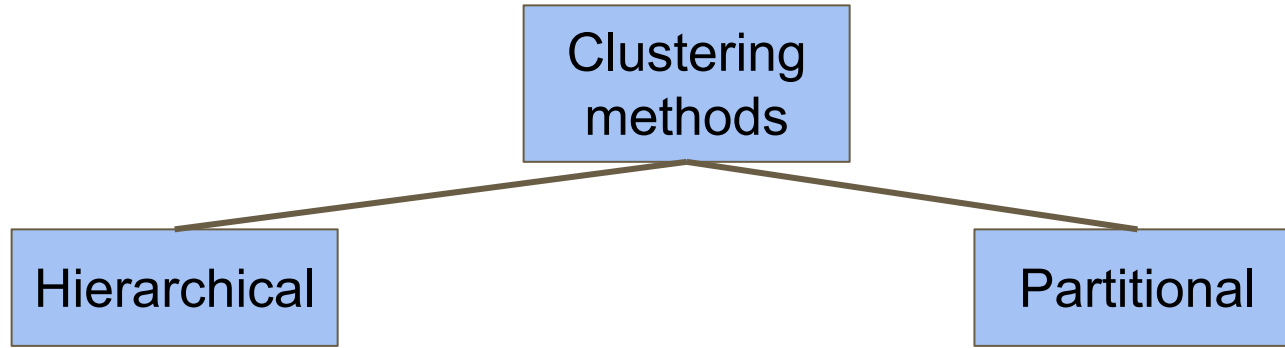
# Unsupervised versus supervised learning

- **Supervised learning**
  - Trying to predict label or value of data points
- **Unsupervised learning**
  - Unlabeled input data

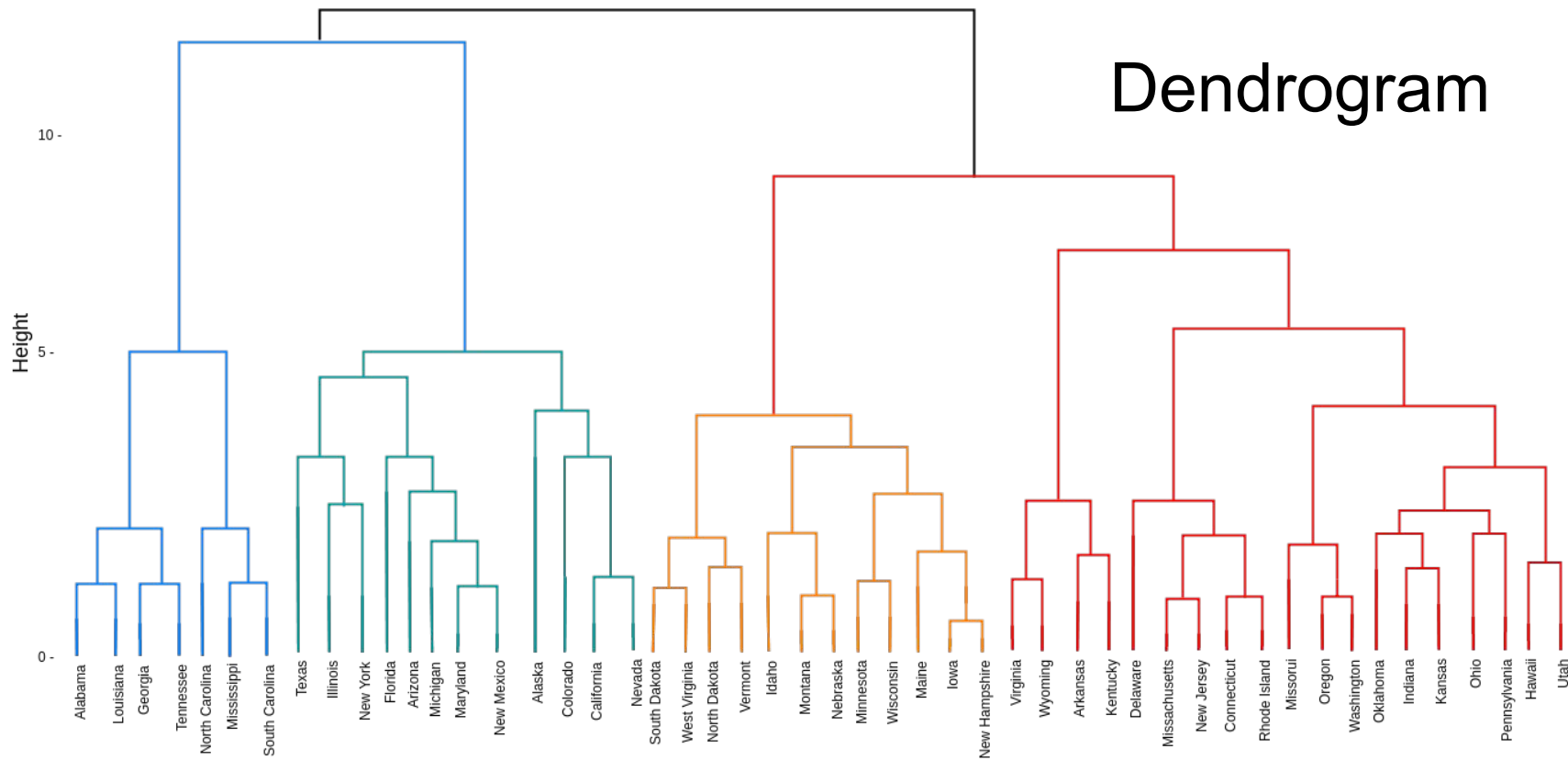
# Why do we use clustering?

- Group data points based on their similarities using the given features
  - Identify similar data points (within the same group)
  - Identify dissimilar data points (within different groups)
- Some methods also identifies outliers and points that cannot be assigned to any cluster
- Investigating differences between the groups
  - Survival of breast cancer patients

# Categories of clustering methods that we focus on



# Agglomerative hierarchical clustering



# Steps of agglomerative hierarchical clustering

1. Computing dissimilarity or similarity between every pair of data points
2. Using linkage function to group objects into hierarchical cluster tree
  - a. linkage function determines the distance between sets of data points as a function of the pairwise distances between data points in the groups.
3. Deciding where to cut the hierarchical tree into clusters.

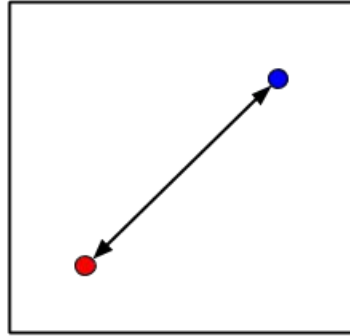
# Common distance measures used in clustering

- Euclidean distance
- Manhattan distance
- Minkowski distance
- Chebychev distance
- Cosine similarity
- Hamming distance
- Binary distance
  - Jaccard index
  - Hamming distance



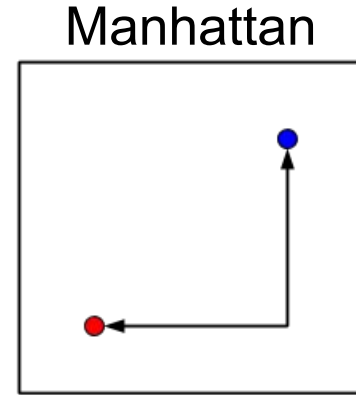
# Euclidean distance

Euclidean



$$d = \sqrt{(x_1 - x_2)^2 + (y_1 - y_2)^2}$$

# Manhattan distance

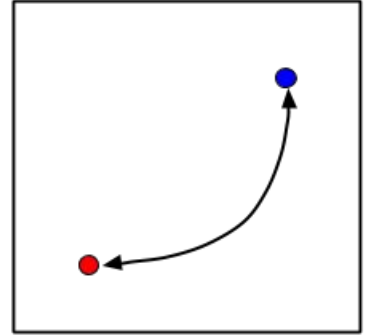


$$d = |x_1 - x_2| + |y_1 - y_2|$$

# Minkowski distance

$$d = (|x_1 - x_2|^p + |y_1 - y_2|^p)^{1/p}$$

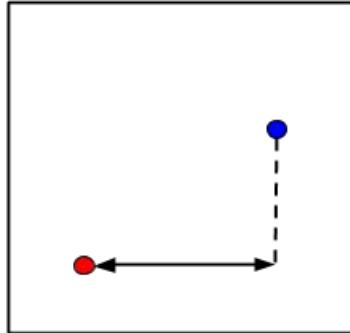
Minkowski



# Chebychev distance

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

Chebychev



# Jaccard index

$$J(A, B) = \frac{A \cap B}{A \cup B}$$

Jaccard



# Hamming distance

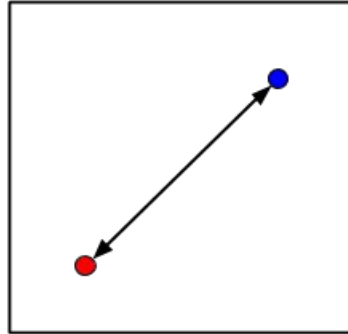
$d = \text{number of bits that they are different}$   
Hamming

"Go lang"

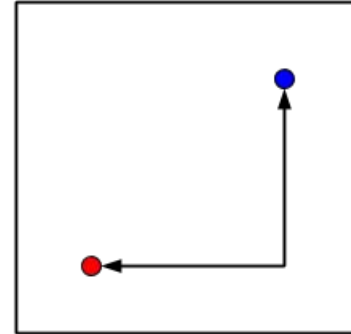
"Go pher"

# Different distance measures

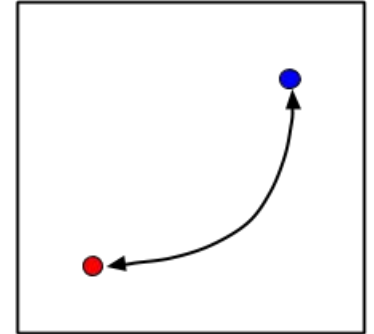
Euclidean



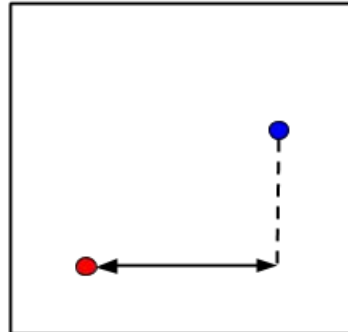
Manhattan



Minkowski



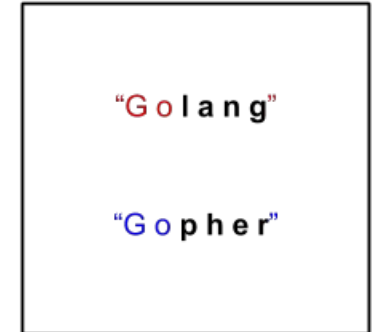
Chebychev



Jaccard



Hamming



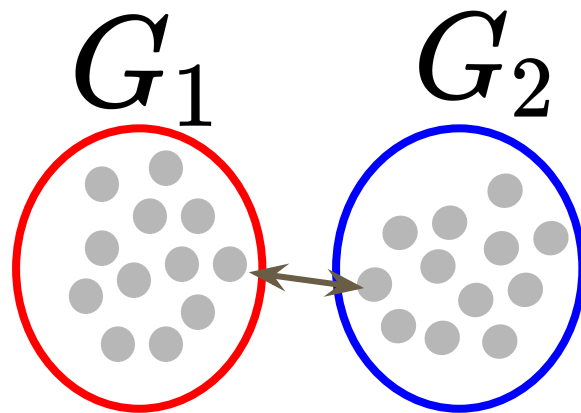
# Some of widely-used linkage functions

- Single
- Complete
- Average
- Median
- Centroid



## Some of widely-used linkage functions

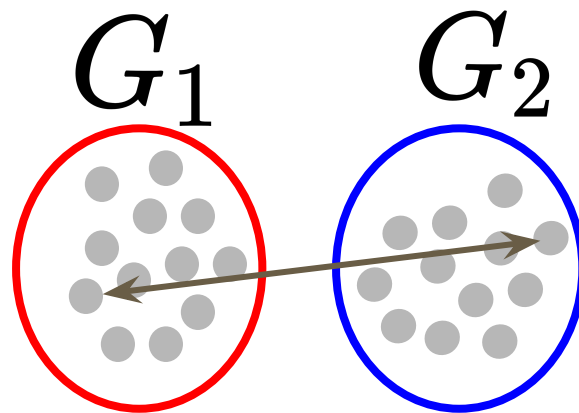
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$$D(G_1, G_2) = \min(d(x, y)), x \in G_1, y \in G_2$$

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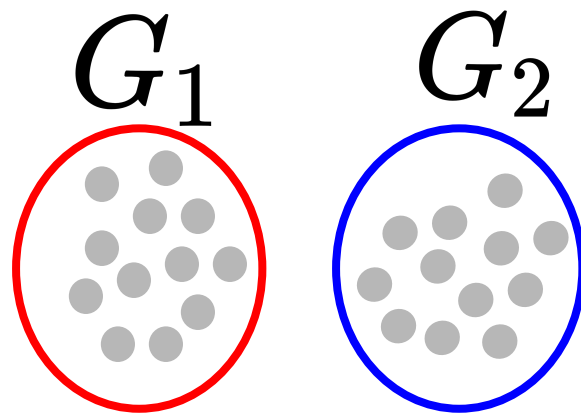
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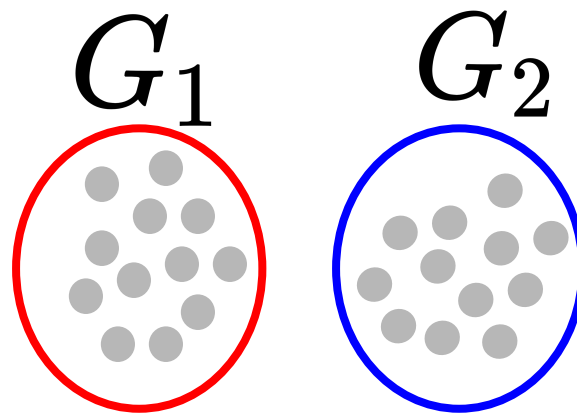


$$D(G_1, G_2) = \frac{1}{|N_{G_1}| |N_{G_2}|} \sum_x \sum_y d(x, y)$$

$$x \in G_1, y \in G_2$$

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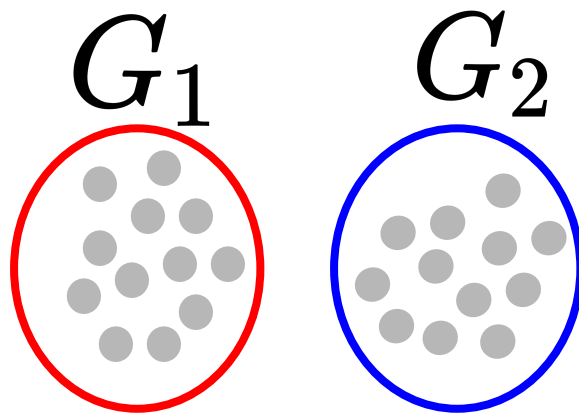
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Weighted center of mass distance (WPGMC)  
*Note.* appropriate for Euclidean distances only

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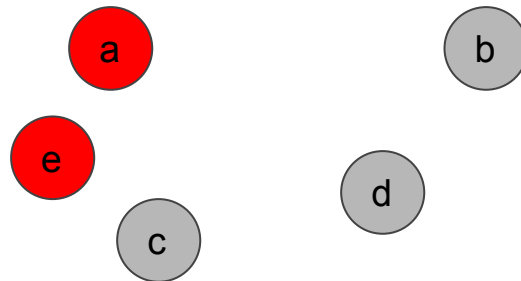
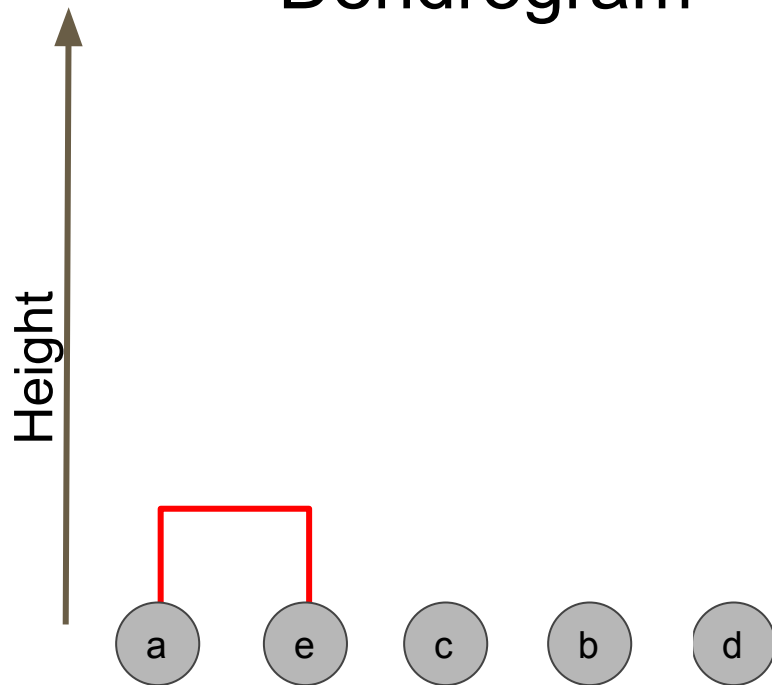


$$D(G_1, G_2) = ||c_{G_1} - c_{G_2}||$$

*Note.* appropriate for Euclidean distances only

# Let's see how agglomerative hierarchical clustering works

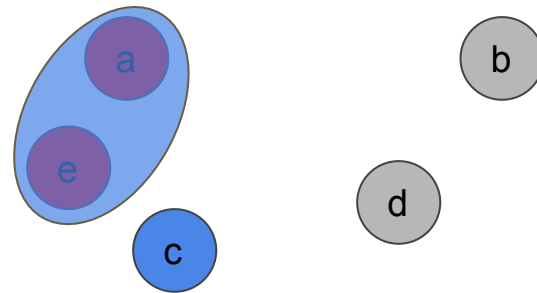
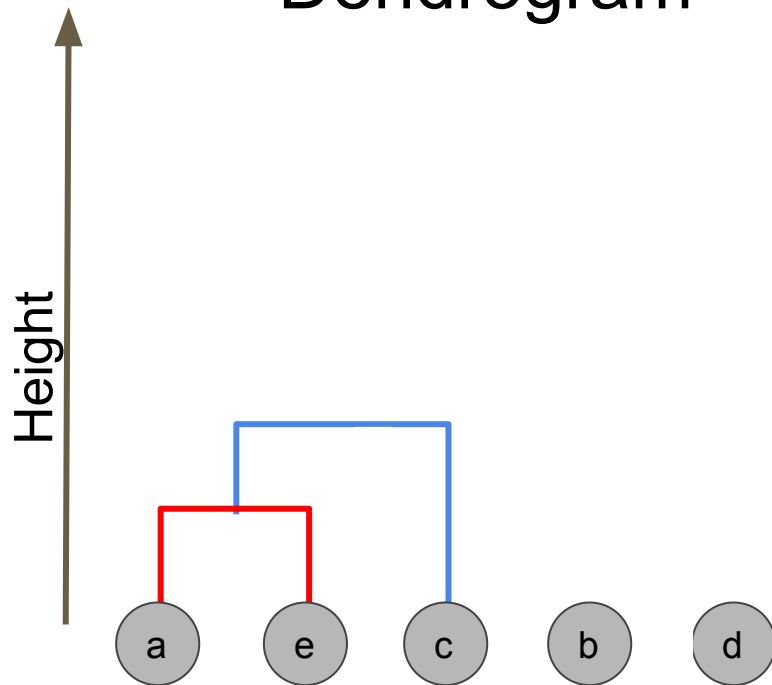
## Dendrogram



Data points should be rearranged for building the dendrogram

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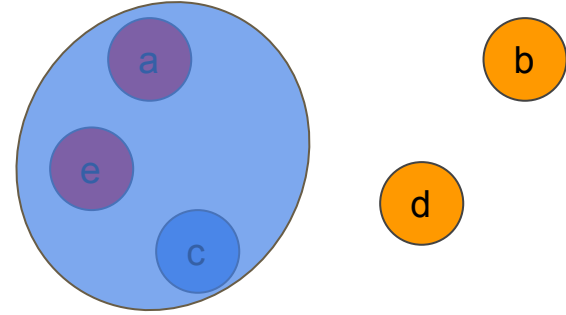
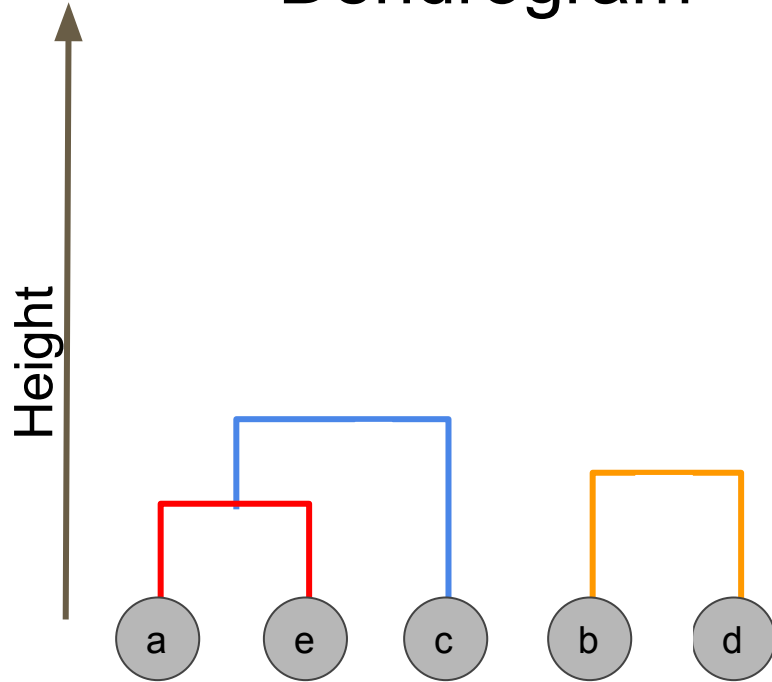
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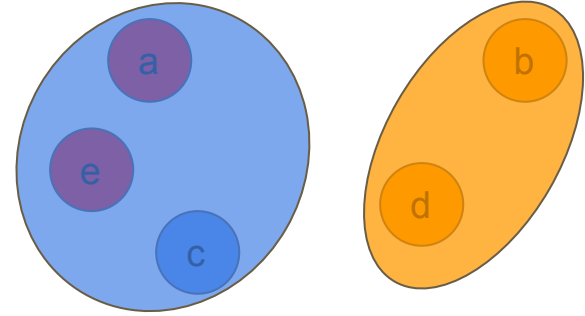
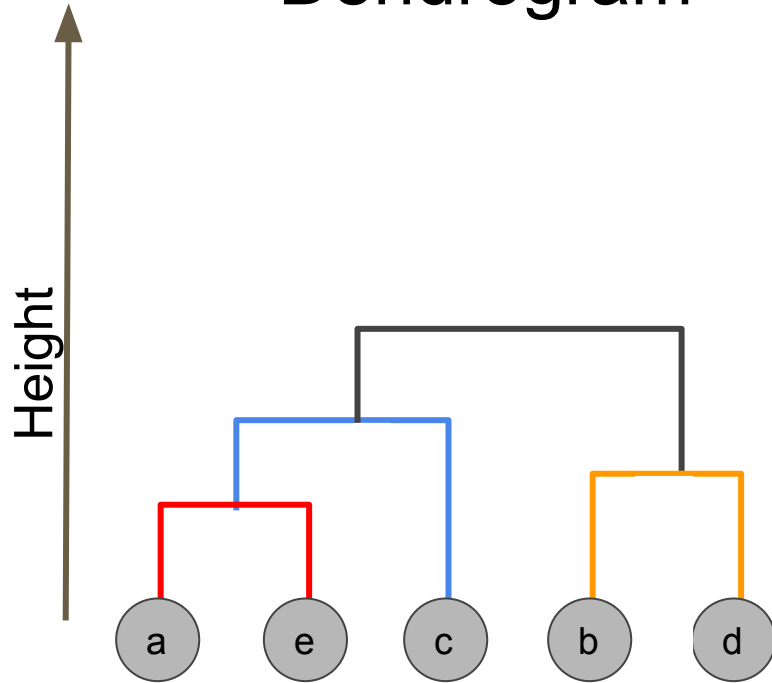
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# Let's see how agglomerative hierarchical clustering works

## Dendrogram



# Hierarchical versus partitional clustering

- Partitional clustering: division of the set of data points into clusters
  - each data object belongs to one subset
- Hierarchical clustering is a set of nested clusters
  - organized as a tree

# K-means clustering

Steps of k-means clustering

- 1) Choosing  $k$

# K-means clustering

## Steps of k-means clustering

- 1) Choosing  $k$
- 2) Randomly selecting  $k$  data points (as initial centers)
  - a) Final result depend on these points
    - i) Because k-means converges to one of many possible local minima

# K-means clustering

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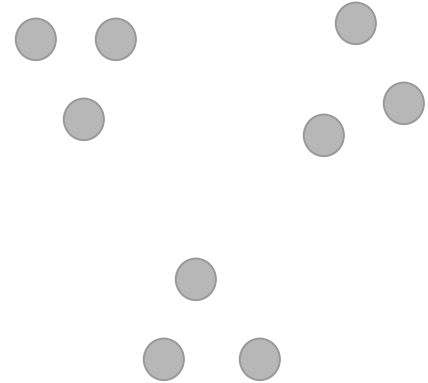
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- 5) Calculate new center of each cluster
- 6) Repeat steps (3) to (5) until convergence
  - a) No point in changing cluster membership



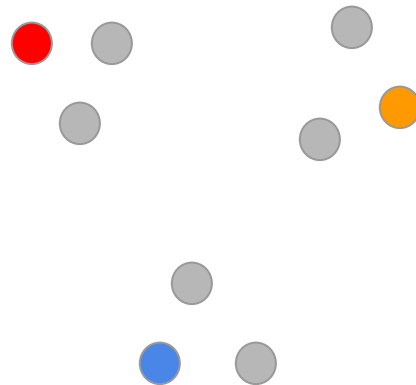
# Implementing k-means manually (simple example)

1) Choosing  $k$  (good guess  $k=3$ )



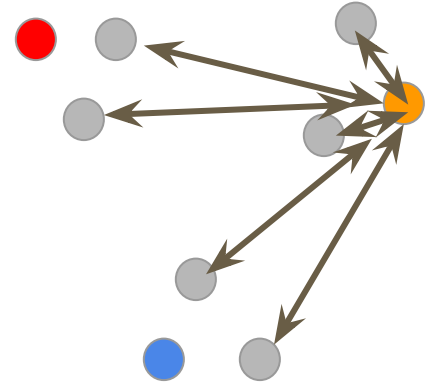
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- 1) Choosing  $k$  (good guess  $k=3$ )
- 2) Randomly selecting 3 data points (as initial centers)



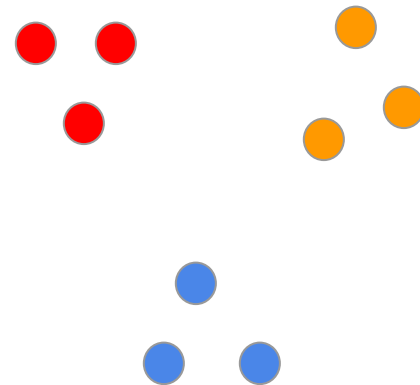
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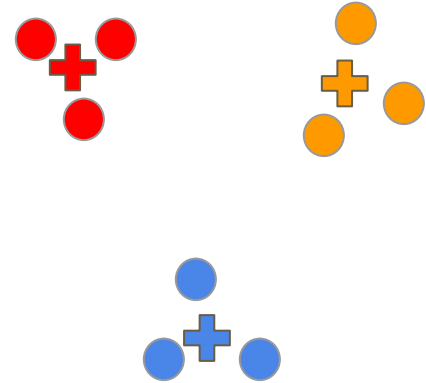
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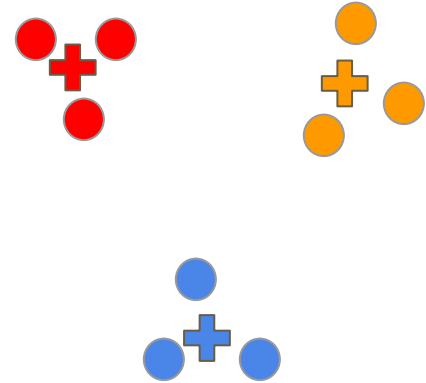
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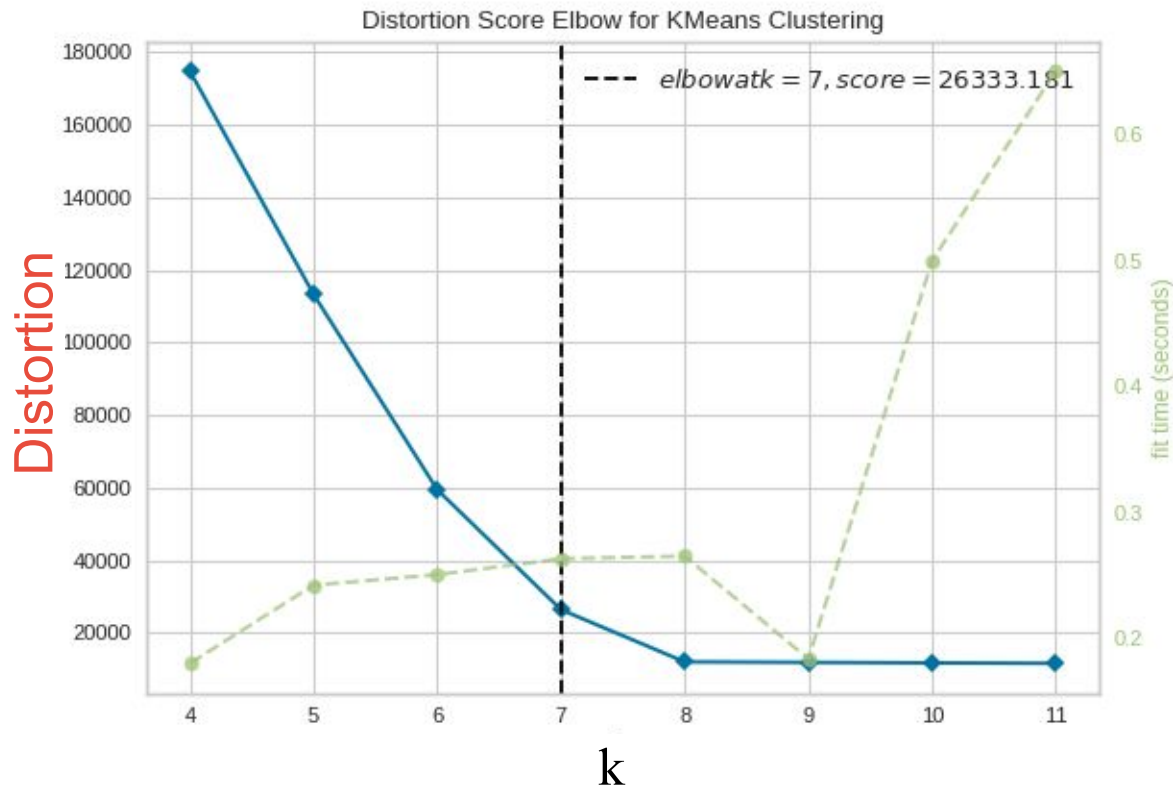
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- 5) Calculate new center of each cluster
- 6) No need for repetition
  - a) Good choice of initial centers
    - i) Fast convergence



# Elbow method for selecting optimal K

**Distortion** is the sum of squared distances from each point to its assigned center



# Hierarchical k-means (divisive hierarchical clustering)

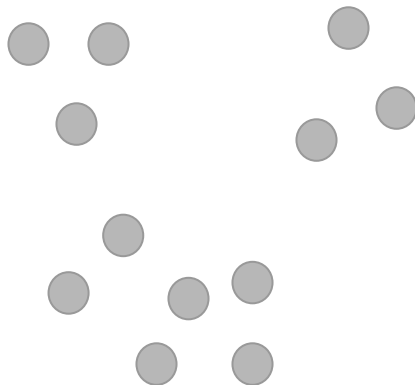
- Start with all the data points
- Implement k-means ( $k=2$ ) in an iterative manner
- Until reaching singletons (data points)



# Hierarchical k-means (divisive hierarchical clustering)

Issue with this process:

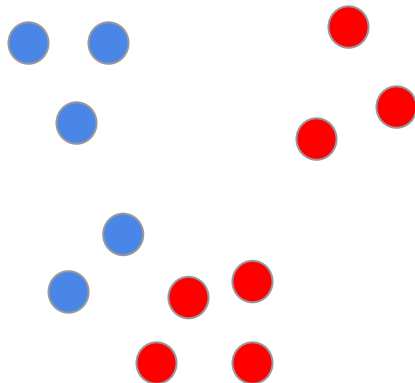
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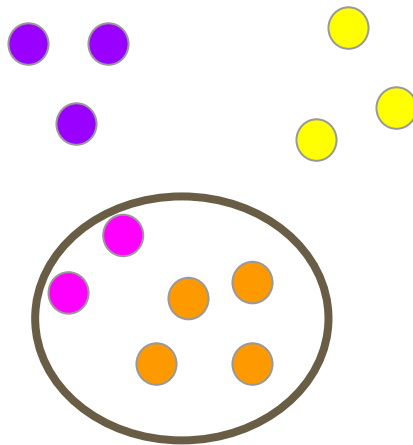


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We like these to  
be one cluster



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- Iterative message passing
  - Data points compete to become exemplars
    - Exemplars are real data points not centers as in k-means
- Initialization independent



# Message matrices

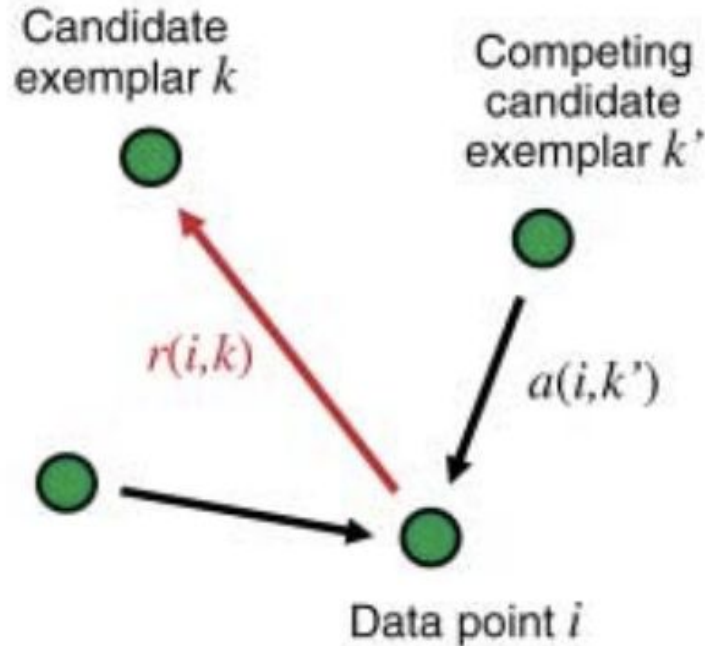
- **Responsibility matrix:**  $R(i,k)$ 
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# Message matrices

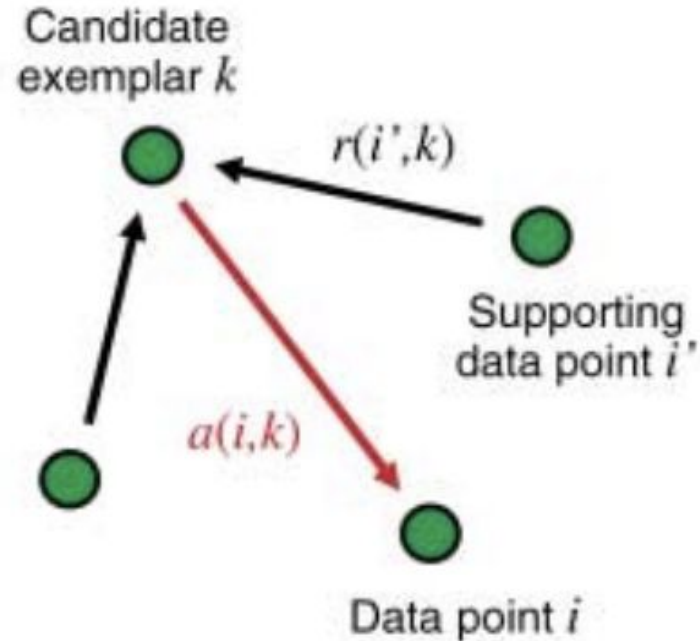
- **Responsibility matrix:**  $R(i,k)$ 
  - Responsibility message from  $i$  to  $k$ 
    - Accumulated evidence for how well-suited  $k$  is to serve as the exemplar for  $i$
- **Availability matrix:**  $A(i,k)$ 
  - Availability message from  $k$  to  $i$ 
    - Accumulated evidence for how appropriate it is for  $i$  to choose  $k$

# Sending responsibility and availability

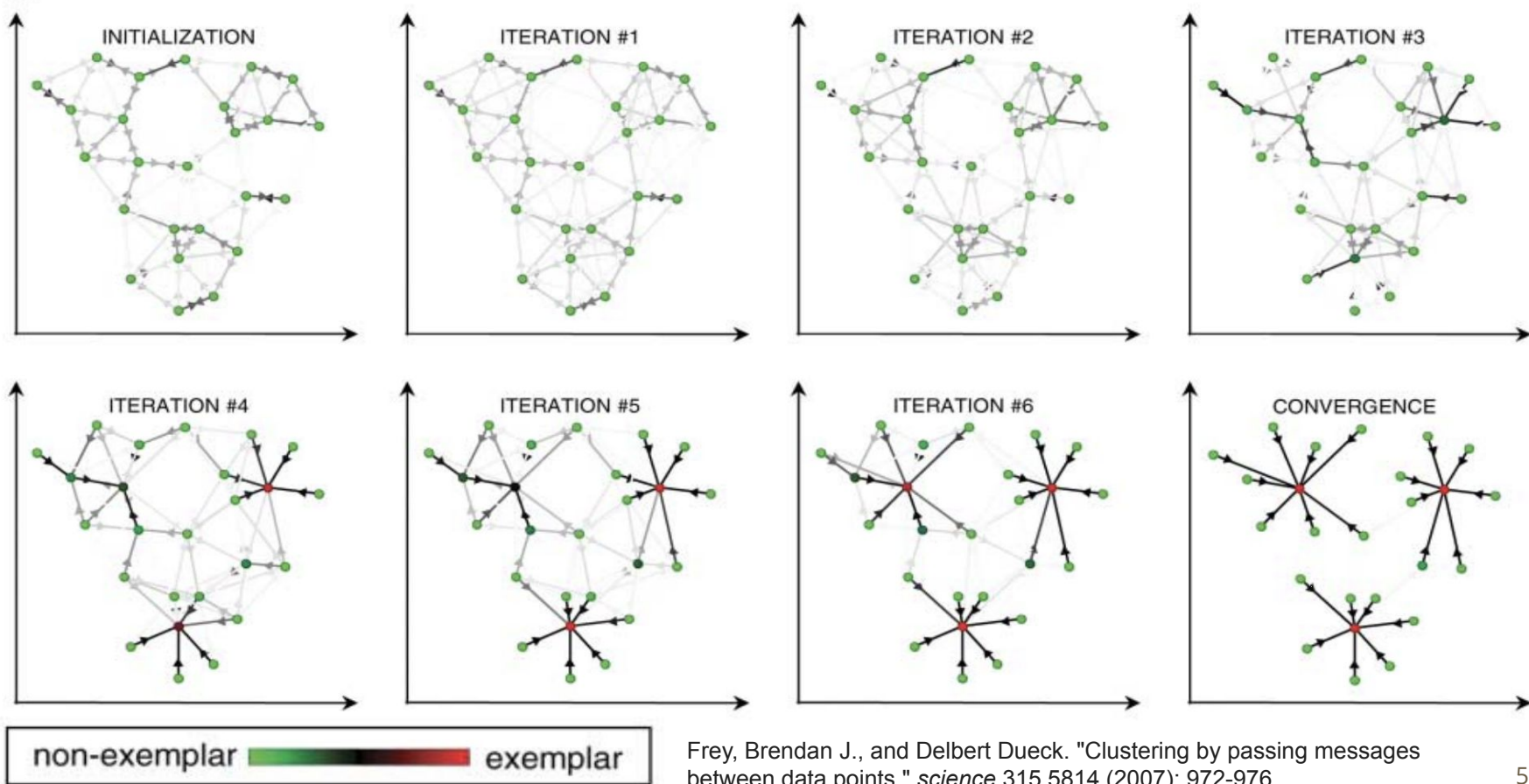
## Sending responsibilities



## Sending availabilities



# Iterations of affinity propagation for 2 dimensional data points



Frey, Brendan J., and Delbert Dueck. "Clustering by passing messages between data points." *science* 315.5814 (2007): 972-976.

# DBSCAN (Density-Based Spatial Clustering of Applications with Noise)

Let's look at clusters from another point of view:

- Clusters is a maximal set of density-connected points

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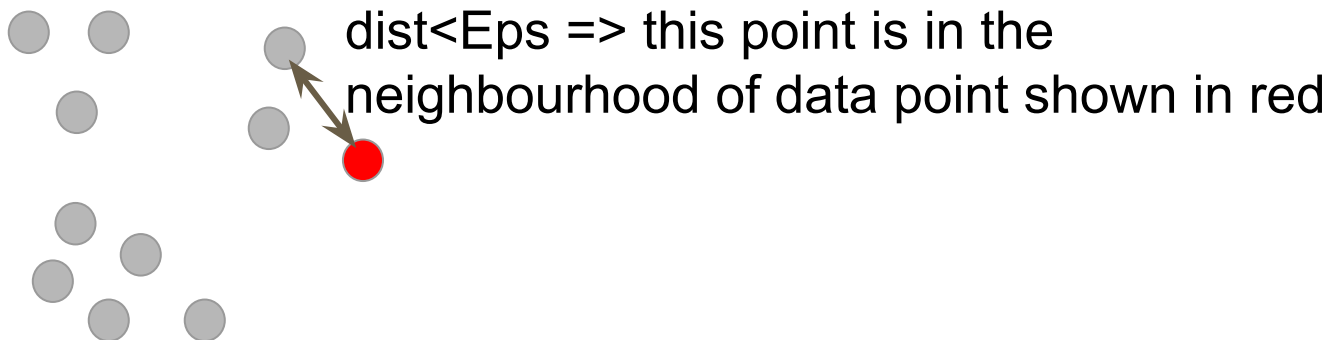
Two main parameters:

- **Epsilon (Eps)**: Maximum radius of neighbourhood
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Two main parameters:

- **Epsilon (Eps):** Maximum radius of neighbourhood
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  - Then that data point will be called a **core** point

Different types of data points:

- **Core**
- **Border:** within epsilon distance does not meet MinPts criteria
  - But there is at least 1 core point within the epsilon distance
- **Noise (Outlier):** Not assigned to any clusters

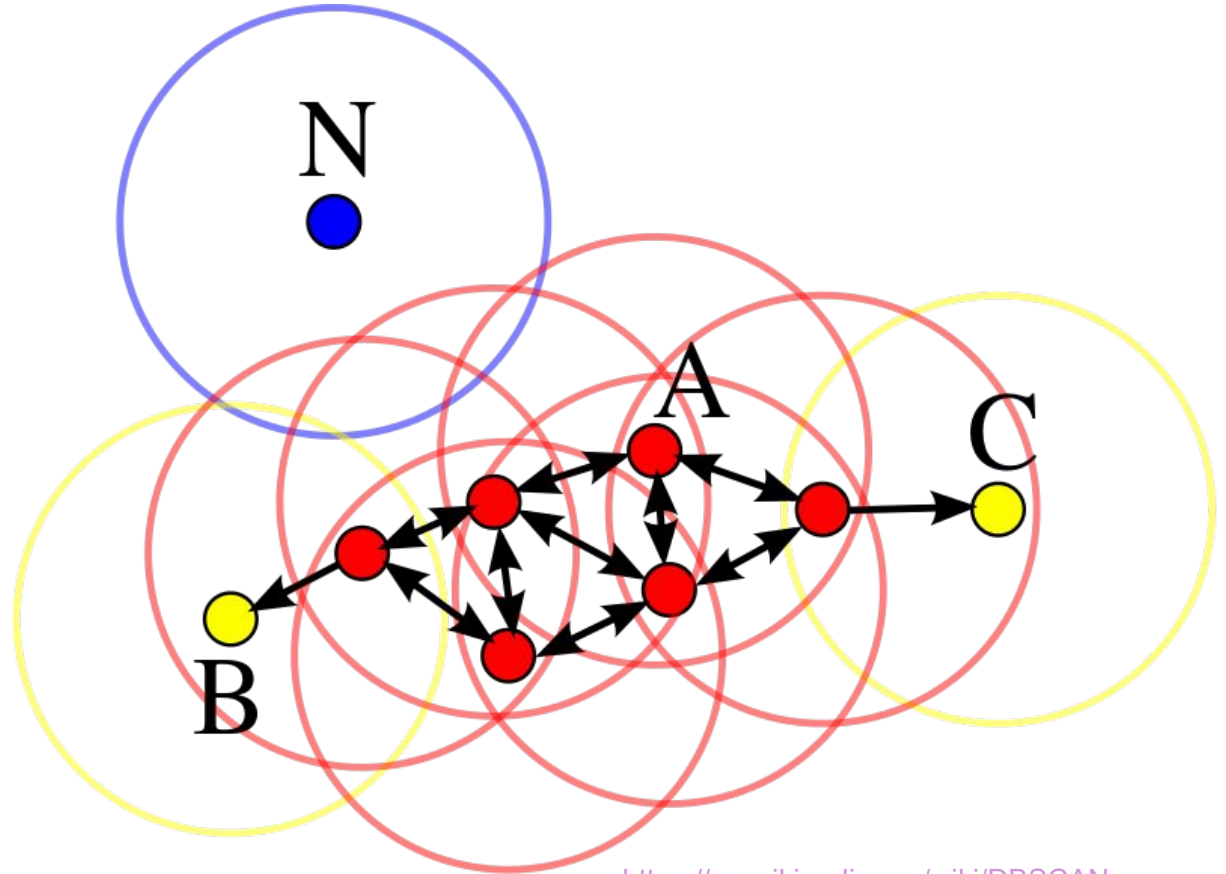


# Visualizing different types of data points in DBSCAN

**A:** Core

**B** and **C:** Border

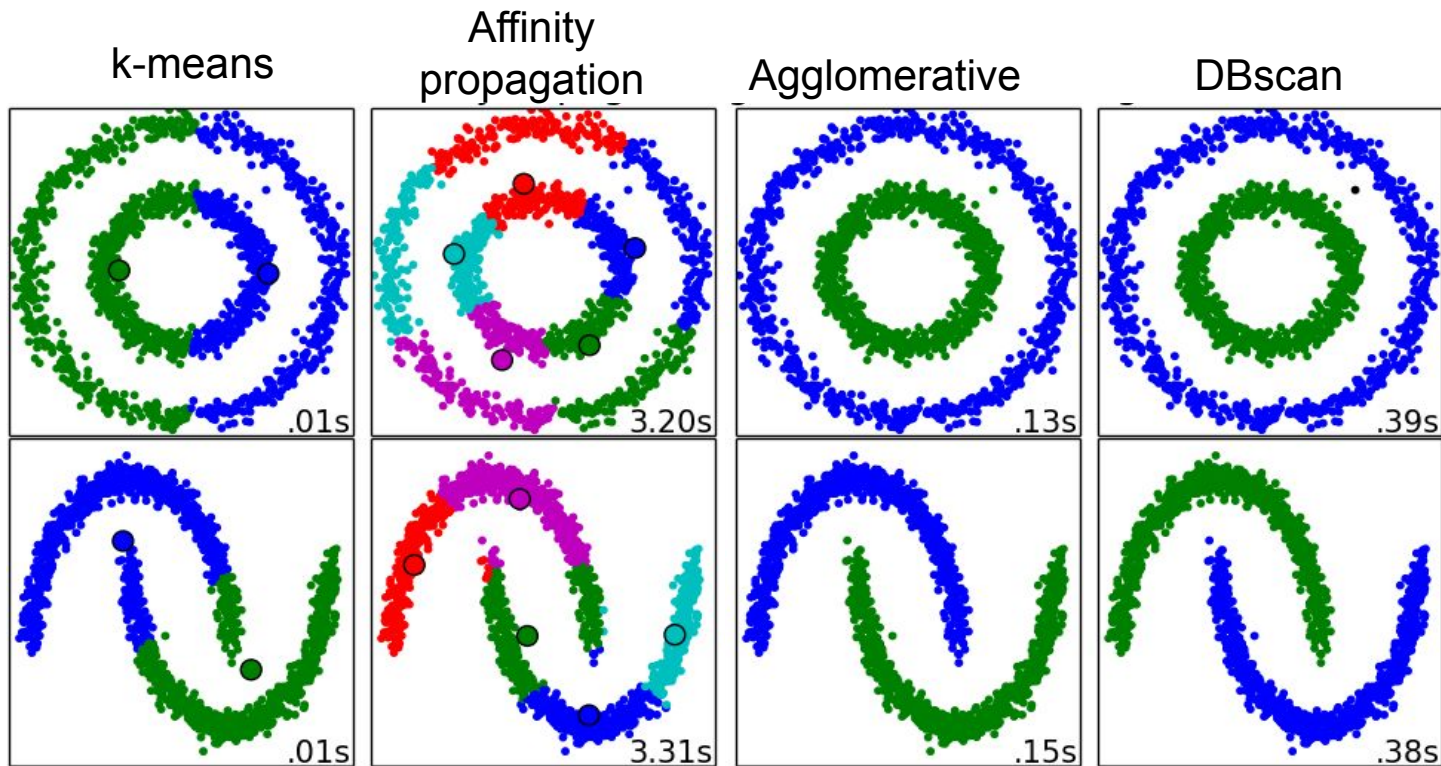
**N:** outlier



## Some disadvantages of DBSCAN

- May not work well with clusters of similar densities
  - But great for separating clusters of low and high densities
- May not be great with very high dimensional data

# Comparison of clustering methods



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