Unsupervised Learning

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Unsupervised Learning

 Unsupervised learning involves training a model to find patterns in a dataset without pre-labeled responses.

Supervised vs. Unsupervised Learning

- Supervised Learning: Labeled data, goal is to predict outcomes.
- Unsupervised Learning: Unlabeled data, goal is to find hidden patterns.

Unsupervised Learning

Three common unsupervised learning algorithms:

- Clustering: Grouping similar items.
- Hierarchical Clustering: to build a hierarchy of clusters
- Dimensionality Reduction: Simplifying data without losing important information.

Clustering

- Clustering is a data mining technique which groups unlabeled data based on their similarities or differences.
- Clustering algorithms are used to process raw, unclassified data objects into groups represented by structures or patterns in the information.
- Clustering algorithms can be categorized into a few types, specifically
 - Exclusive
 - Hierarchical

Clustering in market segmentation



Clustering

Cluster 1: High income/high property value

Cluster 2: Middle income/middle property value

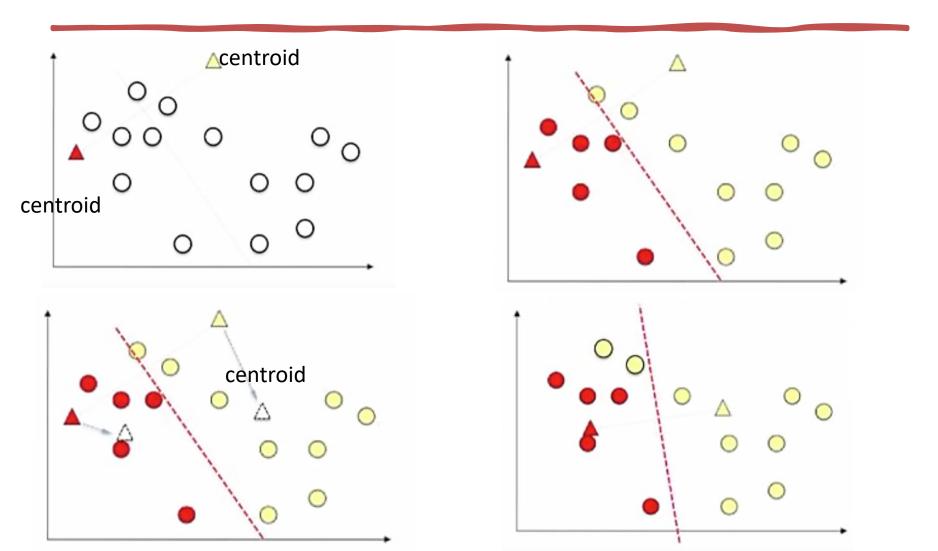
Cluster 3: High income/low property value

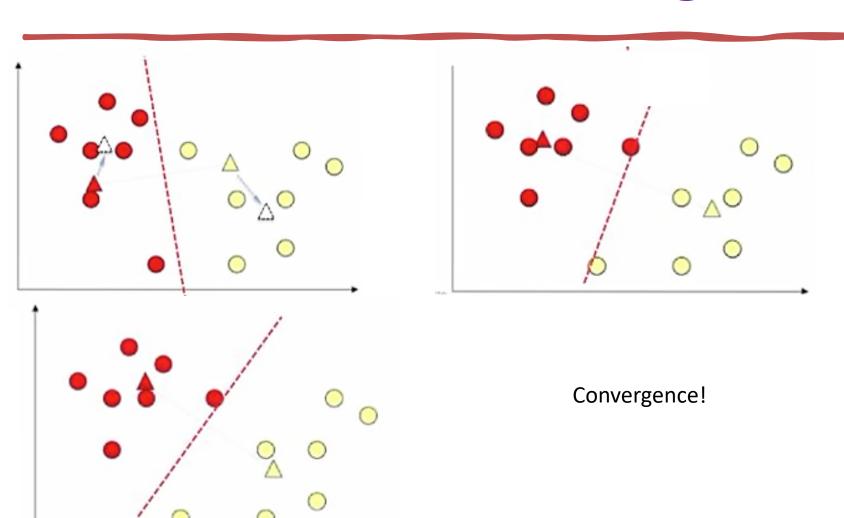
- The K-means clustering algorithm is an example of exclusive clustering.
- K-Means clustering aims to group n data points into k clusters in which each observation belongs to the cluster with the nearest mean.
- A larger K value will be indicative of smaller groupings whereas a smaller K value will have larger groupings and less granularity.
- K-means clustering is commonly used in market segmentation, document clustering, and image segmentation.

- input k, set of points $x_1, x_2, ..., x_n$
- starts by placing K points (centroids: $c_1, c_2, ..., c_k$) at random locations in space.
- repeat until convergence:
 - for each point x_i :
 - find nearest centroid c_i
 - assign the point x_i to cluster j

$$c_j(a) = \frac{1}{n_j} \sum_{x_j \to c_j} x_i(a)$$
 for $a = 1, ...$

- for each cluster j = 1, 2, ..., k:
 - new centroid c_j = mean of all points x_i assigned to cluster j in previous step





KMeans

```
class sklearn.cluster.KMeans(n_clusters=8, *, init='k-means++',
n_init='auto', max_iter=300, tol=0.0001, verbose=0, random_state=None,
copy_x=True, algorithm='lloyd')
```

initialization method of centroids

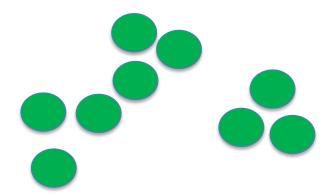
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Number of clusters

Selecting K! Question of granularity

How coarse or fine-grained is the structure in your data?

The main problem of K-means is what a good k? No clustering algorithm can pick k.



How many clusters do you see?

Instead of picking K, let's find a hierarchy of the structure:

At the top level – coarse grained At low levels – fine grained

How to cluster:

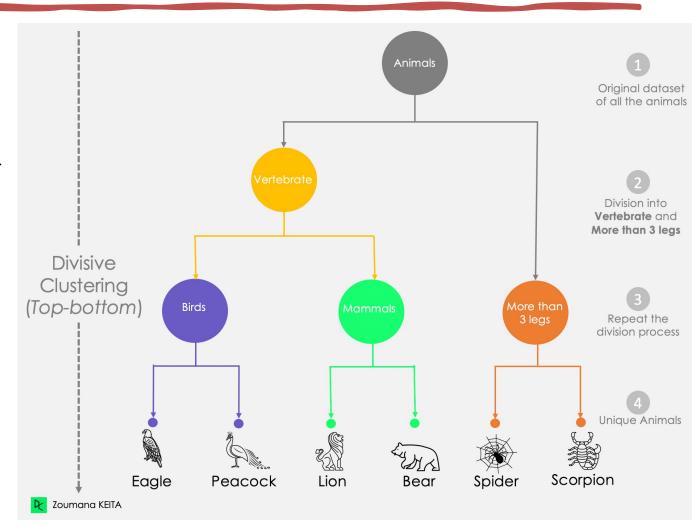
Topmost cluster → have every points in the cluster

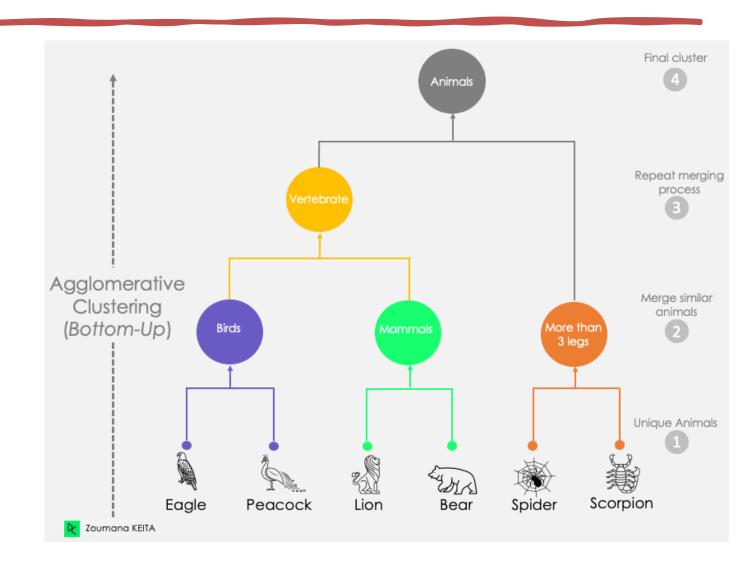
Bottom clusters \rightarrow have n set of singletons

So, you will end up with a tree!

Two types of algorithm

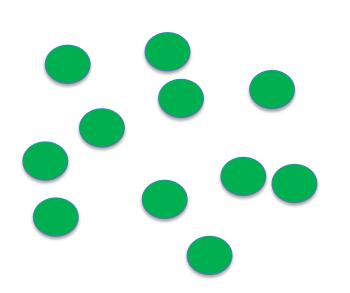
- Divisive(Top-bottom)
- agglomerative (Bottomup)

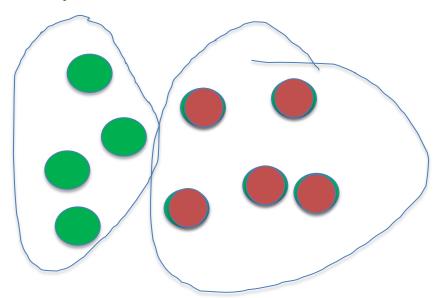




Hierarchical algorithm (top-down approach):

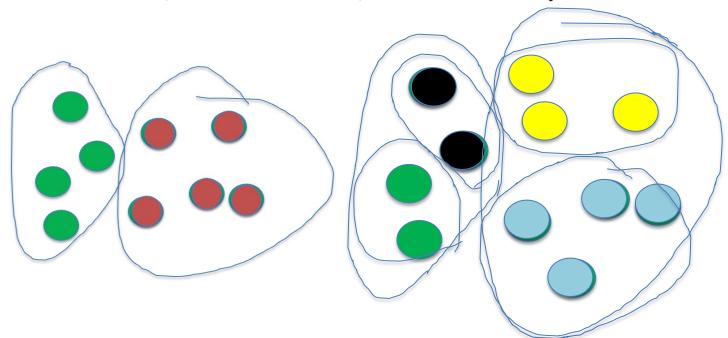
run K-means algorithm on the original data $x_1, x_2, ..., x_n$ for each of the resulting clusters c_i , i = 1, 2, ..., k recursively run K-means on points in cluster c_i





Hierarchical algorithm (top-down approach):

- run K-means algorithm on the original data x_1 , x_2 , ... , x_n
- for each of the resulting clusters c_i , i = 1, 2, ..., k
 - recursively run K-means on points in cluster c_i



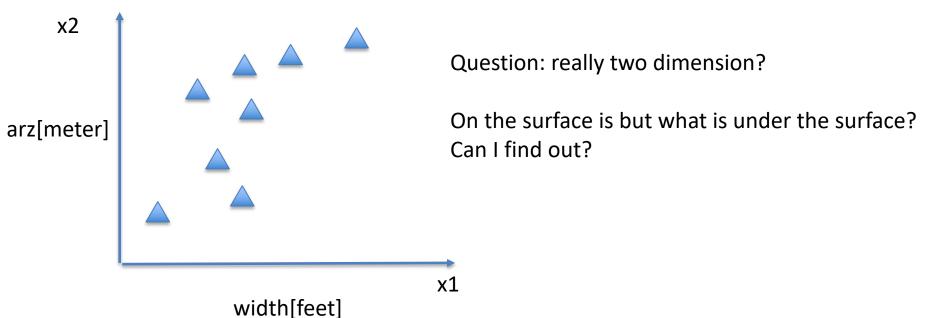
Applications of Hierarchical Clustering

- Biology
 - The clustering of DNA sequences is one of the biggest challenges in bioinformatics.
- Image processing
 - Hierarchical clustering can be performed in image processing to group similar regions or pixels of an image in terms of color, intensity, or other features.
- Social network analysis
 - Hierarchical clustering can be used to identify groups or communities and to understand their relationships to each other and the structure of the network as a whole.

scipy.cluster.hierarchy.linkage(y, method='sin gle', metric='euclidean', optimal_ordering=Fal se)

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- Suppose we have a dataset and we want to find the dimensionality of the data
- Each data point represent a pair (x, y), where x and y are real numbers.



- A dataset including 5 features:
 - X1: number of car accidents
 - X2: number of burst water pumps
 - X3: number of school closure
 - X4: number of patients with broken legs or arms
 - X5: snowplow cost
 - All of these are strongly influence by one factor! What is that?

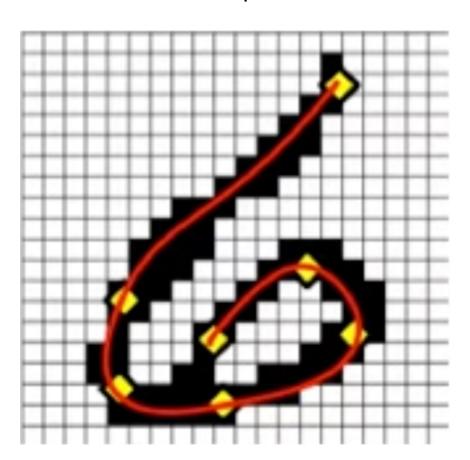
Applications:

Texts - each word a separate feature

Image - each pixel a separate feature

Millions of features!!!
High dimentional

Example 1

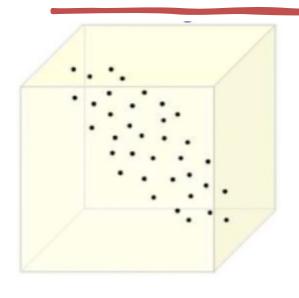


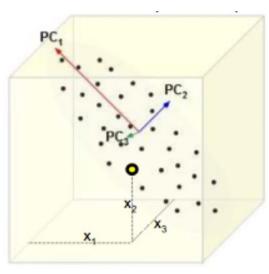
Example 2

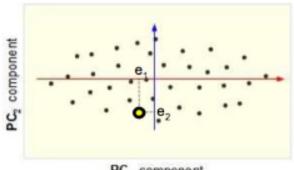
"Grouping, clustering, and categorizing data points into nested, ordered, and structured layers exemplifies the process of hierarchical algorithms."

"Grouping data into nested layers illustrates hierarchical algorithms."

- We suppose we have our low dimensional data embedded in a high dimension space.
- Defines a set of principal components
 - 1st: direction of the greatest variability in the data
 - 2nd: perpendicular to 1st, greatest variability of what's left
 - Repeat this until d (original dimensionality)
- First m<<d components become m new dimensions
 - Change coordinates of every datapoint to these dimensions







PC, component

PCA

```
class sklearn.decomposition.PCA(n_components=None, *, copy=True, whiten=False, svd_solver='auto', tol=0.0, iterated_power='auto', n_oversa mples=10, power_iteration_normalizer='auto', random_state=None)
```

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