# Kmeans talk – Unsupervised learning- you do not know the number of clusters or cluster membership

#### First: Kmeans is not KNN

K-means is a clustering algorithm that tries to partition a set of points into K sets (clusters) such that the points in each cluster tend to near each other. It is unsupervised because the points have no external classification.

K-nearest neighbors is a classification (or regression) algorithm that in order to determine the classification of a point, combines the classification of the K nearest points. It is supervised because you are trying to classify a point based on the known classification of other points.

### **Problems**

K-means is vulnerable to outliers

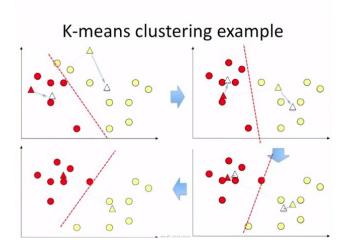
K-means works best with spherical clusters

K-means requires you to pick the number of clusters you have before you run the algorithm (how to know)

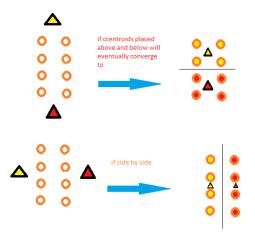
K-means is not guranteed to produce the same result every run (it depends on initial cluster centers)

# **Algorithm**

- 1. Randomly pick \*k\* centroids from sample points as initial cluster centers
- 2. Assign each sample to the nearest centroid
- 3. Move the centroids to the center of the samples assigned to it
- 4. Repeat steps 2 and 3 until cluster membership stops changing



# It is not guaranteed to converge to same solution every time



## sklearns implementation

use kmeans++ to initialize (picks the k points in the data furthest from each other)

talk about inertia (Sum of squared distances of samples to their closest cluster center)

### Pick ideal number clusters, the elbow method

calculate the inertia for every value of K, plot these inertias, look for point where K starts to increase rapidly (the elbow)

## Pick ideal number clusters, the Silhouette score

Plots how tightly grouped the samples in the clusters are. How to calculate the <u>silhouette</u> <u>coefficient of a single sample</u> in a dataset. Do this for all samples then plot per cluster.

- 1. ai cluster cohesion- average distance between sample and all other points in the cluster
- 2. bi seperation- average distance between sample and all other points in nearest cluster

Silhouette score for point= (bi-ai)/max(bi,ai)

- -1<Si<1
- =0 if ai=bi then point is right on edge of being mis classified
- =1 ideal bi>>ai very far away from nearest clustering
- <0 then point probably misclassified</p>

Do this for every point, then plot the scores per point and per cluster