Machine Learning



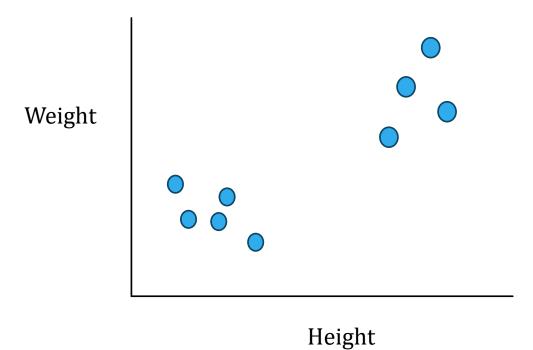
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Department of Computer Science & Engineering

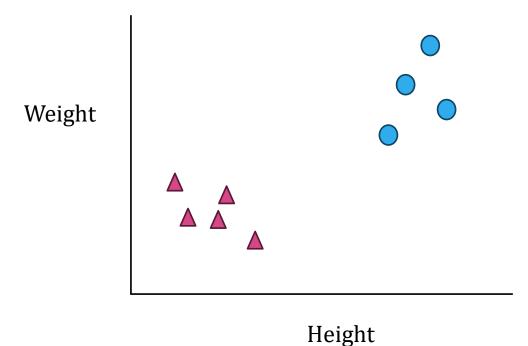
Clustering

What is Clustering?



I plot some sample data of a few animals

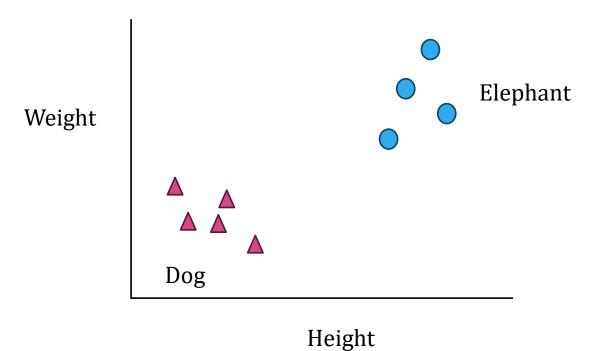
What is Clustering?



I can clearly see two groups here

In clustering, we create groups of samples in such a way that the samples in a group are more similar to each other than the samples in other groups

Classification

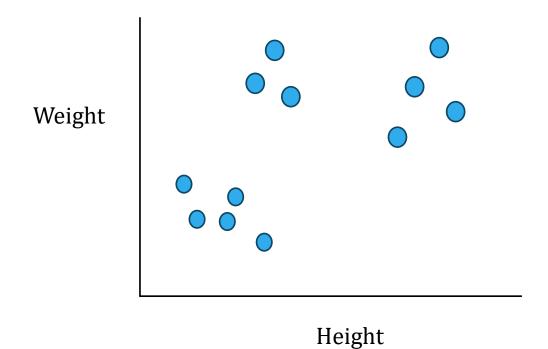


In supervised learning, we assign labels to each group

Clustering Techniques

- Centroid models: k-Means, k-medoids
- Graph-based models: Spectral Clustering
- Distribution models: Gaussian Mixture Models
- Density models: DBSCAN
- Connectivity Based: Hierarchical Clustering
- Neural models: Self-organizing map

Take the data points

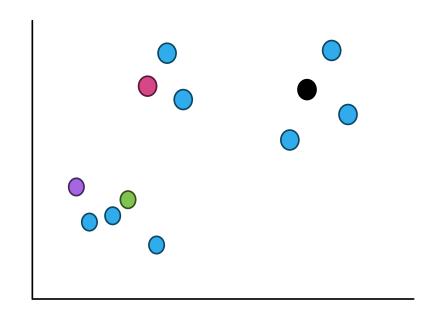


- Let's say, I want *m* clusters
- Here, let's take m = 4

Weight

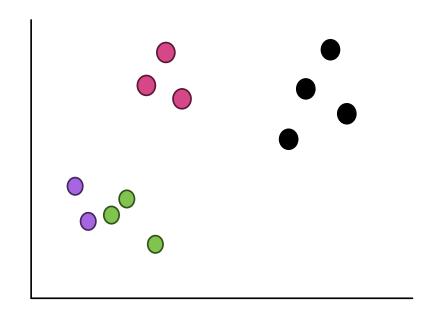
Height

- Let's say, I want *m* clusters
- Here, let's take m=4
- Randomly initialize m = 4 cluster centers



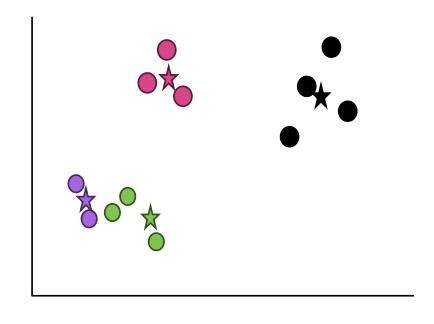
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- Let's say, I want m clusters
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- Randomly initialize m = 4 cluster centers
- Assign each of the other points to the nearest cluster center



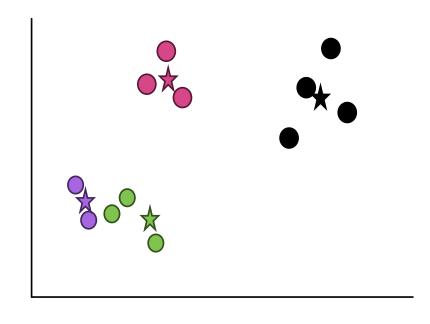
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- Let's say, I want m clusters
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- Assign each of the other points to the nearest cluster center
- Recalculate cluster mean



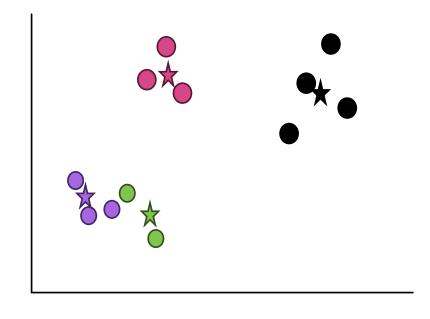
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- Randomly initialize m = 4 cluster centers
- Assign each of the other points to the nearest cluster center
- Recalculate cluster mean
- Reassign points based on updated mean



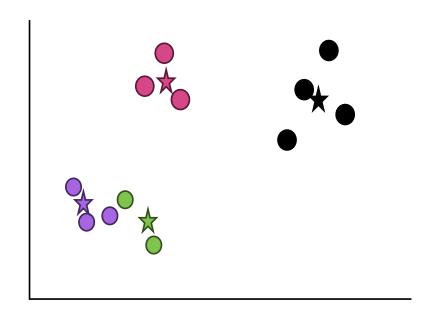
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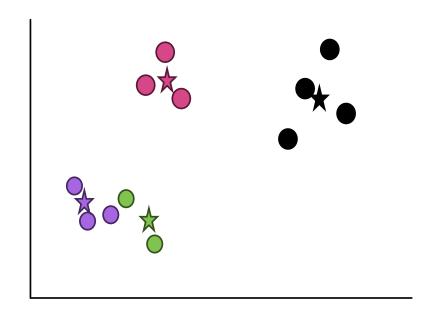
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- 5. Recalculate cluster mean
- 6. Reassign points based on updated mean
- 7. Go to step 5 if there is any change in the assignment. Otherwise, stop



Height

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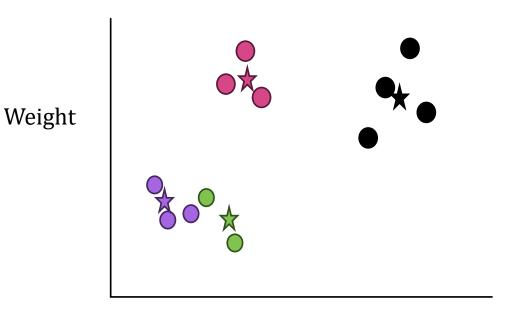
Weight



Height

Not guaranteed to converge

Assignment: How to find K?



Height

K-means: Limitations

Not guaranteed to converge

• Difficult to kind K

Sensitive to outliers

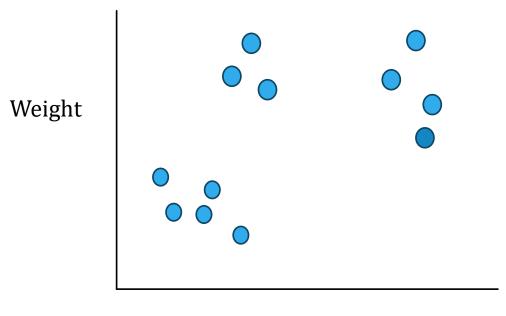
May be significantly affected by initialization

 A Medoid is a point in the cluster from which the sum of distances to other data points is minimal

 A Medoid is a point in the cluster from which dissimilarities with all the other points in the clusters are minimal.

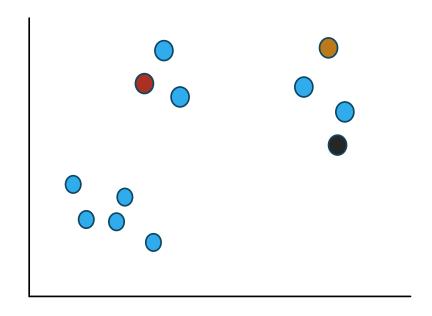
 Instead of centroids as reference points in K-Means algorithms, the K-Medoids algorithm takes a Medoid as a reference point

Take the data points



Height

- Take the data points
- Randomly assign medoids
- For each non-medoid data point, find the nearest medoid and assign the data point to the corresponding cluster



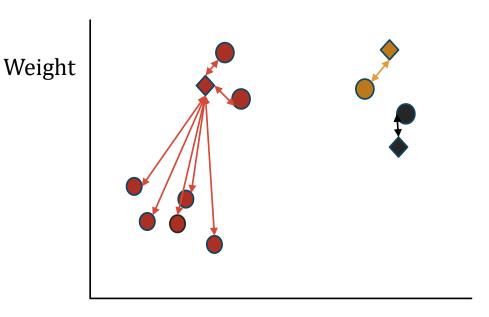
Height

- Take the data points
- Randomly assign medoids
- For each non-medoid data point, find the nearest medoid and assign the data point to the corresponding cluster
- Calculate the distance of every data point from the corresponding medoid. Sum of all these distances is called the cost

Weight

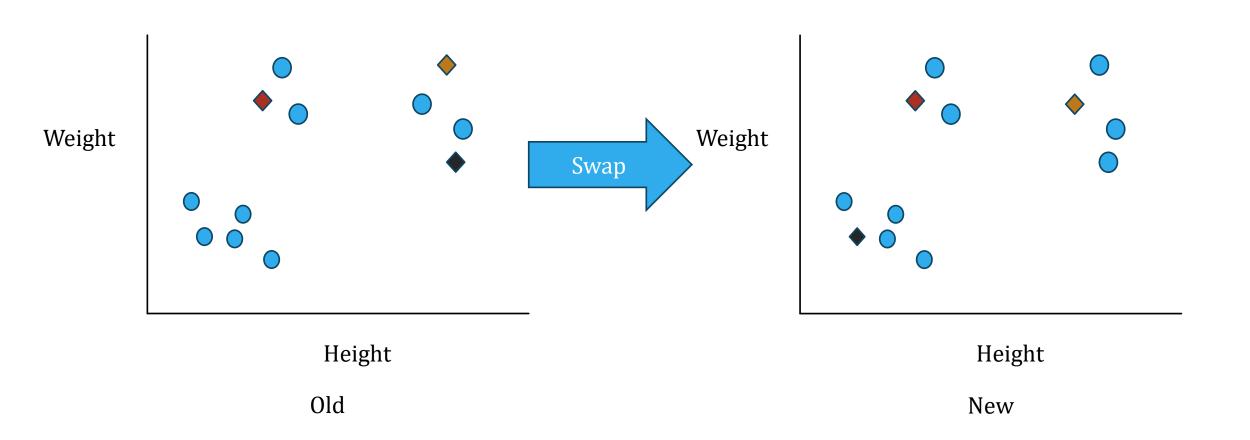
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- Randomly assign medoids
- For each non-medoid data point, find the nearest medoid and assign the data point to the corresponding cluster
- Calculate the distance of every data point from the corresponding medoid. Sum of all these distances is called the cost
 - Cost= total red distance + total yellow distance + total black distance

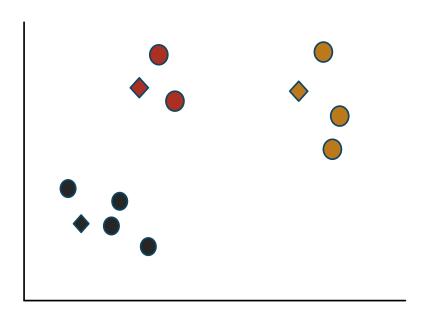


Height

Randomly swap the medoids

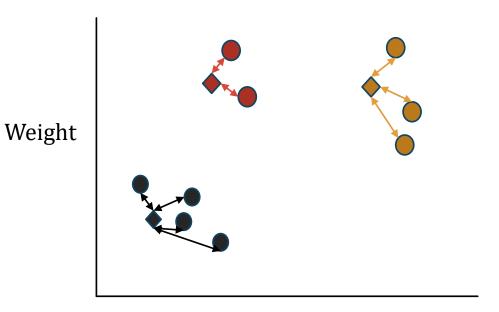


With the new medoids, create the clusters using nearest medoids



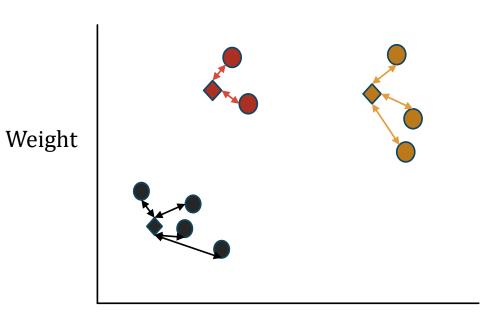
Height

- With the new medoids recalculate the cost
- Calculate the distance of every data point from the corresponding medoid. Sum of all these distances is called the cost
 - Cost= total red distance + total yellow distance + total black distance



Height

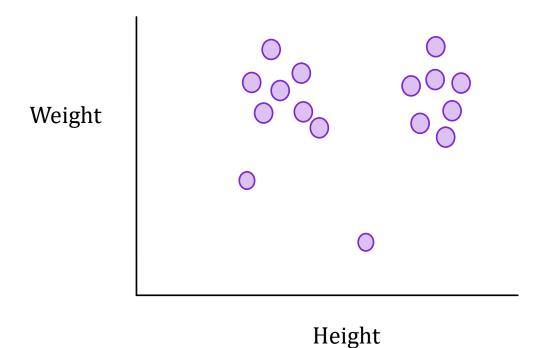
- With the new medoids recalculate the cost
- Calculate the distance of every data point from the corresponding medoid. Sum of all these distances is called the cost
 - Cost= total red distance + total yellow distance + total black distance
- If new cost > old cost, discard the new medoids and go back to the old medoids. The algorithm converges.
 - Else, keep the new medoids and redo the random swapping



Height

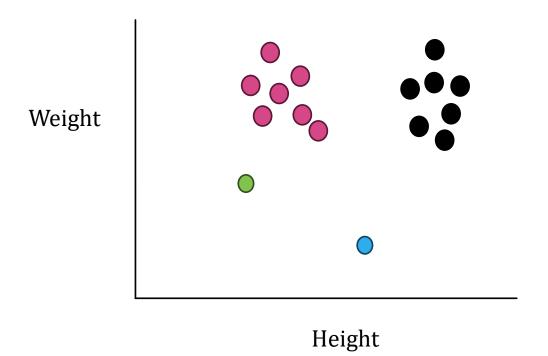
- **1. Select** *k* **random points from the dataset.** Select *k* random points from the dataset as the initial medoids. The medoids that are chosen are used to define the initial *k* clusters.
- **2. Assign data points to the cluster of the nearest medoid.** Assign each non-medoid to the cluster corresponding to the closest medoid.
- 3. Calculate the total sum of distances of data points from their assigned medoids for each medoid. Calculate the cost. Cost is given by the sum of the distances from a data point to the assigned (nearest) medoid.
- **4. Swap a non-medoid point with a medoid point and recalculate the cost.** Swap a non-medoid point with the medoids and repeat step 2 and 3 to calculate the cost with the new medoids.
- **5. Undo the swap if the recalculated cost with the new medoid exceeds the previous cost.** Check if the cost with new medoids is more than the cost with the old medoids. If that is the case, undo the swap, and the algorithm converges. Otherwise, go to step 4.

Clustering Techniques: Use of Density



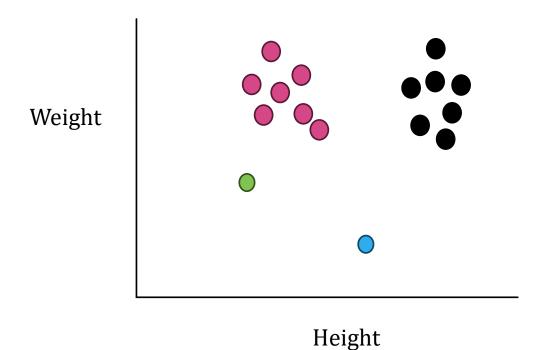
Can we say which points should form clusters?

Clustering Techniques: Use of Density



How did we do this visually?

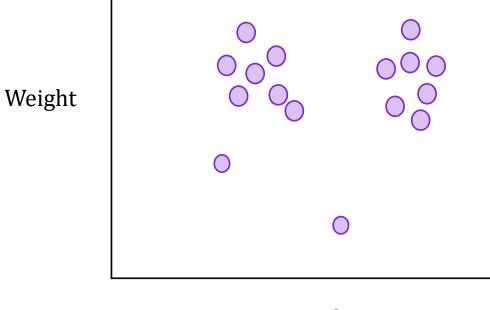
Clustering Techniques: Use of Density



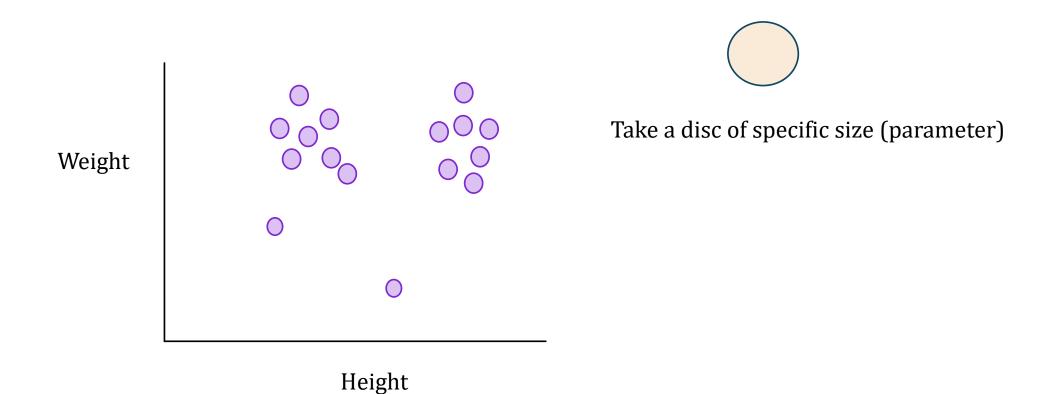
How did we do this visually?

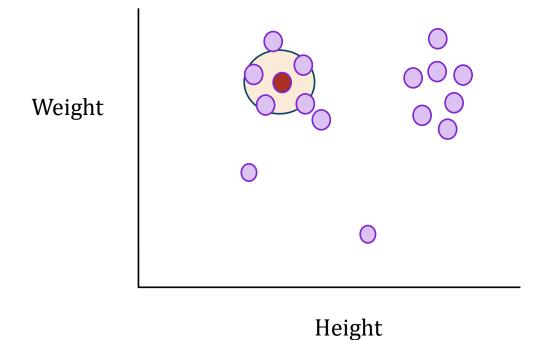
Based on density

DBSCAN: Density-Based Spatial Clustering of Applications with Noise



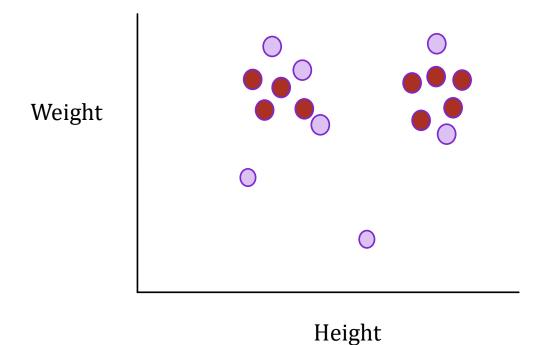
Height





Take a disc of specific size (parameter)

Put it across every point. If there are m number of points within the disc when positioned on point x_i , the point x_i will be called a core point. m is a parameter.



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Assign a cluster label in a core point (green)

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Put every point within the disk inside the first cluster

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Put every point within the disk inside the first cluster

Apply the disk from every point in the cluster

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Take a disc of specific size (parameter)

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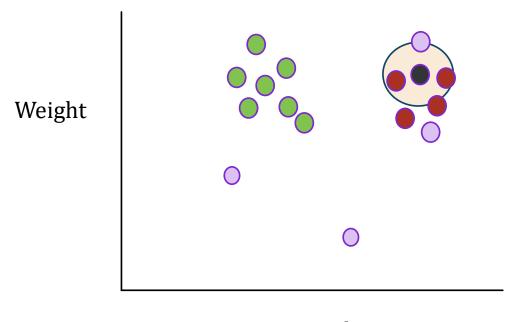
Red: core points

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Put every point within the disk inside the first cluster

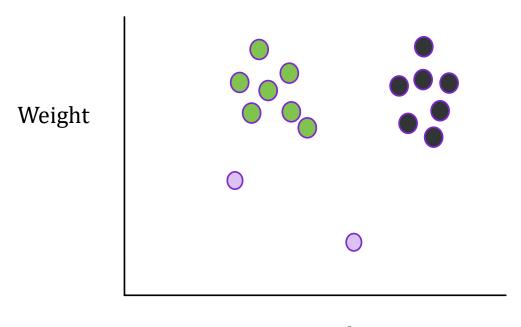
Apply the disk from every core point in the cluster

Repeat it for other core points

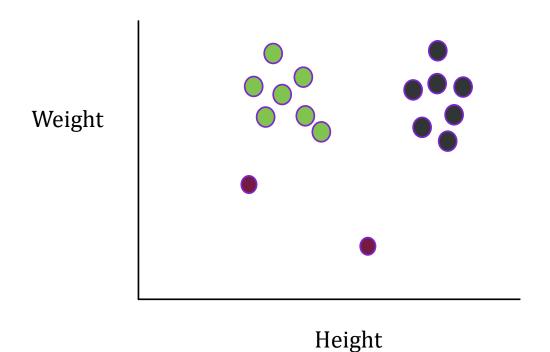


Height

Repeat it for other core points



Height

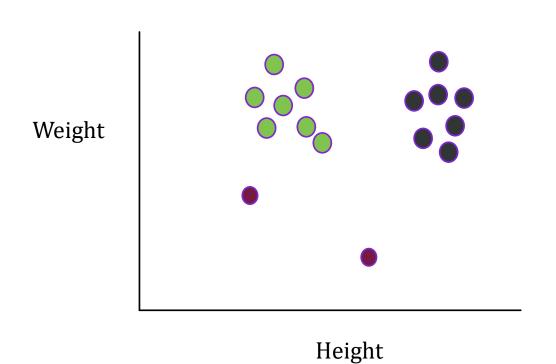


Repeat it for other core points

When we are done with all core points and left with only non-core points that can't be added to any clusters, we are done.

These non-core points are called outliers.

DBSCAN: Advantage



We don't need to know the number of clusters apriori

Weight

Height

The red point is a non-core point. So it can't expand a cluster.

Suppose, at start, the green point is chosen for expansion.

Weight

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The red point is a non-core point. So it can't expand a cluster.

Suppose, at start, the green point is chosen for expansion.

The red point will get included in the first cluster

Weight

Height

The red point is a non-core point. So it can't expand a cluster.

Suppose, at start, the green point is chosen for expansion.

The red point will get included in the first cluster

Weight

Height

The red point is a non-core point. So it can't expand a cluster.

But, at start, if the blue point is chosen for expansion,

Weight

Height

The red point is a non-core point. So it can't expand a cluster.

But, at start, if the blue point is chosen for expansion, the red point will get included in the second cluster

Weight

The red point is a non-core point. So it can't expand a cluster.

But, at start, if the blue point is chosen for expansion, the red point will get included in the second cluster

Height

Density-Based Spatial Clustering of Applications with Noise

• Let $S = \{x_1, x_2, ..., x_n\}$ be a set of points to be clustered

The goal is to identify points that form groups (nested clusters)

- Let $S = \{x_1, x_2, ..., x_n\}$ be a set of points to be clustered
- 1. Choose m > 0 and r
- 2. Let A_i be the set of points that lies withing a disc of radius r from x_i . Do it for every x_i
- 3. If $A_i < m$, we will not consider this x_i for our calculation
 - Points other than these points are called core points
- **4.** Take union of A_i and A_j if $A_i \cap A_j \neq \phi$
 - Go on doing it no more union operation is possible