

Clustering

Standard problem:

Regression and classification:

We want to find the function that satisfies the training data. In Case of clustering the task is clustering.

Task: It is to group the similar data points.

"Clustering:"

classification & regression

✓ $\mathcal{D} = \{x_i, y_i\}$ $y = f(x)$

✓ $y_i \in \{0, 1\} \rightarrow$ 2-class classification

✓ $y_i \in \mathbb{R} \rightarrow$ regression

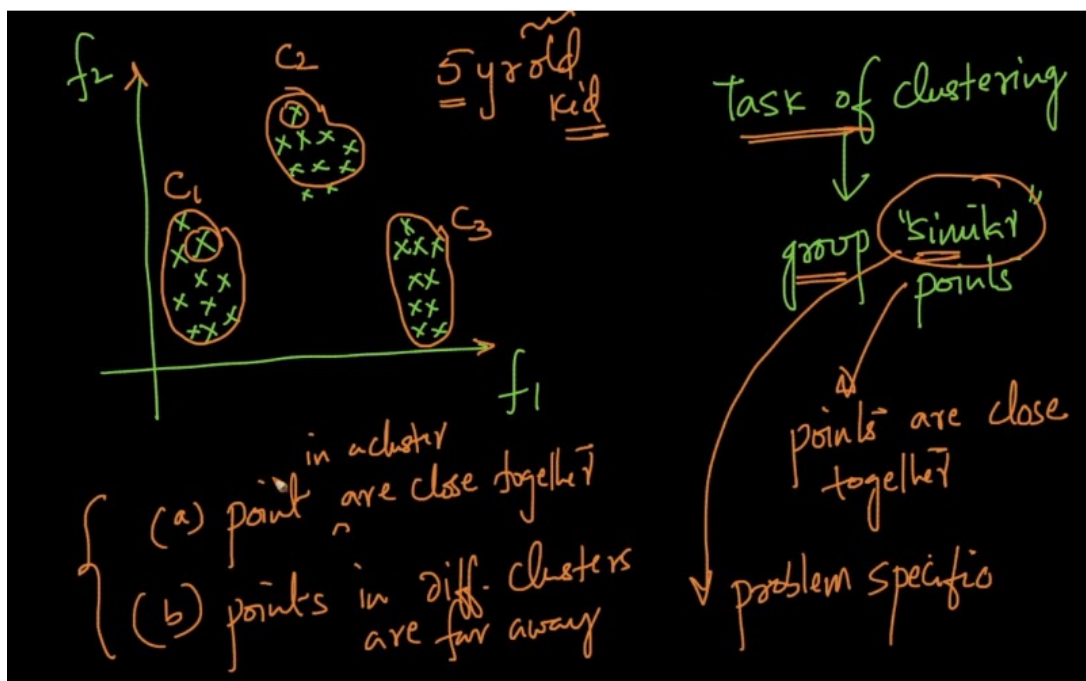
Clustering

$\mathcal{D} = \{x_i\}$
no y_i 's

Task: group/cluster
"similar"
data points

Consider the data points as follows:

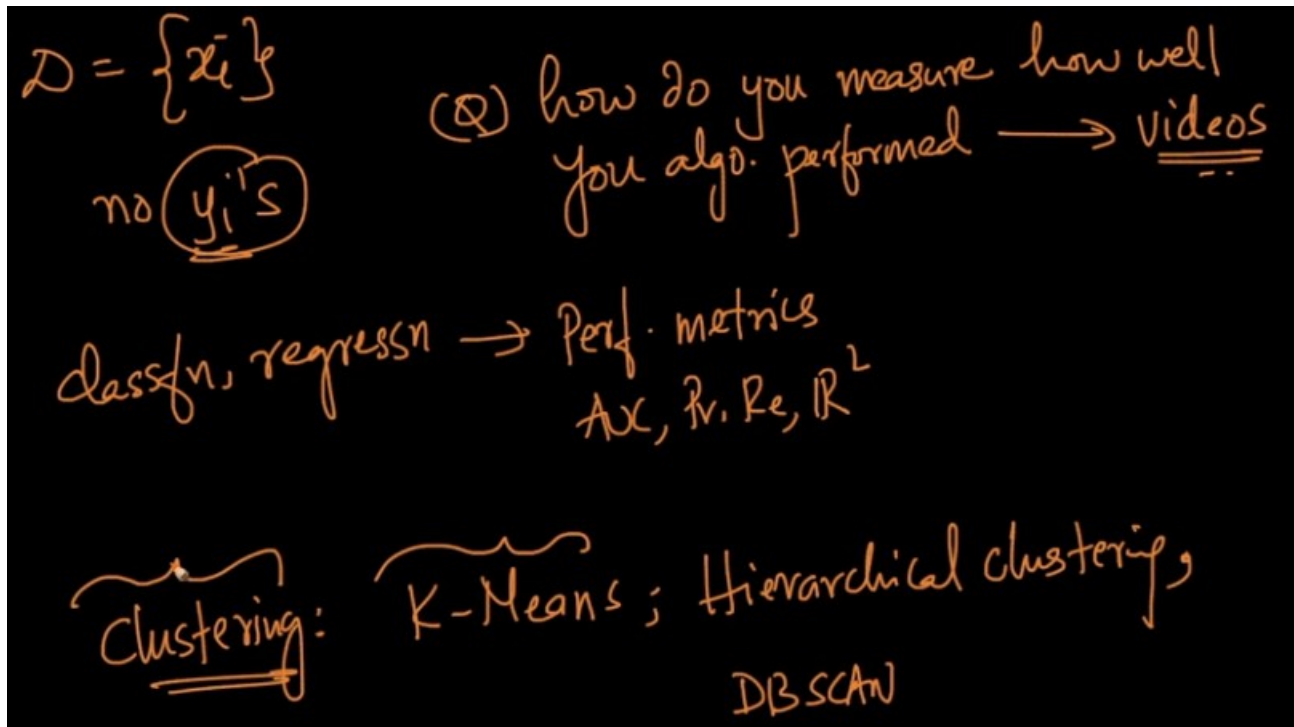
The task of clustering is to group the similar data points. All these points are grouped together. The points are grouped together. The similar is very much task specific.



Here all the data given are the data points in the mathematical form.

Measuring the performance of the clustering algorithm. There are various metrics that can measure the performance of clustering.

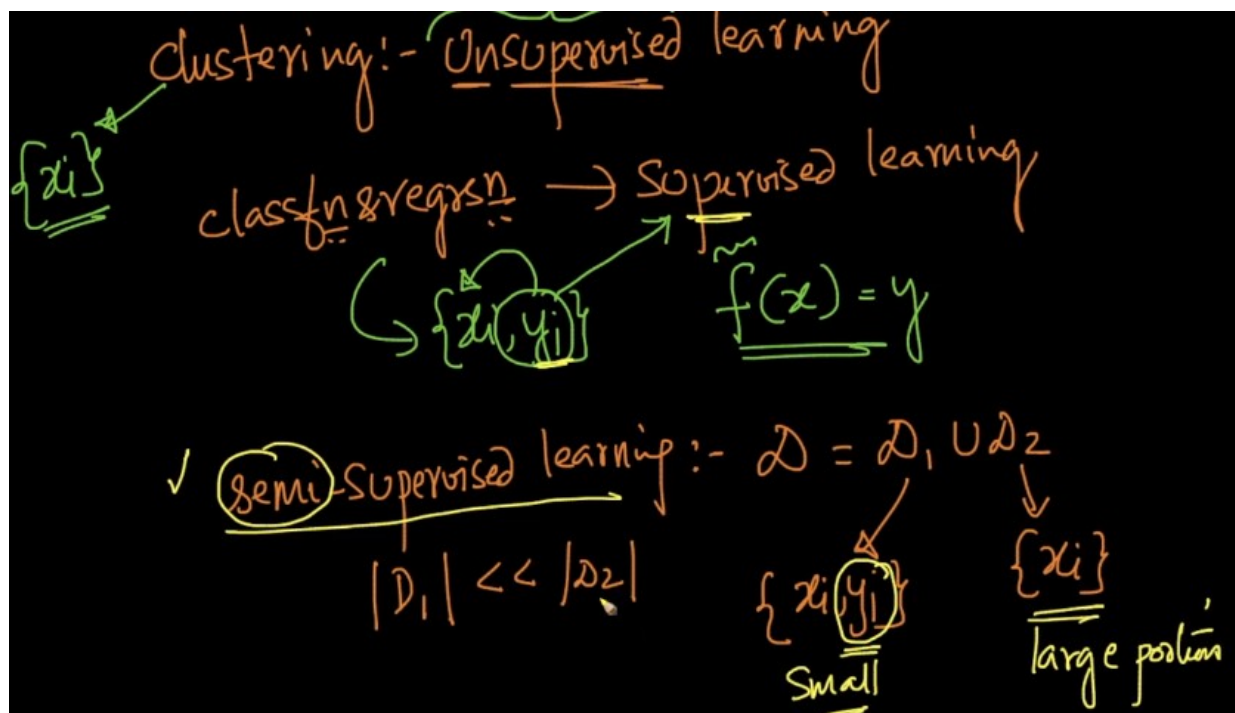
Clustering algos: K- Means, Hierarchical, DBSCAN.



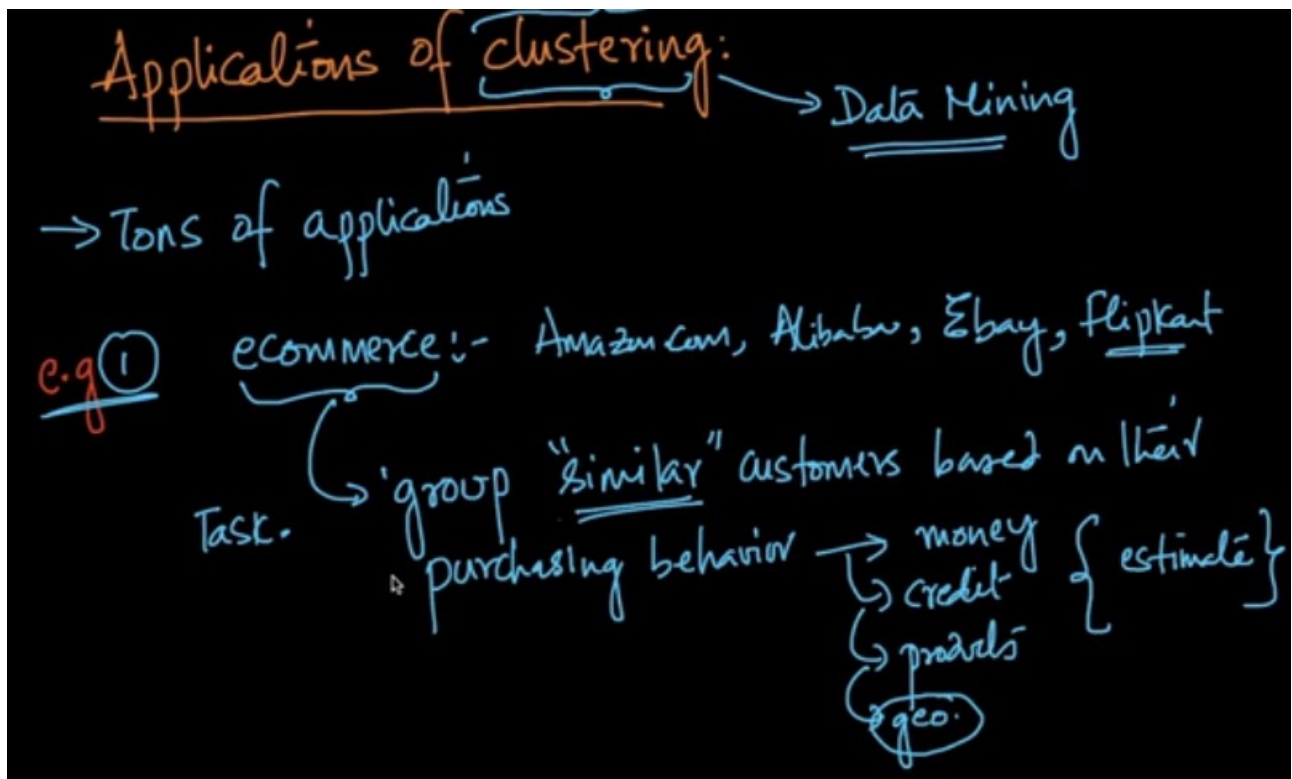
These are the most used algos. There are algos for various data.

Unsupervised learning: Clustering is referred to as unsupervised learning. Both classification and regression are the supervised learning.

There is also an area called semi-supervised learning, the data sets are union of D_1 and D_2 . This happens when the cost of labeling the data is expensive.



Applications of clustering: These are more studied in Data mining, than machine learning.
There are many applications of clustering.
They want to group the similar customers based on the purchasing behavior.
They can decide the income level using the purchasing behavior.



Assume the clusters are grouped to various classes. By grouping customers we can give different offers.

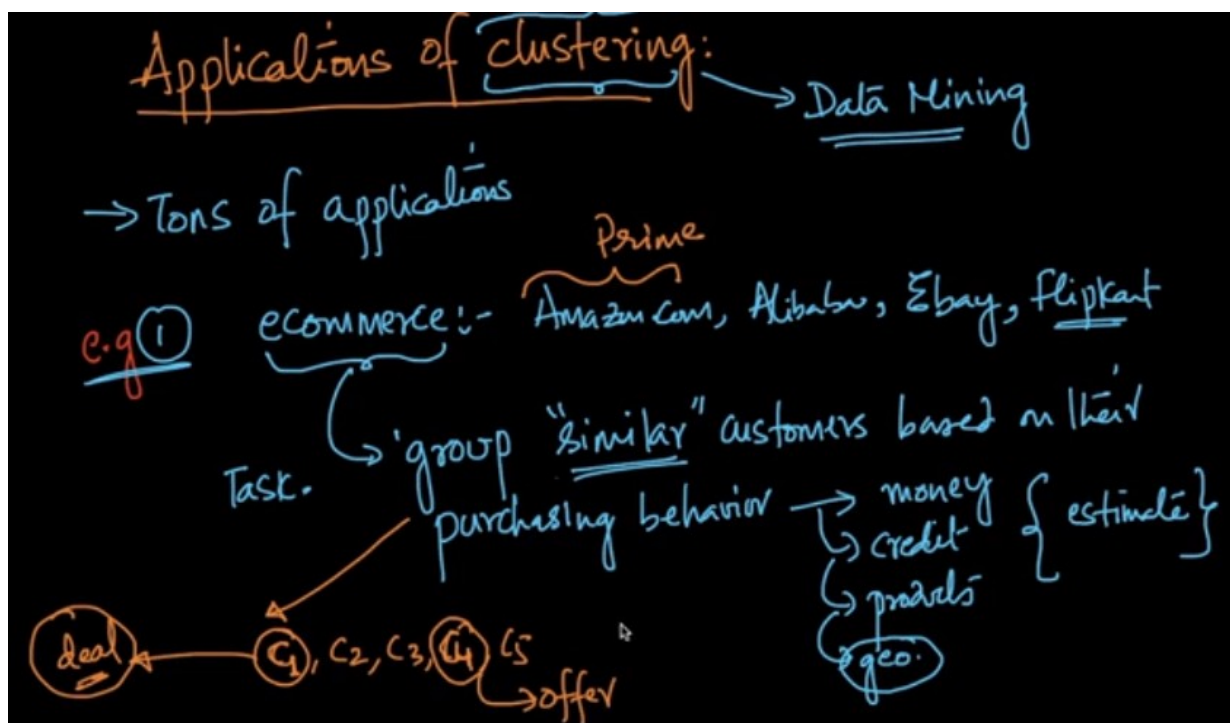
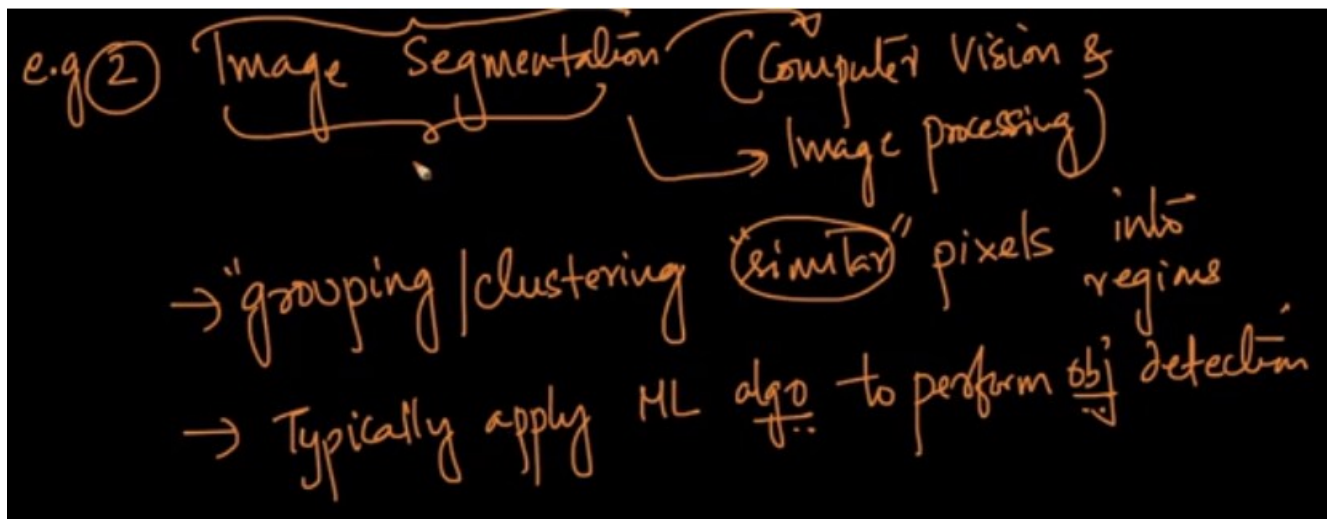
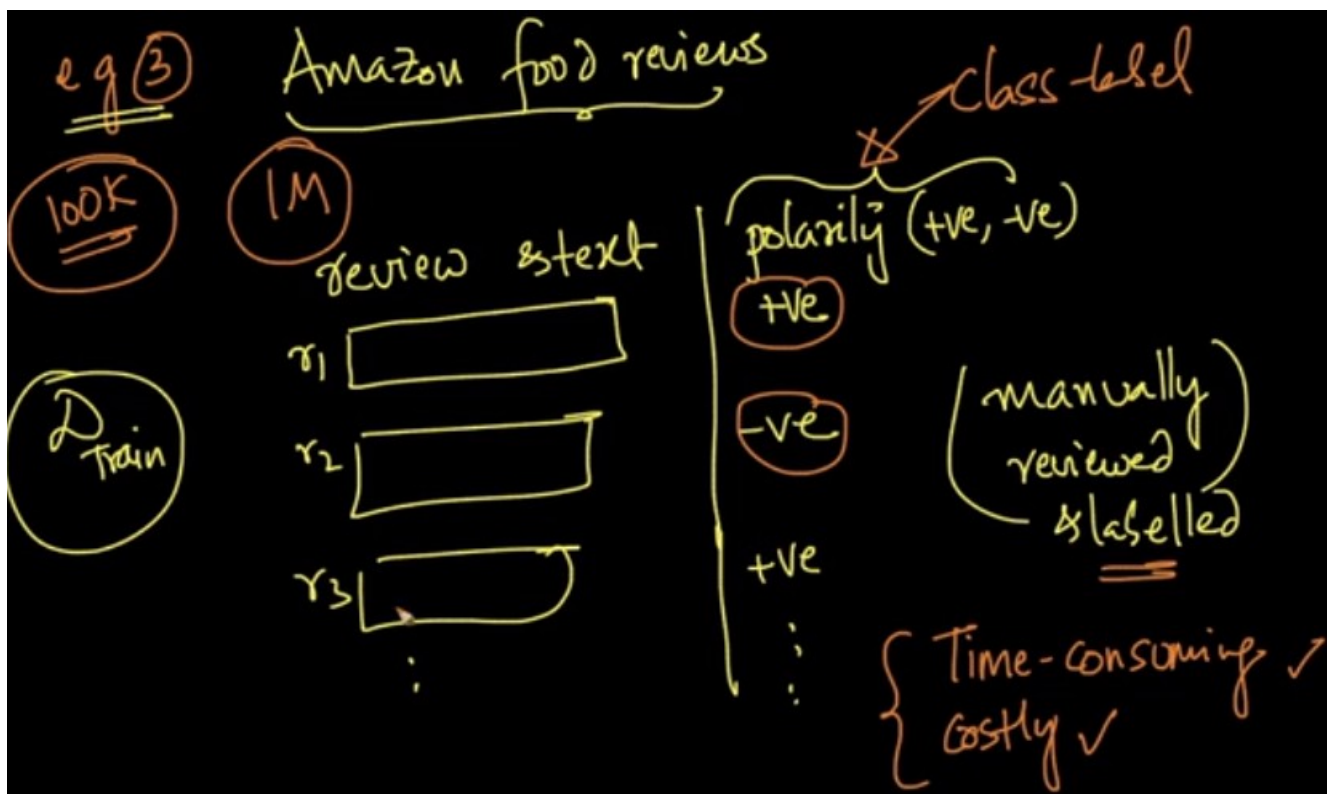


Image segmentation: This is a problem in computer vision and image processing.

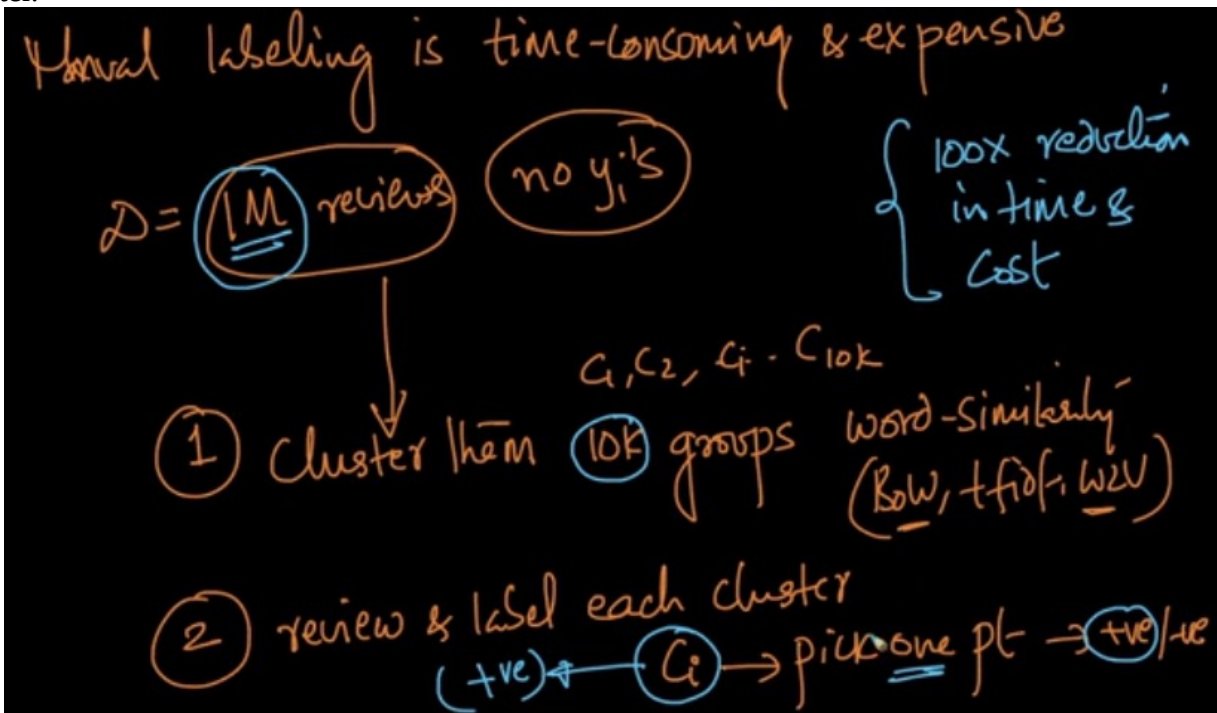
This is all about segmenting the pixels of the image. The clustering can help in seg the image. We can apply the ML algo to apply the object detection.



The polarity is the class label. This is very time consuming. We can apply the clustering algorithm to get the labels of the reviews.



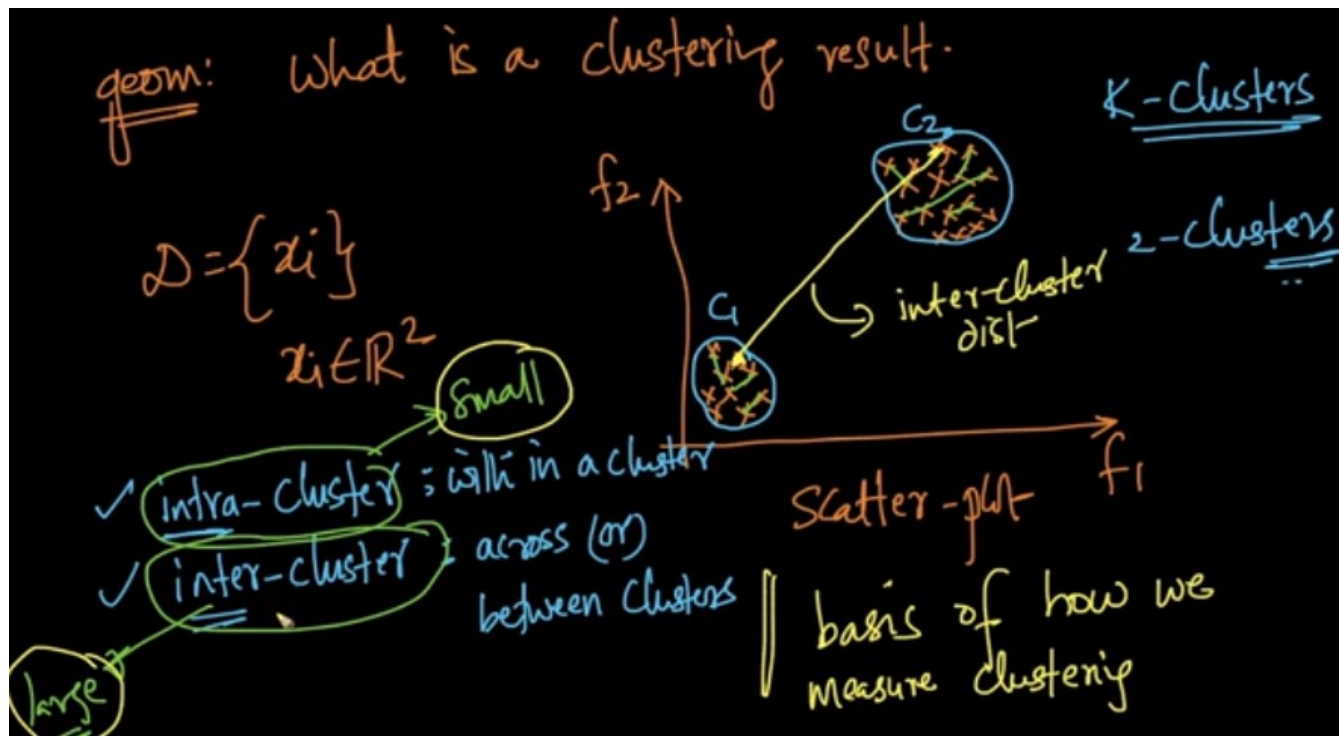
The manual labeling is time consuming and expensive. We can pick a cluster it is review and label each cluster.



Now, we can train the machine learning algorithms.

Metrics for measuring Clustering: What is a good clustering result?

All the points inside the cluster is called the intra clusters. All the outside the cluster is called the inter clusters. The intra cluster distance is kept small and the inter cluster distance kept large. This leads to the good clustering. **This is the basis of clustering effectiveness.**



In an ideal world, we want the inter clustering distance to be very high and intra to be lvery less.

Dunn – index: The numerator is the max inter cluster distance. The denominator is the intra cluster distance.

ideal:- inter-cluster dist \rightarrow v. high
intra-cluster dist \rightarrow v. low

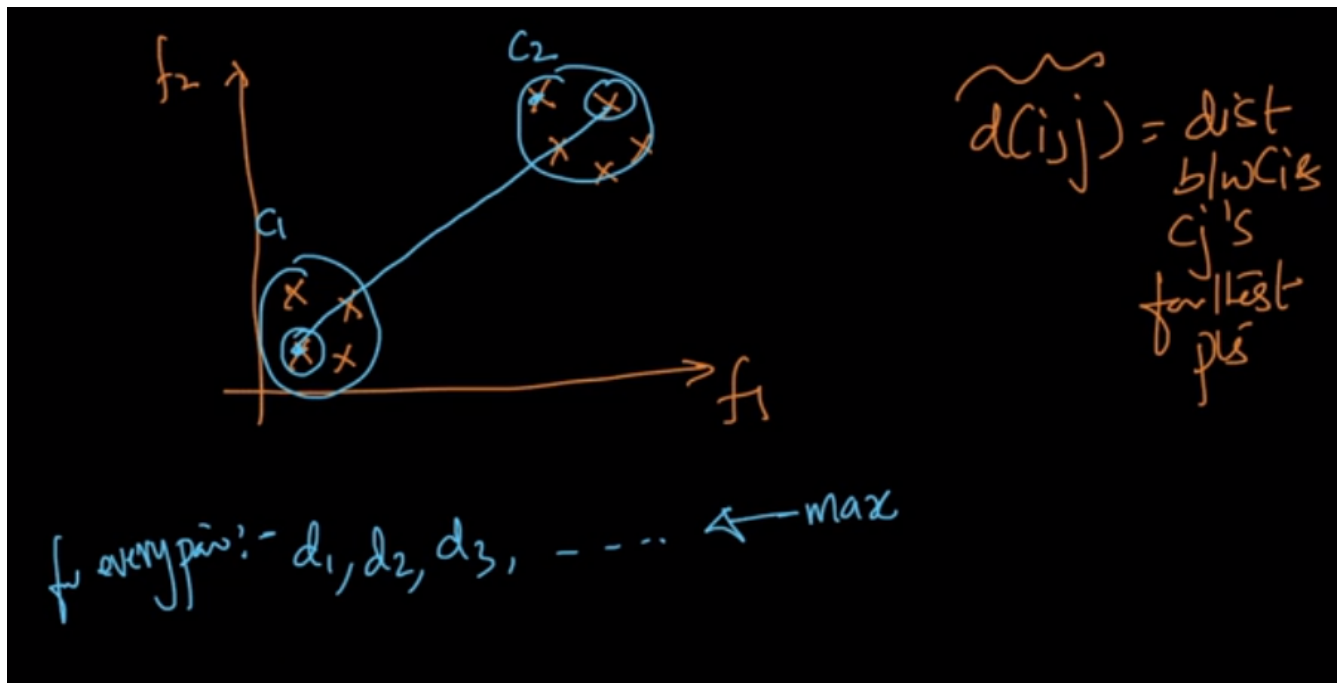
Dunn-index: $D = \frac{\max_{i,j} d(i,j)}{\max_K d''(K)}$

$d(i,j)$ is dist c_i & c_j K -clusters $\{c_1, c_2, \dots, c_i, \dots, c_K\}$
maximal inter-clust dist

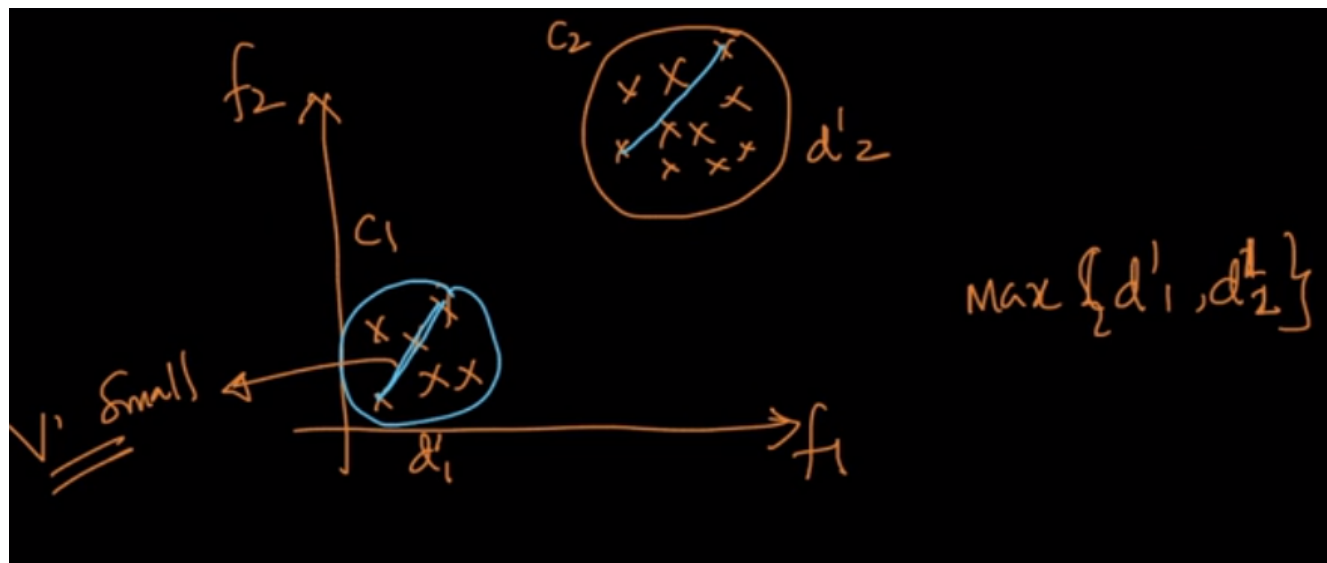
$d''(K)$ is intra-cluster dist (K)
max K

D is high \Rightarrow good clustering

The distances are calculated as follows and the farthest points are chosen.



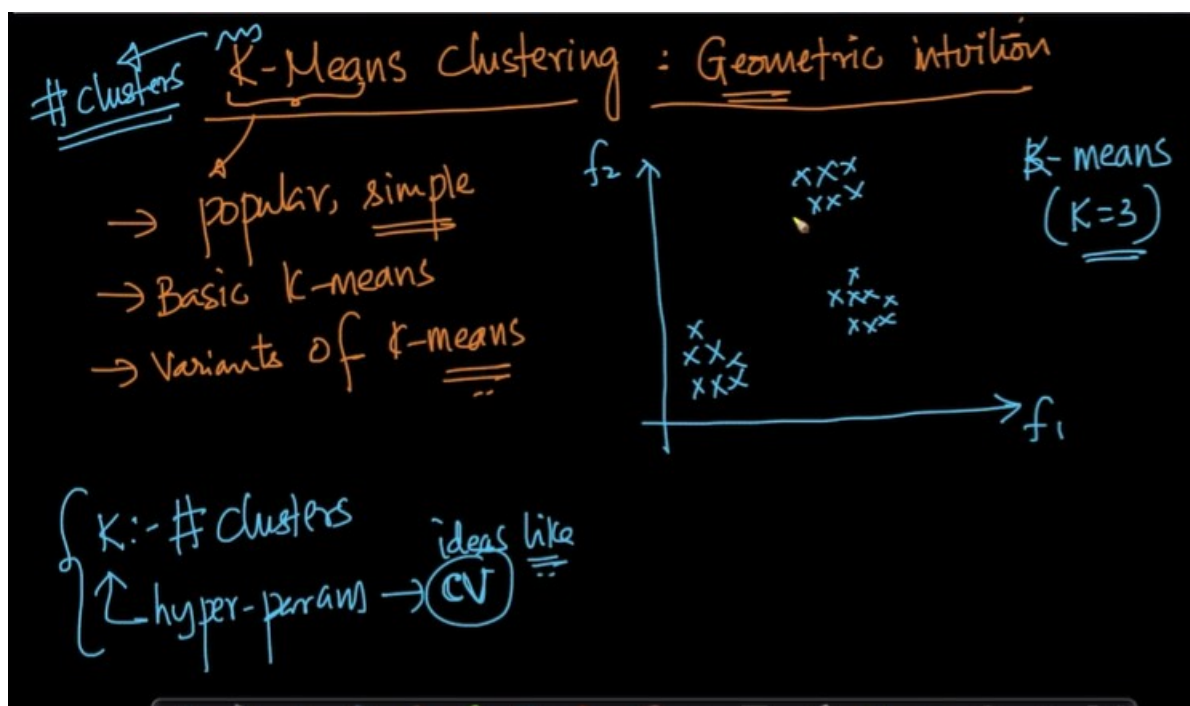
Denominator: The farthest distance of the cluster is computed, then we take the max over the distances d'_1 and d'_2 .



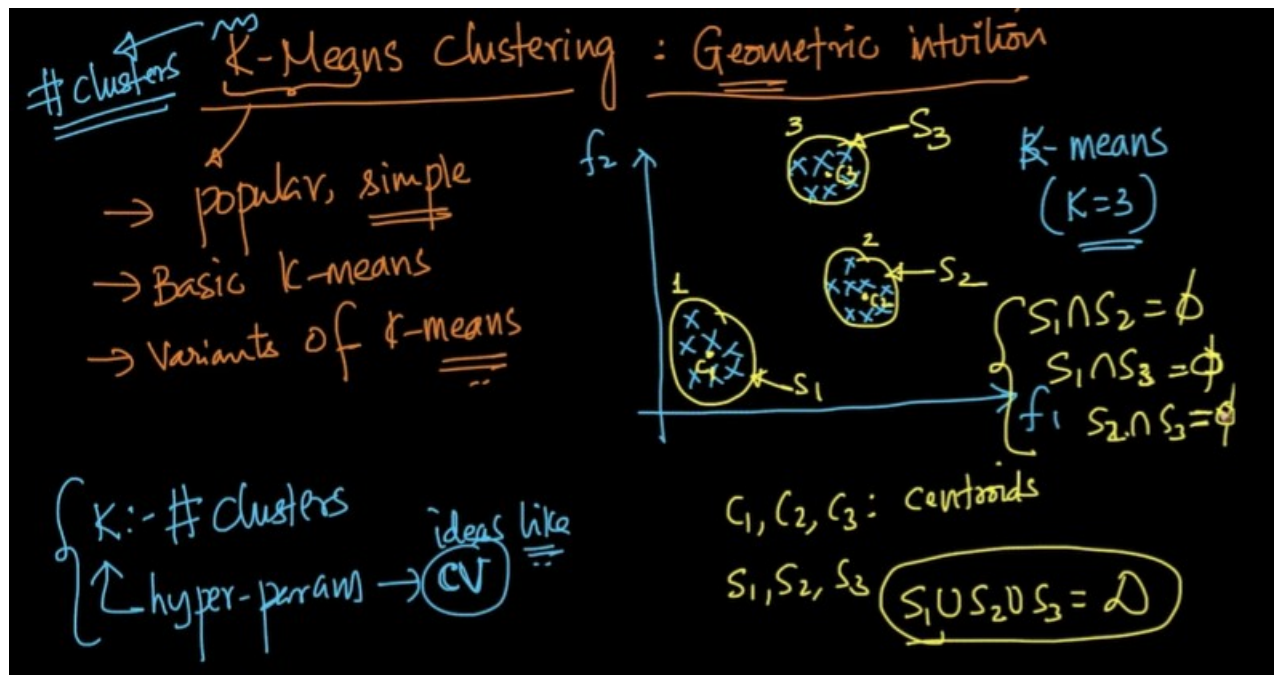
K – Means: Geometric intuition, Centroids

It is the popular clustering algorithm, variants of K- Means.

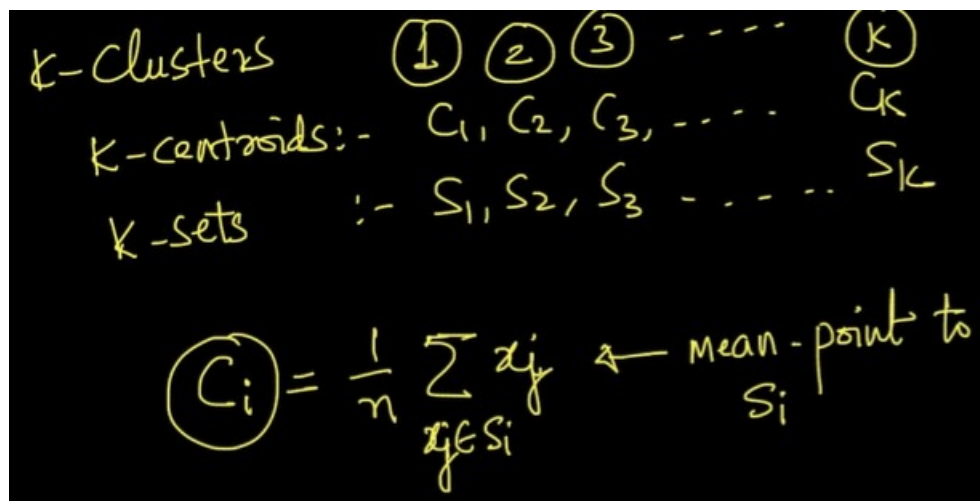
The K is the number of clusters which is the hyper parameter in K – Means.



K-Means groups every cluster, In this case we have three clusters. The intersection of the clusters is the null set.

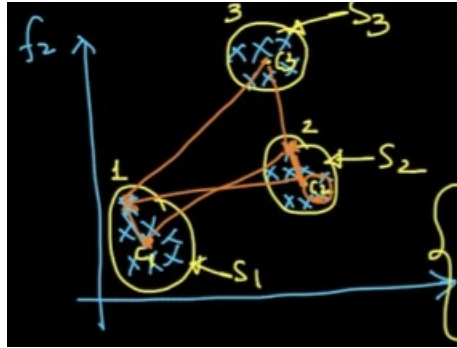


In K-Clusters, we have K centroids. K- clusters, K-sets of points. It is the geometric mean of the each cluster.

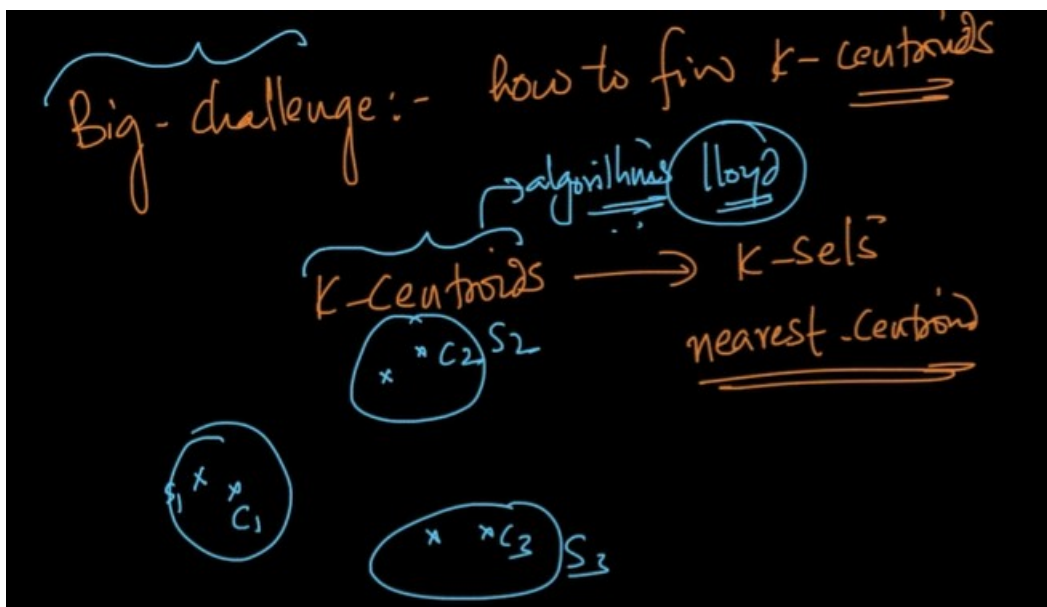


K – Means clustering is the centroid based scheme.
 There are other clustering called

Heriarical based clsutersing
 DBSCAN.



The challenge is how to find the K – Centroids. Once we get the K-centroids, we can compute the K-set nearest centroid. There are algos to find the K centroids.



To find the K central points and assign those points to the cluster.

K - Means: Mathematical formulation: Objective function

K-means : Mathematical formulation

$$D = \{x_1, x_2, \dots, x_n\}$$

Task: K-centroids: c_1, c_2, \dots, c_k

Sets: S_1, S_2, \dots, S_k

$$\begin{cases} \forall i, x_i \in S_j \\ \forall i, j, S_i \cap S_j = \emptyset \end{cases}$$

This is the optimization problem -

$$\arg \min_{c_1, c_2, \dots, c_k} \sum_{i=1}^k \sum_{x \in S_i} \|x - c_i\|^2$$

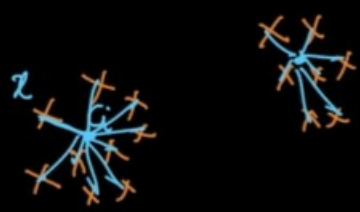
$$\text{s.t. } \begin{cases} x \in S_i \\ S_i \cap S_j \neq \emptyset \end{cases} \rightarrow \text{constraints}$$

$$\begin{matrix} \text{proximity} \\ \downarrow \\ c_1, c_2, \dots, c_k \\ \downarrow \\ S_1, S_2, \dots, S_k \end{matrix}$$

$$\text{inter-clust. dist}$$

$$\text{intra-clust. dist is minimized}$$

$$\text{dist of } x \text{ fr } c_i$$



This is sum of squared distances in the cluster I,

$$\text{argmin}_{c_1, c_2, \dots, c_k} \sum_{i=1}^k \sum_{x \in S_i} \|x - c_i\|^2$$

all clusters

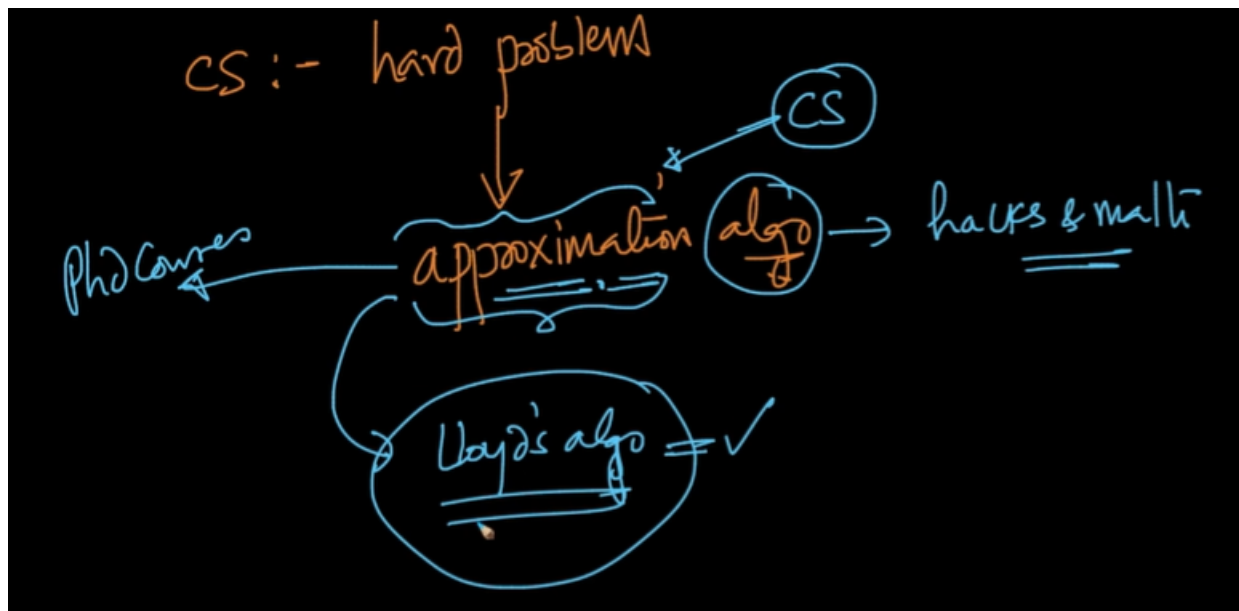
Sum of sq. dist for centroid in cluster i

V.V. hard to solve

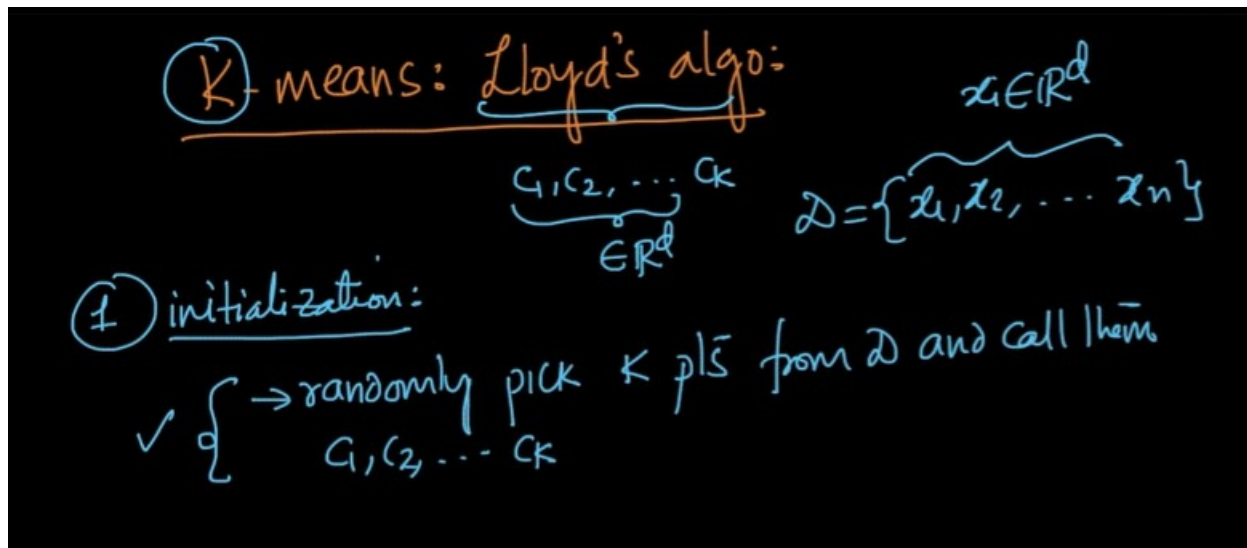
st $\setminus x_i \in S_i$
 $S_i \cap S_j = \phi$

This problem is very hard to solve. If there is a hard problem then we use the approximation of algo. Using some hacks and math.

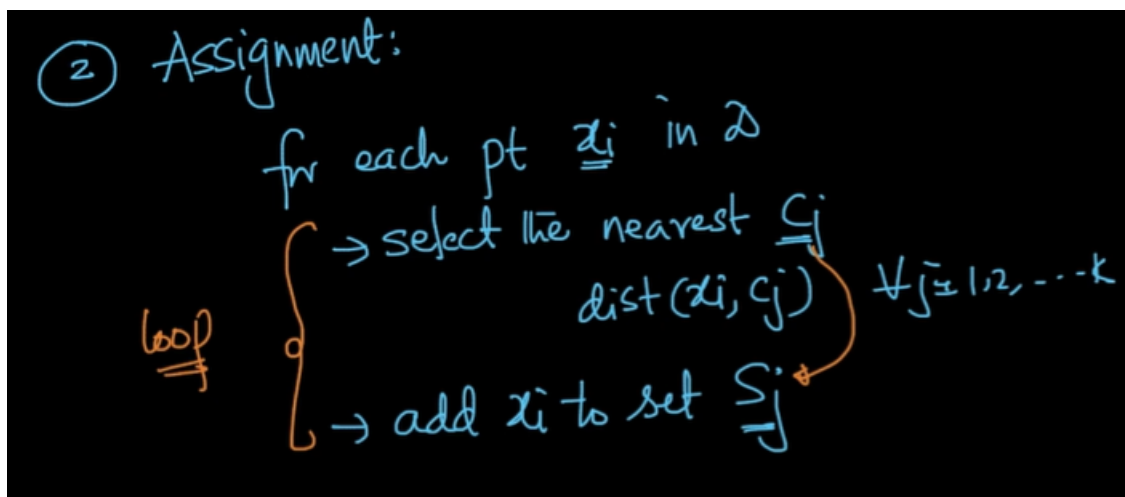
Lloyds algorithm: This solves the problem.



Lloyd's algorithm: K-means (we find the K centroids randomly)



The second step is assignment -



The third stage is called recompute centroid state: This is also called the update stage of centroids.

③ Recompute centroid / update
→ recalculate / update C_j 's as follows

$g \rightarrow s_j$

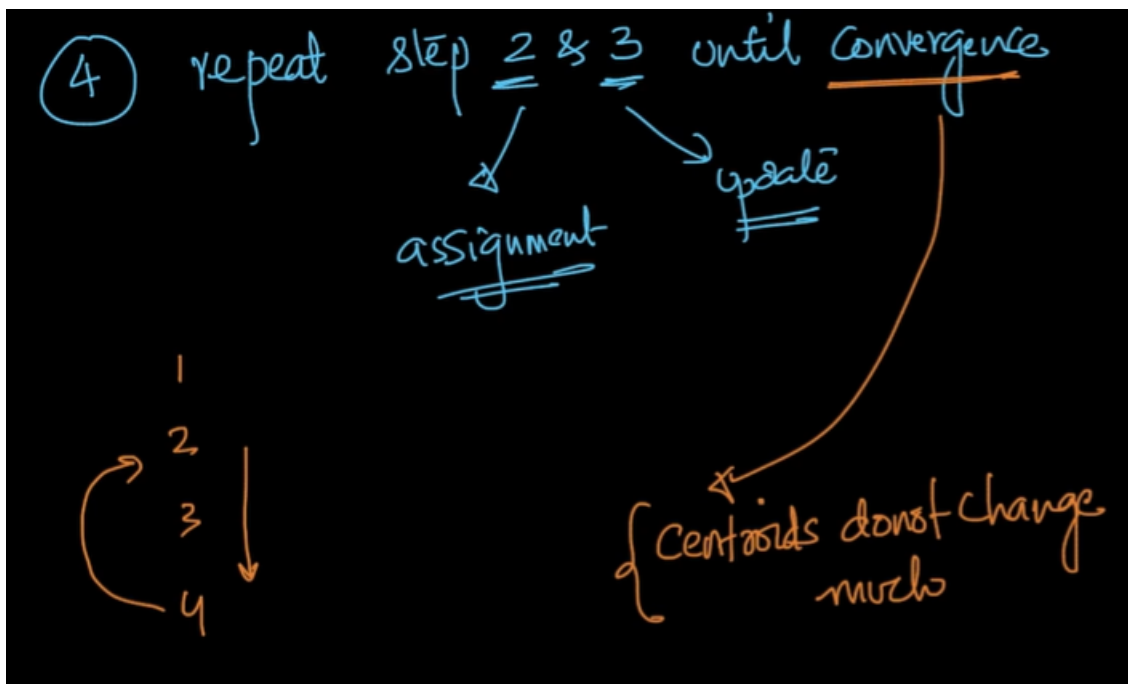
$$C_j = \frac{1}{|S_j|} \sum_{x_i \in S_j} x_i$$

C_1, C_2, \dots, C_k

pts in set S_j mean-pt

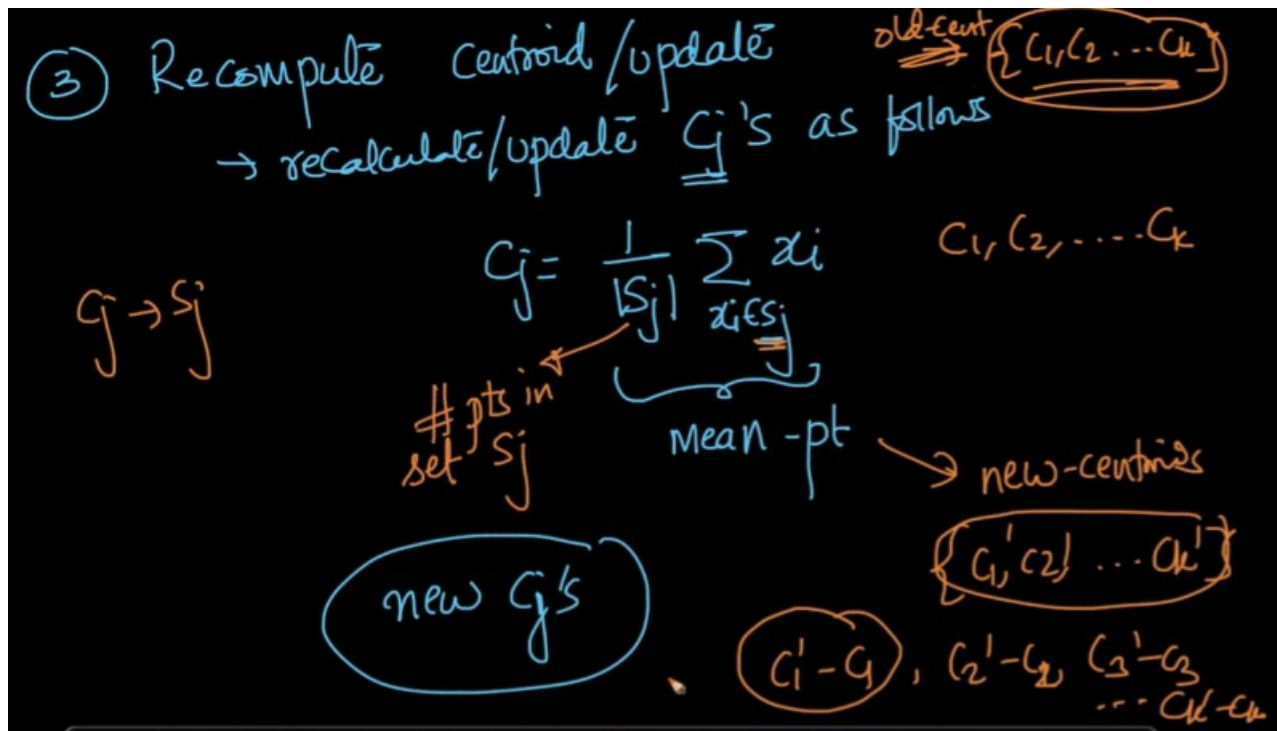
Step - 4:

We can repeat the step -2 and step -3, until convergence.

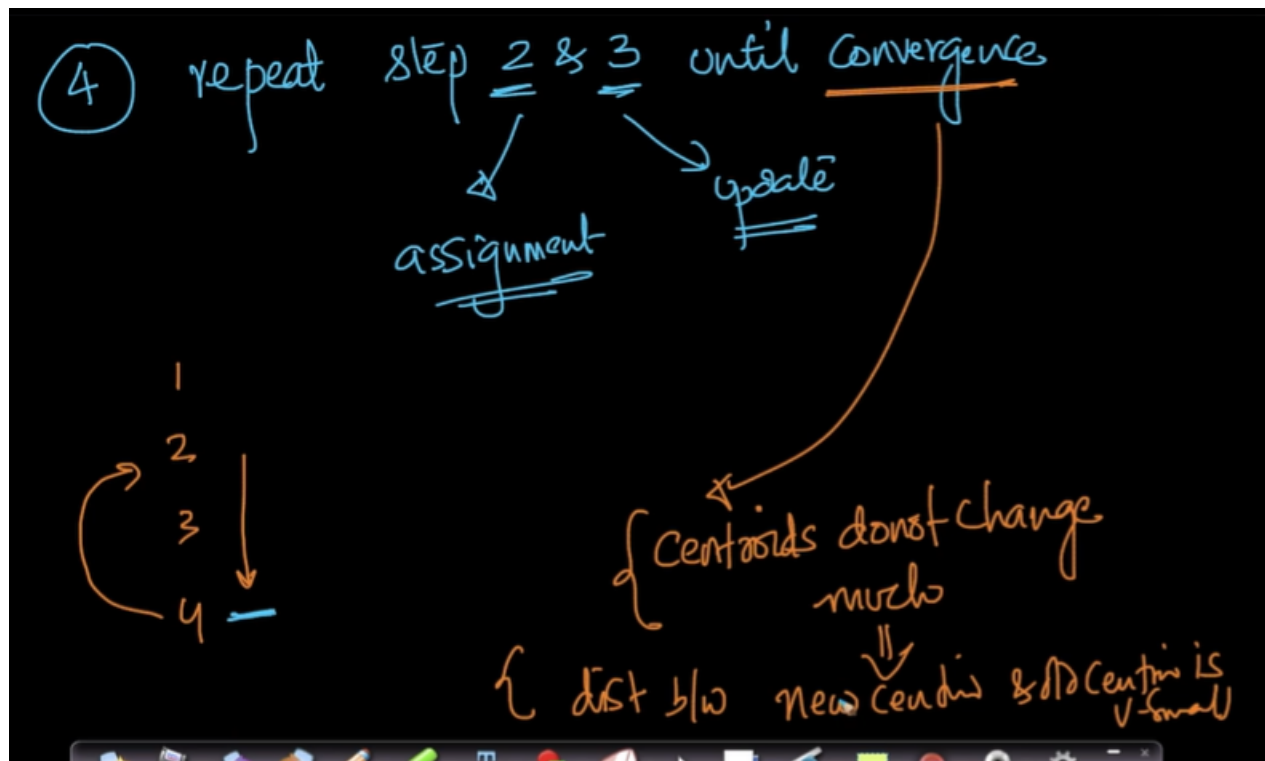


Here, we have the set of centroids, of new and old.

If there are no much change in the distance of the old and new points distances.



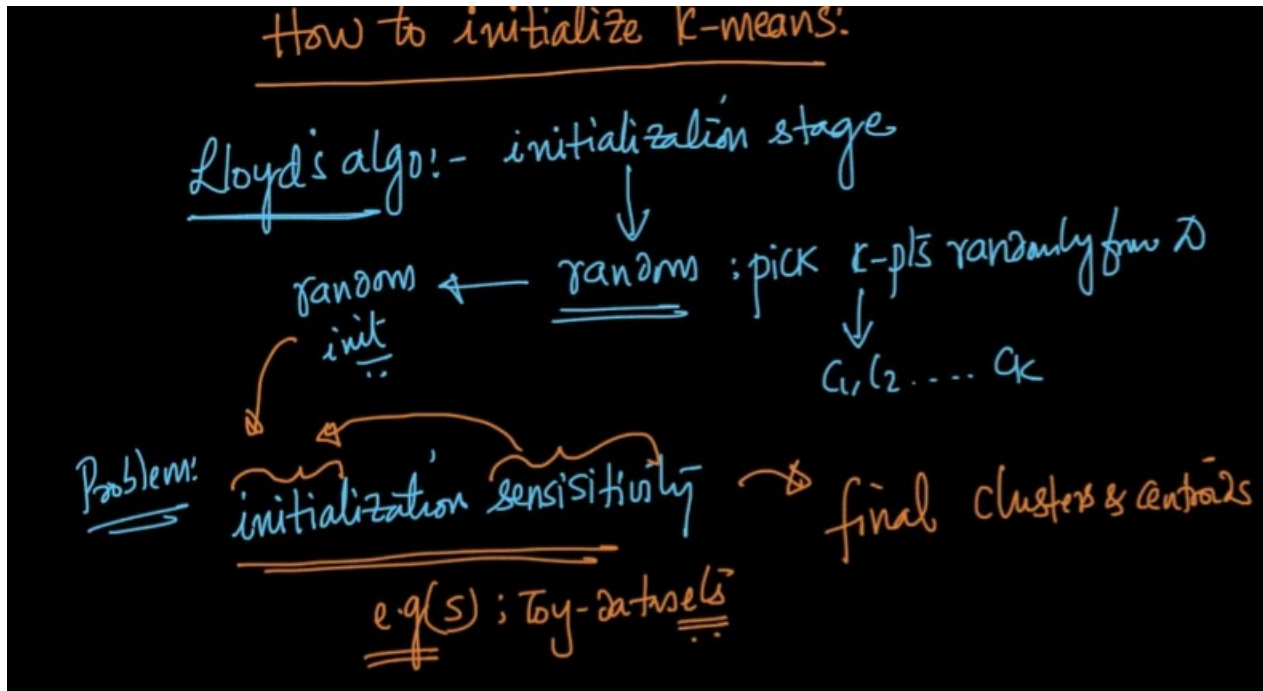
Distance between new centroids and old has no change.



The actual mathematical formulation is very hard. This is the Lloyd algorithm.

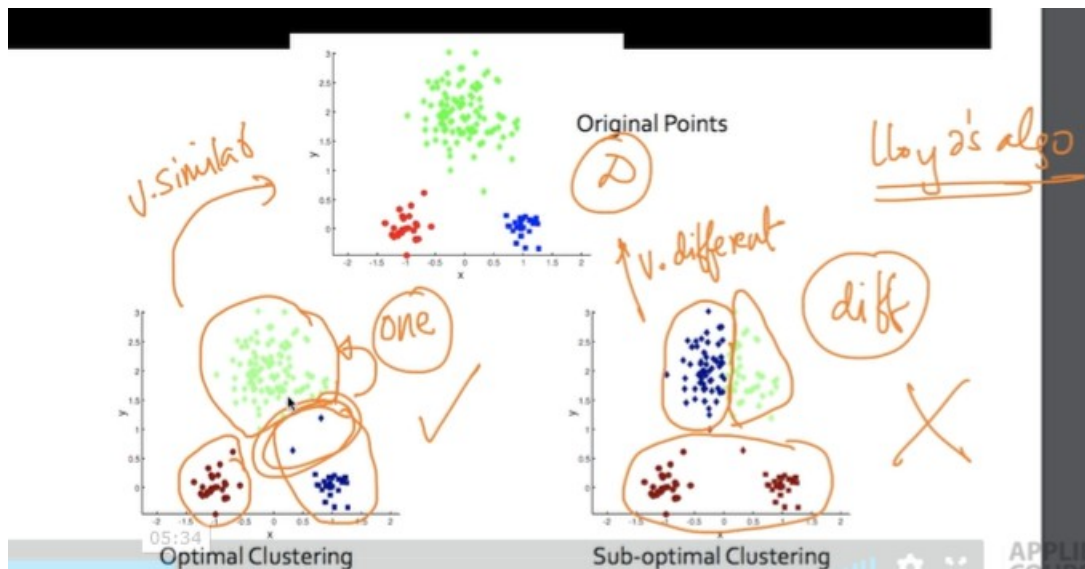
How to initialize: K-Means++

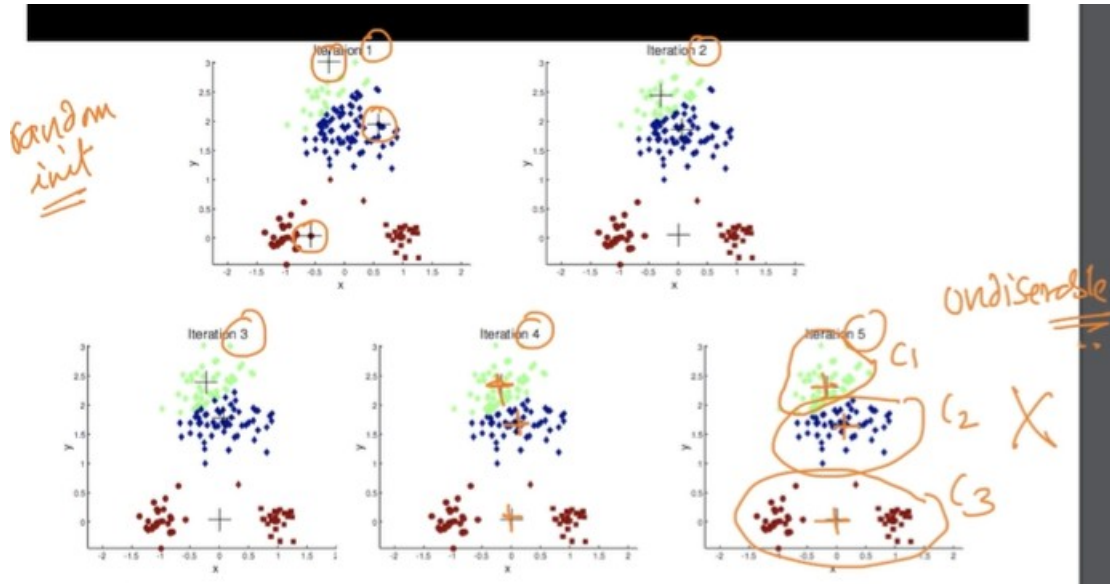
We can do the random initialization from the dataset and make the K centroids. There is one problem, initialization sensitivity.



Given this ideal dataset, when we initialize the centroids. We get the optimal and sub-optimal clustering states.

Applying the same Lloyd algorithm, we get the complete different solution.





- ① repeat k-means multiple times with different initializations
- ↳ pick the best clustering based on
- ✓ { smaller intra-clust }
 - ✓ { larger inter-clust }

The second way is K-Means ++, Instead of using the random initialization we use the smart initialization.

Step - 1:

Pick K centroids, we will pick the first centroid randomly C_1 .

② K-means++ $C \rightarrow C_{++}$

random-init \rightarrow smart init

initialization in k-means: (Task) pick $\{c_1, c_2, \dots, c_k\}$

① pick the first centroid randomly $\rightarrow C_1$
from D

② $\forall x_i \in D$ distribute as follows.

Step - 2:

② $\forall x_i \in D$

dist:- $x_i \rightarrow \text{dist}^2(x_i, \text{nearest centroid})$

$D - \{C_1\}$

C_2

C_1

{probabilistic approach}

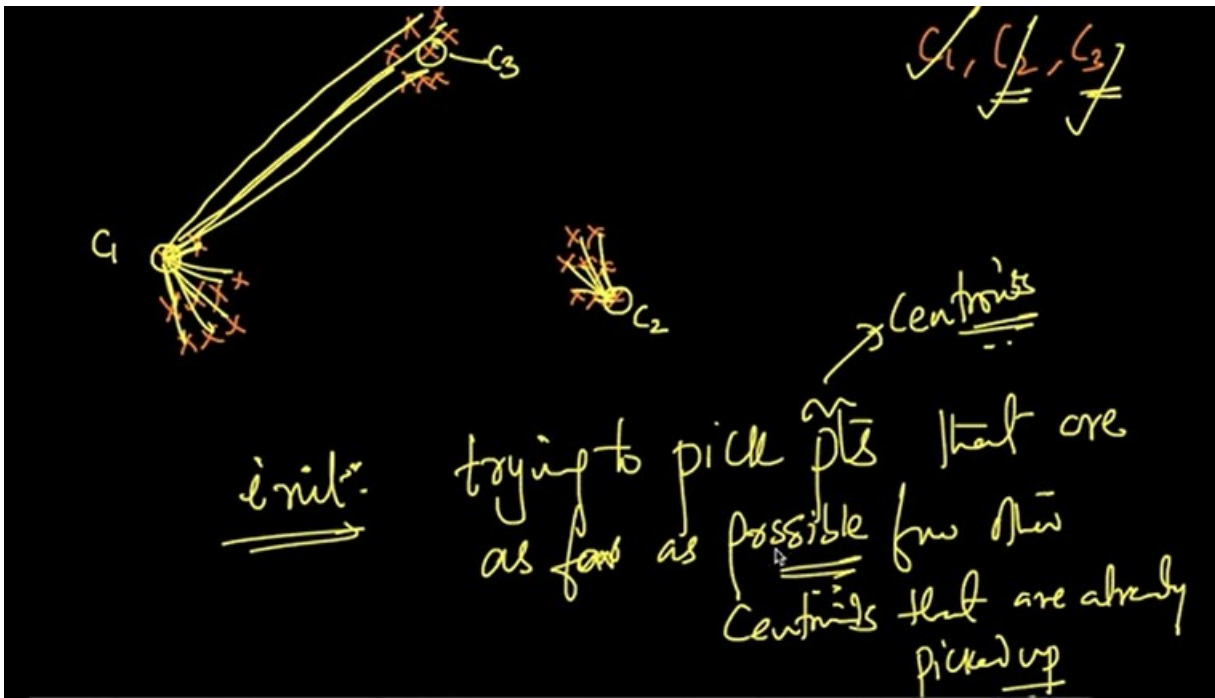
x_1	d_1
x_2	d_2
x_3	d_3
\vdots	\vdots
x_n	d_n

$\|(x_i - C_1)\|^2$

pick a pt fr $D - \{C_1\}$ with a prob. prop. to d_i

The chances of the points near to the centroid is very low.

In the initialization we want to pick the points that are far as possible from other centroids picked up.

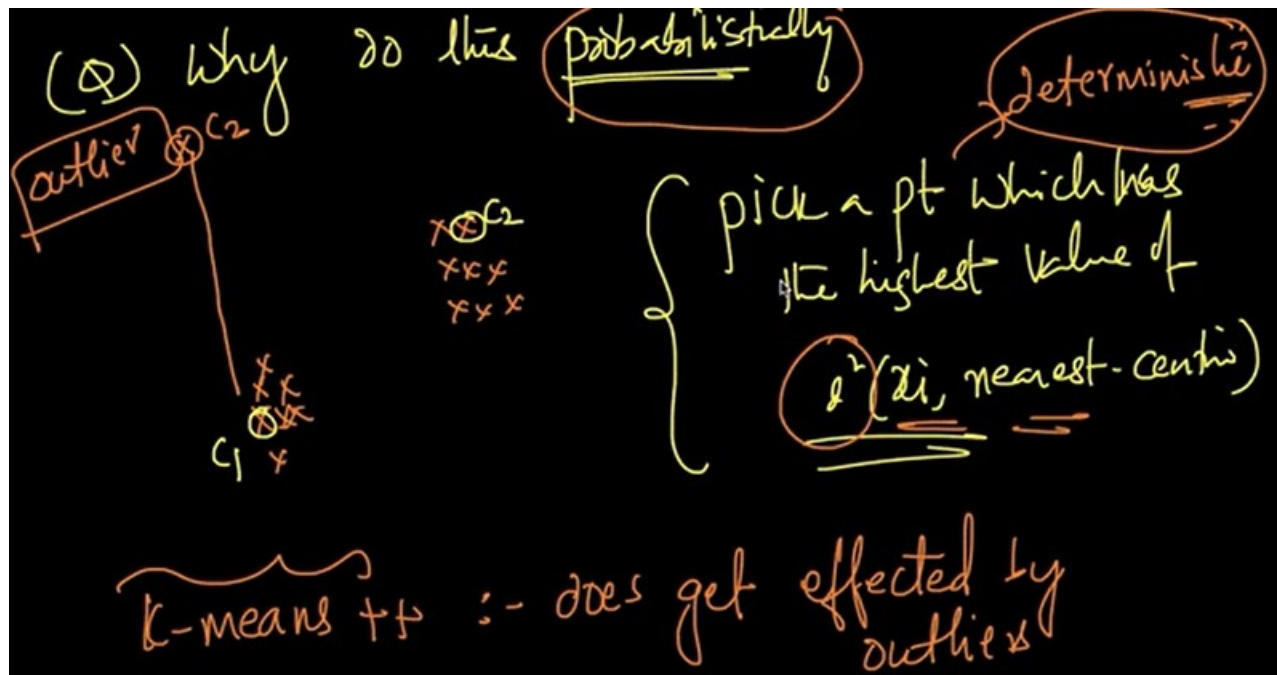


The points far to the chosen center has more probability to be the centroid.

We are trying to pick the points as far as possible that are already picked up.

Why cant I just pick the point that has highest value from the nearest center. We can have outliers.

We will pick the outliers as the centroid. K - Means ++ does get affected by outliers.

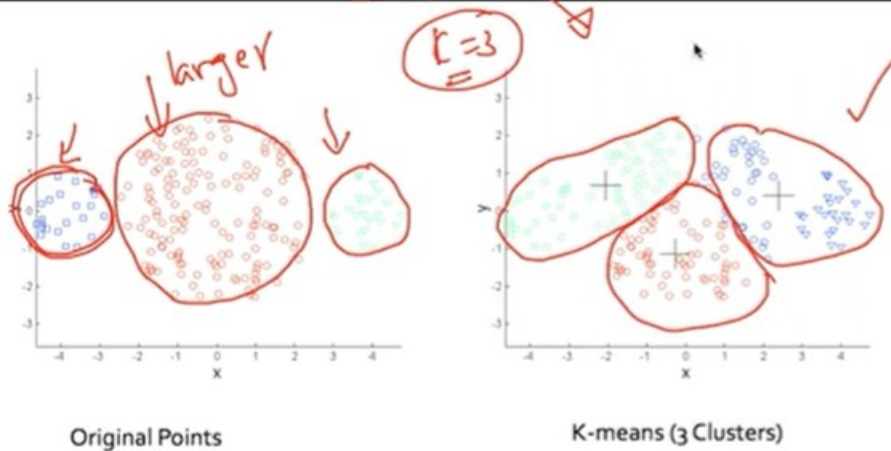


That is the whole reason we do it probabilistic-ally. To reduce the mitigation of being an outlier.

Failure cases/Limitations:

When we have clusters of different sizes, different Densities and Non – gobular shapes.

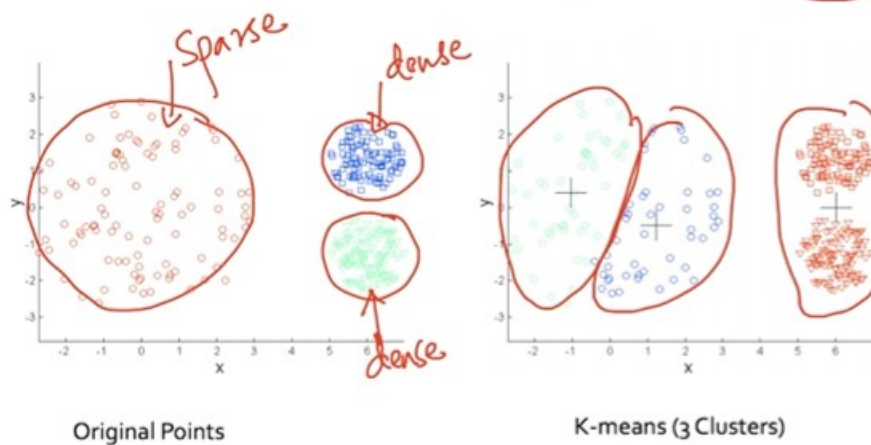
Limitations of K-means: Differing Sizes



Different densities:

K means tend to fail in case of different densities.

Limitations of K-means: Differing Density

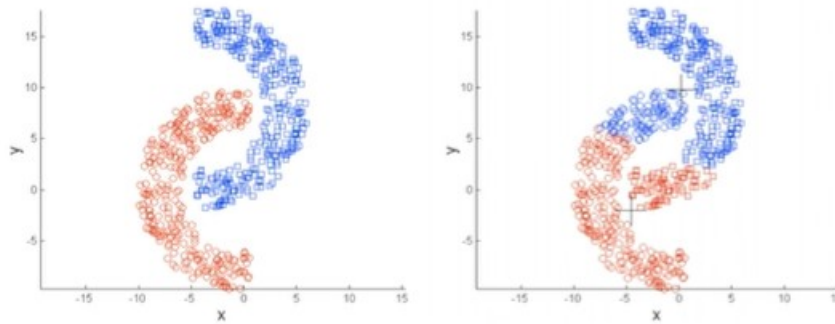


Non – gobular shapes:

If we give this data to the K – Means, we cannot work with gobular data.

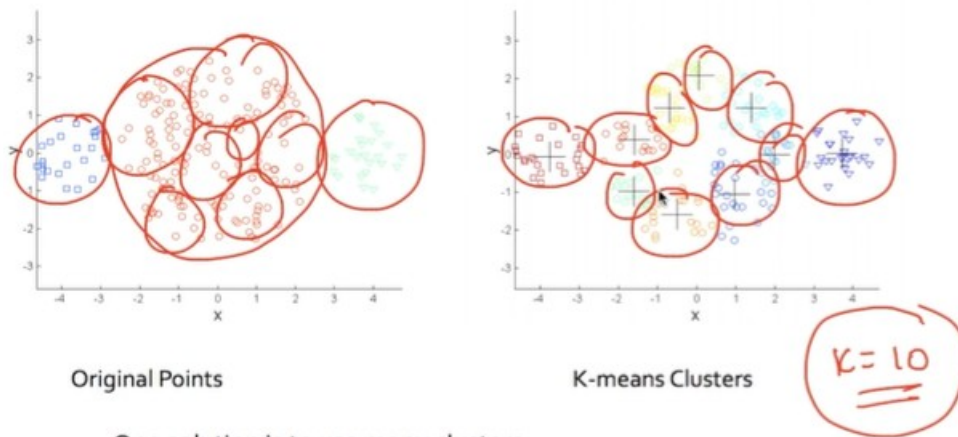
Limitations of K-means: Non-globular Shapes

non-convex set of pts



If we are continuing this K means then, if we keep $k = 10$, w get clusters like as follows:

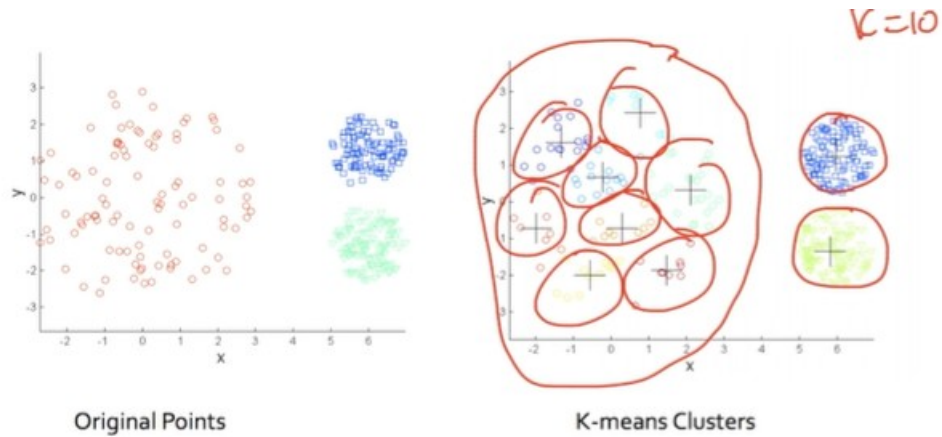
Overcoming K-means Limitations



One solution is to use many clusters.
Find parts of clusters, but need to put together.

TO find the parts of cluster, but need to put together.

Clusters of different densities:



K means is never perfect algorithm.

In case of non – globular structures, this is the best solution.

Evaluating clustering is not easy, there is no ground truth.

Limitations of K-means

- K-means has problems when clusters are of different

- Sizes
- Densities
- Non-globular shapes

- K-means has problems when the data contains outliers.

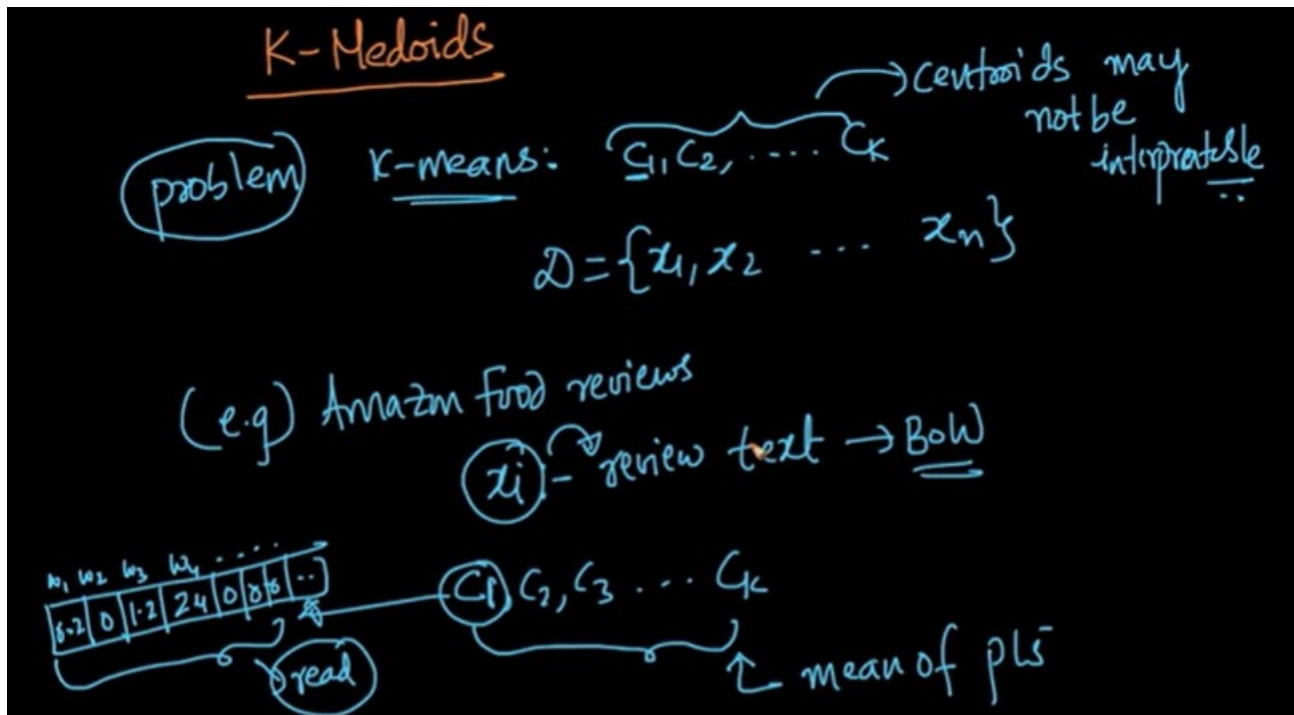
evaluating clustering
not easy

intracluster
intercluster

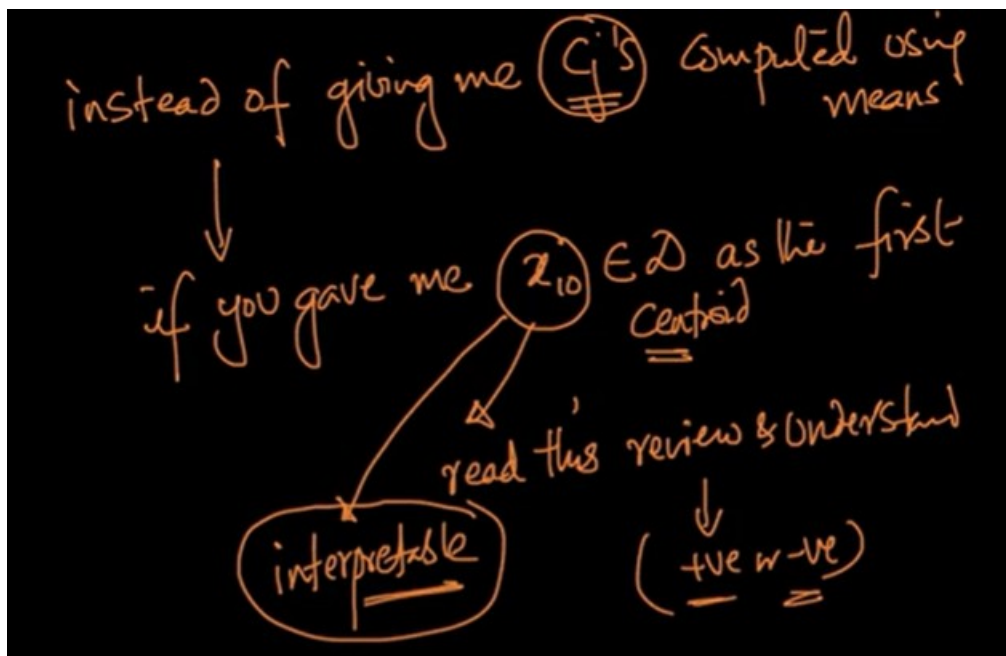
no y_i → ground truth

In the case of classification and regression there is no uncertainty in the results obtained.

K - Medoids: The centroids may not be interpret-able.



That means the centroids may be not interpretable in case of K-medoids.



When we represent the centroid as the data point in the dataset. Then it is called K-Medoids.

Especially in case of interpretation.

K - Medoids:

Partitioning around medoids(PAM) algorithm:

Partitioning around medoids (PAM) : K-medoids

① Initialization :- K-Means++ → probabilistic method
pick k pts from D

② Assignment :- closest medoid → same as in K-means
 $\{x_i \in S_j \text{ if medoid } j \text{ is the closest medoid to } x_i\}$

We swap each medoid with a non-medoid point in the dataset.

③ Update / recompute
✓ K-means :- $C_j = \frac{1}{|S_j|} \sum_{x_i \in S_j} x_i$ ✗

→ K-medoids (PAM)

(a) swap each medoid with a non-medoid pt

(b) if loss decreases, keep the swap else undo the swap

④ $k=2$

M_1 (x_1) x_2, x_3, x_4, x_5 M_2 (x_6) x_7, x_8, x_9, x_{10}
non-medoid non-medoid

What is loss in K-Means?

loss - in k-means

$$\min \sum_{i=1}^K \sum_{x \in S_i} \|x - m_j\|^2$$

\uparrow
medoid j

After swapping again compute the loss values.

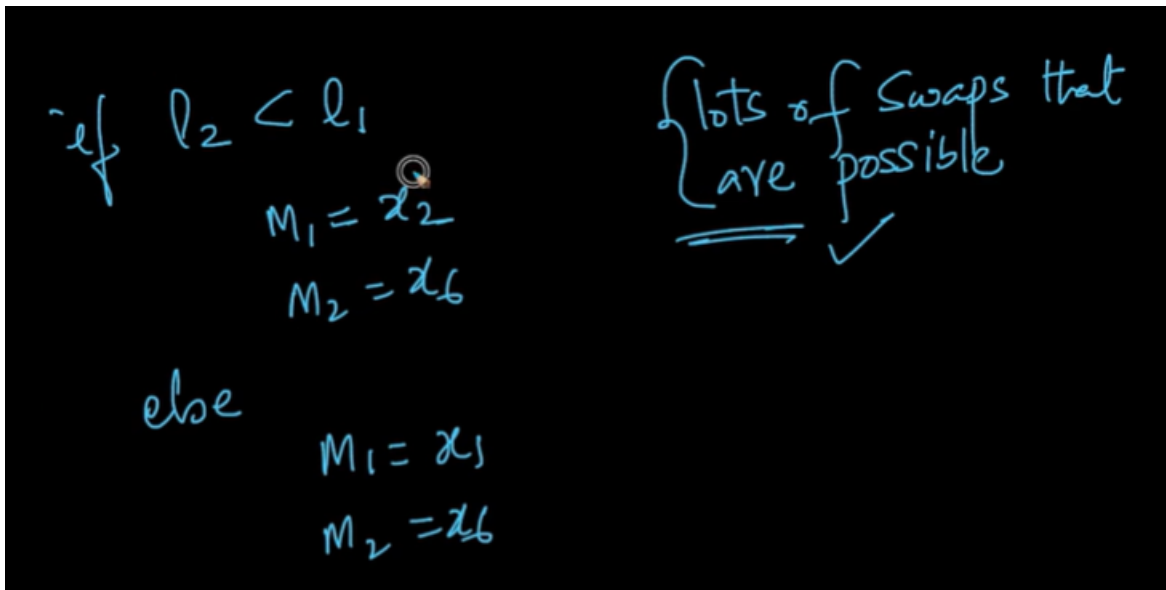
$m_1 \rightarrow x_1, x_2, x_3, x_4, x_5, x_6, x_7, x_8, x_9, x_{10}$
 $m_2 \rightarrow x_6$

(a) loss-value $x_1 = m_1; x_6 = m_2 \rightarrow l_1$

(b) swap $m_1 \leftrightarrow x_2$
 $m_1 = x_2; x_1$ as a non-medoid pt
 $m_2 = x_6$

(c) loss-value $m_1 = x_2$
 $m_2 = x_6 \rightarrow l_2$

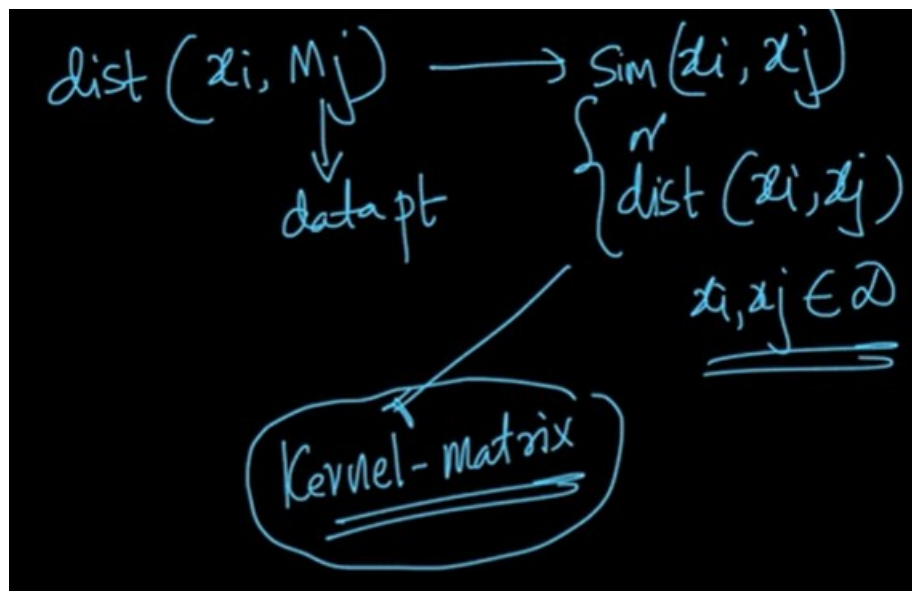
If loss decreases by swapping then we keep the point that decreases the loss. There are a lot of swaps possible.



The swap is successful, when the loss decreases. Medoid is also a data point.

We can use the kernel matrix or distance matrix and apply K – Medoids. The massive advantage is

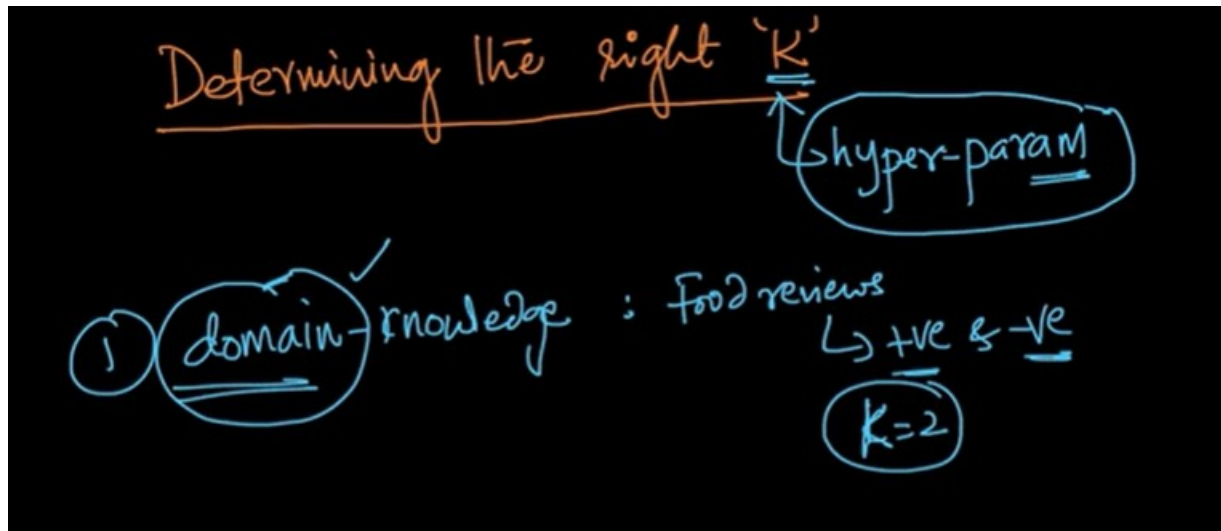
1. More interpretation.
 2. Kernelization.
- It is trivially kernelizable.



Determining the right K:

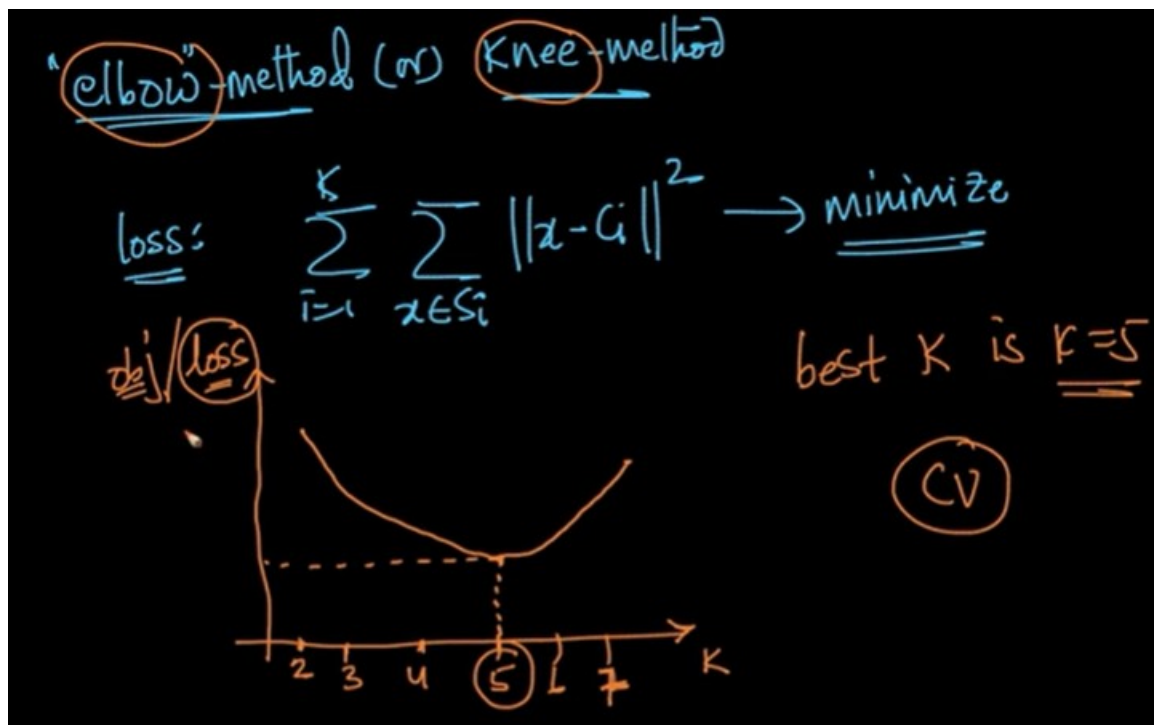
In K – means, K is the hyper parameter,

1. Domain knowledge – we can know the clusters.



Then we use the elbow-method (or) Knee-method -

We want to minimize the loss function is k means.



We only have data $D=\{x_i\}$.

Code examples:

Sklearn – sklearn.cluster.Kmeans

Examples

```
>>> from sklearn.cluster import KMeans
>>> import numpy as np
>>> X = np.array([[1, 2], [1, 4], [1, 0],
...               [4, 2], [4, 4], [4, 0]])
>>> kmeans = KMeans(n_clusters=2, random_state=0).fit(X)
>>> kmeans.labels_
array([0, 0, 0, 1, 1, 1], dtype=int32)
>>> kmeans.predict([[0, 0], [4, 4]])
array([0, 1], dtype=int32)
>>> kmeans.cluster_centers_
array([[ 1.,  2.],
       [ 4.,  2.]])
```

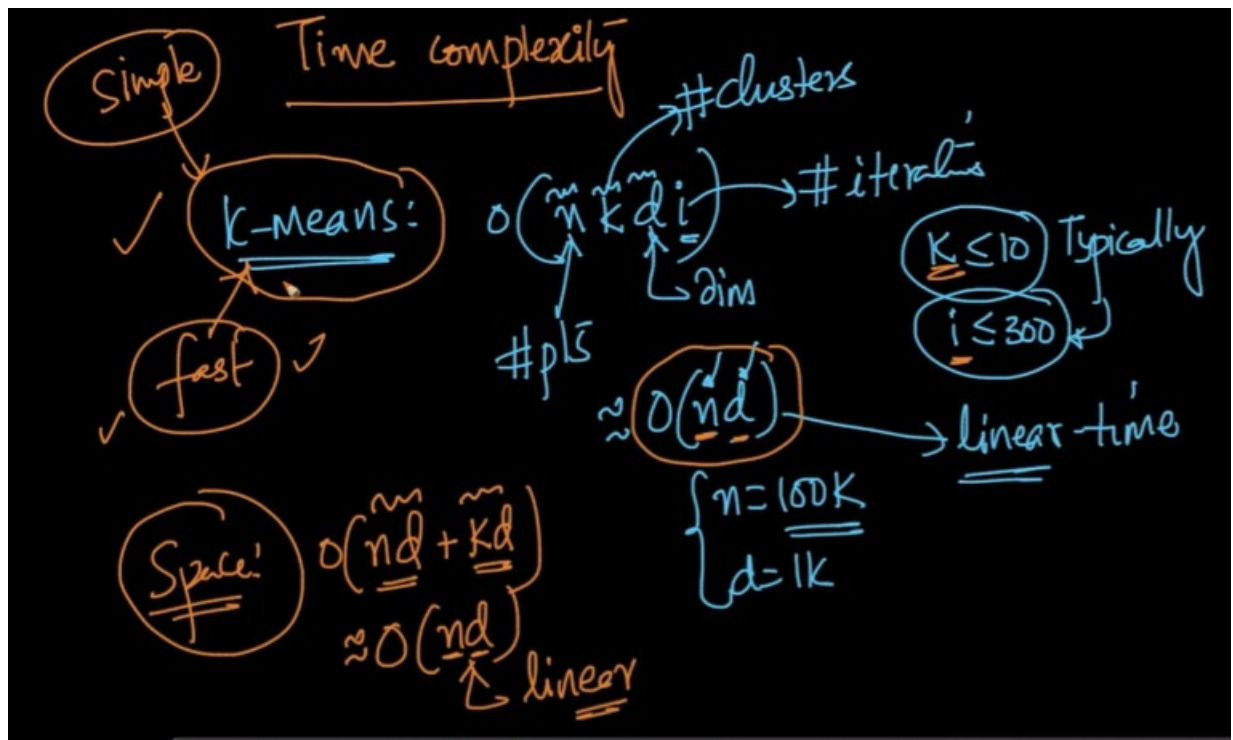
Methods

Time and space complexity:

K-Means: $O(nkdi)$ This is still a linear in time.

As far as space concerned we need to store $nd + kd$. Which is also linear.

It is fast and simple to understand.



Cluster Amazon reviews: