

Machine Learning - Session 2

Unsupervised Learning





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 is no known output, no teacher to instruct the learning algorithm.
- In unsupervised learning, the learning algorithm is just shown the input data and asked to extract knowledge from this data.
- In contrast with supervised learning, in unsupervised learning, there is no direct measure of success.





Some tasks include

- Clustering
- Anomaly Detection
- Density estimation
- Dimensionality Reduction
- Association Rules





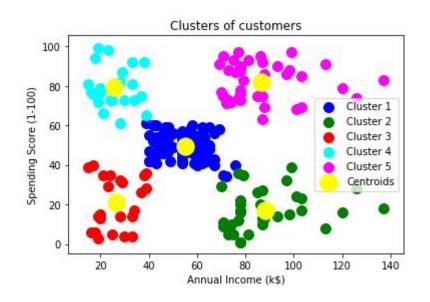
Clustering:

- The goal is to group similar instances together into clusters.
- Clustering is a great tool for data analysis, customer segmentation, recommender systems, search engines, image segmentation, semi-supervised learning, dimensionality reduction, and more.





Clustering







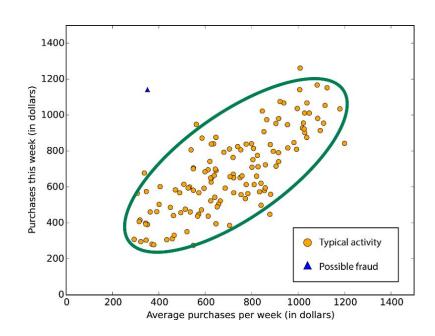
Anomaly Detection:

 The objective is to learn what "normal" data looks like, and then use that to detect abnormal instances, such as defective items on a production line or a new trend in a time series.





Anomaly Detection





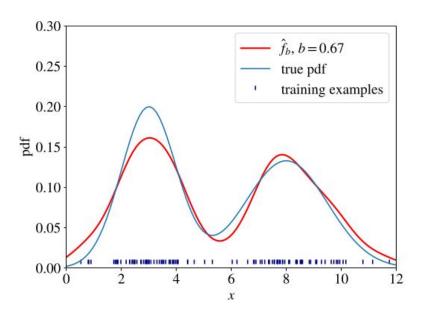
Density estimation:

- This is the task of estimating the probability density function (PDF) of the random process that generated the dataset.
- Density estimation is commonly used for anomaly detection: instances located in very low-density regions are likely to be anomalies.
- It is also useful for data analysis and visualization.





Density estimation







Dimensionality Reduction:

- Considers Latent Variables that explain observed variables.
- Decomposition techniques.
- Linear, nonlinear, parametric and non parametric.





Dimensionality Reduction

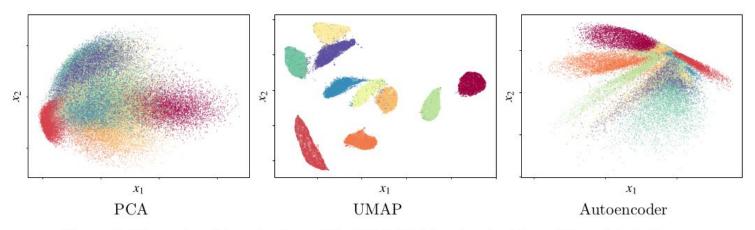


Figure 7: Dimensionality reduction of the MNIST dataset using three different techniques.





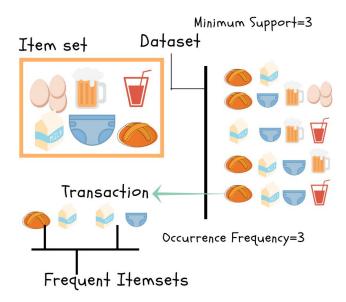
Association Rules:

- Association rule analysis has emerged as a popular tool for mining commercial databases.
- It is most often applied to binary-valued data where it is referred to as "market basket" analysis.





Association Rules:





K-means

Defines an objective function J, sometimes called a distortion measure, that computes the sum of the squares of the distances of each data point to assigned vector of centroids μ .

The goal is to find values for the clusters assignations r and the μ so as to minimize the objective function.



K-means

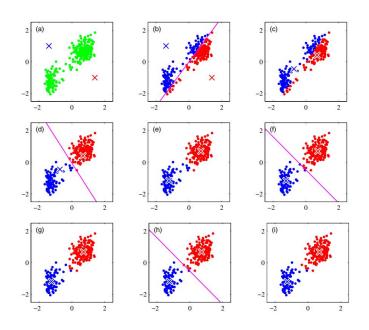
Steps:

- First we choose some initial values for the μ .
- Then in the first phase we minimize J with respect to the r, keeping the μ fixed.
- In the second phase we minimize J with respect to the μ , keeping r fixed.
- This two-stage optimization is then repeated until convergence.





K-means iteration illustration







Gaussian Mixtures

- A Gaussian mixture model (GMM) is a probabilistic model that assumes that the instances were generated from a mixture of several Gaussian distributions whose parameters are unknown.
- Each cluster can have a different ellipsoidal shape, size, density, and orientation





Gaussian Mixtures

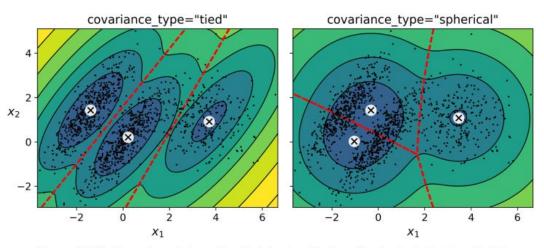


Figure 9-18. Gaussian mixtures for tied clusters (left) and spherical clusters (right)



DBSCAN (density-based spatial clustering of applications with noise)

The idea behind DBSCAN is that clusters form dense regions of data, separated by regions that are relatively empty. Also:

- It does not require the user to set the number of clusters a priori
- It can capture clusters of complex shapes.
- It can identify points that are not part of any cluster.
- Works by identifying points that are in "crowded" regions of the feature space, where many data points are close together.





Others:

- Spectral Clustering
- HDBSCAN
- Agglomerative Clustering
- BIRCH





Evaluation

 Because the dataset is completely unlabeled, deciding on whether the learned model is optimal is much more complicated than in supervised learning.





Evaluation

K-means:

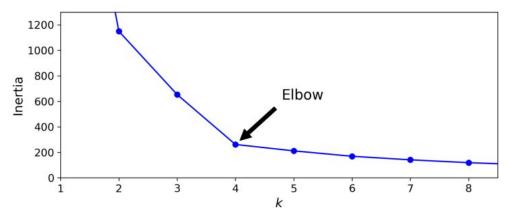
For evaluating the quality of clustering, we need to use intrinsic metrics—such as the within-cluster SSE (distortion)—to compare the performance of different k-means clusterings.





Evaluation

- The so-called elbow estimates the optimal number of clusters, k, for a given task.
- The idea behind the elbow method is to identify the value of k where the distortion begins to increase most rapidly





Evaluation

Gaussian Mixtures:

- In practice, the number of clusters is obtained with criterions from the information theory: AIC (Akaike Information Criterion) and BIC (Bayesian Information Criterion)
- Both the BIC and the AIC penalize models that have more parameters to learn (e.g., more clusters) and reward models that fit the data well. They often end up selecting the same model.



Evaluation

Others:

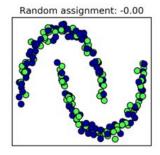
- For k-means and other methods, the silhouette score is frequently used.
- It computes the compactness of a cluster, where higher is better, with a perfect score of 1.
- However, this often don't work well in practice. While compact clusters are good, compactness doesn't allow for complex shapes.

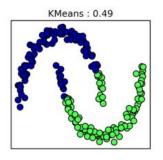


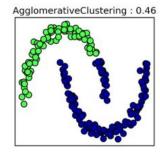


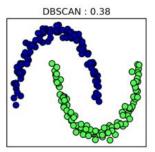
Evaluation

Silhouette scores:











Association rules attempt to construct simple rules that describe regions of high density in the special case of very high dimensional binary-valued data.

For example, suppose a data set of users and purchased items:

$\text{Item} \Rightarrow$	Bread	Butter	Milk	Fish	Beef	Ham
Customer ↓		4 9	2)			
Jack	1	1	1	0	0	0
Mary	0	1	1	0	1	0
Jane	1	1	0	0	0	0
Sayani	1	1	1	1	1	1
John	0	0	0	1	0	1
Tom	0	0	0	1	1	1
Peter	0	1	0	1	1	0



- The goal is to find joint values of the item variables that appear most frequently in the database.
- This is achieved obtaining sets of items that are closely correlated in the transaction database and with the notions of support.

The support of an itemset X is the fraction of transactions in T, of which X is a subset.

In the database example, the sets {Bread, Butter, Milk}, and {Fish, Beef, Ham} have the largest support.



 An association rule is denoted in the form X ⇒ Y, where the "⇒" is intended to give a direction to the nature of the correlation between the set of items X and Y.

For example, the rule that relates {Butter, Milk} ⇒ {Bread}



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The strength of such a rule is measured by its confidence:

The confidence of the rule $X \Rightarrow Y$ is the conditional probability that a transaction in T contains Y, given that it also contains X.



A priori algorithm

A classical algorithm for association rules is the a priori algorithm that considers a threshold for the support. In addition, the lift metric could be employed:

 The lift metric measures how much more often the antecedent X and consequent of Y a rule X ⇒ Y occur together than we would expect if they were statistically independent. It is 1 if X and Y are independent.





Hands-On

