ML5 – unsupervised learning and explainability

Machine Learning – Tools and applications for policy – Lecture 6

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ML5 – unsupervised learning and explainability

- 1. Supervised versus unsupervised learning
- 2. What can we do with unsupervised learners?
 - K-means, t-SNE, DBSCAN, Gaussian mixtures
- 3. Can we transfer knowledge?
- 4. How to open the black box and explain results?

Outline 3

ML5 – unsupervised learning and explainability

Unsupervised Learning *k*-Means clustering

t-SNE

Hierarchical Clustering

DBSCAN

Gaussian mixtures

Transfer Learning

Explainable AI



- K-means animation (Andrey Shabalin) (link)
- K-means clustering (StatQuest) (link)

Outline

ML5 – unsupervised learning and explainability

Unsupervised Learning

k-Means clustering

Hierarchical Clustering

Gaussian mixtures

Transfer Learning Explainable AI

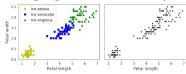


- The vast majority of data is unlabeled → enter unsupervised learning
- Flavors:
 - Clustering: group observations in similar groups for customer segmentation, recommender systems
 - Anomaly detection: what is "normal" so you can detect abnormal observations
 - Density estimation: what is the PDF of a DGP.
 Anomalies are probably in the low density areas



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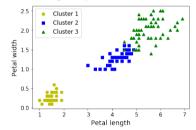
Grouping with labels is easy





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A Gaussian mixture model (covered later) can separate these clusters pretty well

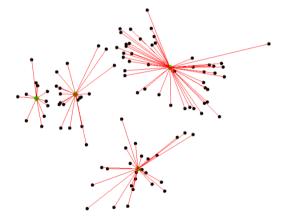




- Goal: to find subgroups or clusters by partitioning dataset into distinct groups that are maximally "different" from one another.
 - Requires a definition of what is similar/different. This is often domain specific
- Types of clustering techniques
 - *k*-means clustering: requires decision for the number of clusters *k*
 - t-SNE: non-linear PCA
 - DBSCAN: looks for "dense" areas in feature space
 - Hierarchical clustering: creates tree-like dendogram that depicts all possible clusters of size 1 to *n*.
 - Gaussian mixtures: data is generated form an unknown mixture of several Gaussian distributions with unknown parameters
 - Agglomerative clustering, BIRCH, mean-shift, affinity propagation, spectral clustering



Starting with 4 left-most points. Click the picture to continue.





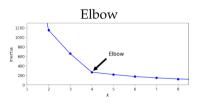
- hard clustering = assigning to a single cluster vs soft clustering = assigning a score
- Inertia is the performance metric: mean squared distance to the closest centroid

Disadvantages

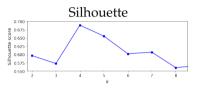
- 1. Guaranteed to converge (mostly quickly) but unclear if clustering is optimal
 - See plotting the inertia attribute. Or increase *n_init*
 - Use MiniBatchKMeans estimator in SKLearn, which reduces computation time significantly with only slight worse quality (See comparison)
- 2. Not easy/impossible to spot visually in more than 3 dimensions
- 3. Requires choosing the number of clusters.



• Plot inertia over *k* and find elbow



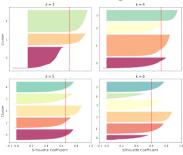
- Plot inertia over *k* and find elbow
- Plot silhouette
 - (b-a) / max(a,b) where a is the mean intra-cluster distance and b is the mean distance to the next cluster
 - Range: -1 to +1





- Plot inertia over *k* and find elbow
- Plot silhouette
 - (b-a) / max(a,b) where a is the mean intra-cluster distance and b is the mean distance to the next cluster
 - Range: -1 to +1
- Plot distribution of silhouette scores
- If you aim for similar sized groups where the observations all contribute roughly equally then k = 5 seems OK

Silhouette diagram

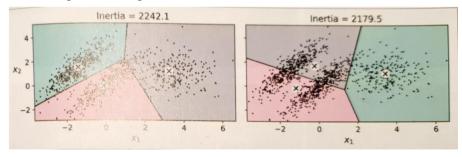




k-Means – limits

K-means does not behave well if:

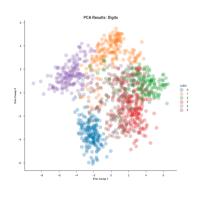
- Clusters have varying sizes
- Different densities
- Nonspherical shapes





t-Distributed Stochastic Neighbor Embedding (t-SNE)

- t-Distributed Stochastic Neighbor Embedding (t-SNE) is an unsupervised, non-linear technique primarily used for data exploration and visualizing high-dimensional data
- Difference between PCA and t-SNE: linear vs non-linear
- t-SNE calculates a similarity measure between pairs of instances in the high dimensional space and in the low dimensional space. It then tries to optimize these two similarity measures using a cost function





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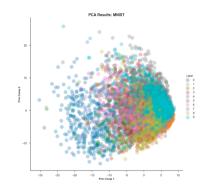




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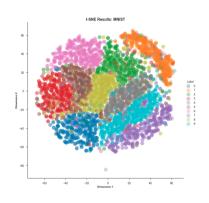




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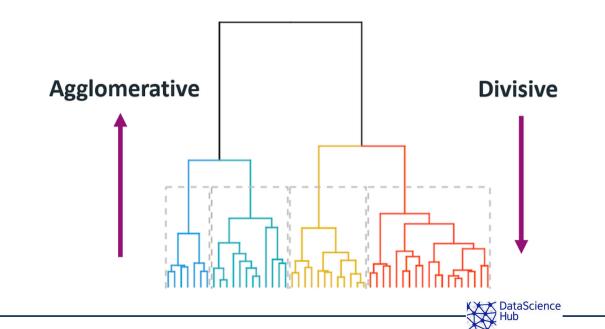
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- A drawback of k-means is the need to pick the total number of clusters a priori.
- Agglomerative Clustering: builds clusters one at a time in a bottom-up fashion.
 Initially, each of the data points is assumed to be a separate cluster. Then, the clusters that are closest to each other are merged to form a merged cluster. Repeat until all data points are merged into one single cluster. This method can help us to determine the hierarchy in which clusters were formed along with their similarity with each other. Dendrograms show this tree-like structure that shows the way in which all the clusters are formed.
- Divisive Clustering: top-down approach where all the data points are assigned to a single cluster. We split the data points into clusters based on their distance. These steps are followed until we have n clusters when n indicates the total number of data points chosen for clustering. Having a large value of n leads to having a higher number of clusters and higher computation costs. It is easier to visualize the results with the help of a dendrogram that also shows similarities between clusters that are used to divide them.



Pros and cons

Pros

- No requirement to specify the number of clusters a priori
- More intuitive as there is a hierarchy in the structure, and it is easier to interpret.

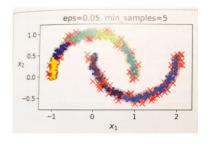
Cons

- Compute to run this algorithm is quite high. For large datasets, it can be better to look for alternative algorithms.
- Sensitive to outliers which affect the way in which clusters are formed or generated.
- Distance metric chosen used to split or merge clusters can have a significant impact



DBSCAN 16

- Looks for "dense" areas in feature space
- Has just 2 hyperparameters: ϵ and $min_samples$
- Works well if dense areas are clearly separated by sparse areas

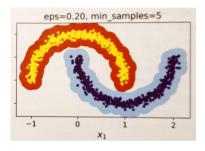


Source: Géron (2019)



DBSCAN 16

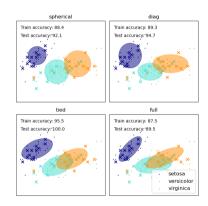
- Looks for "dense" areas in feature space
- Has just 2 hyperparameters: ϵ and $min_samples$
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- Assumes data is generated form an unknown mixture of several Gaussian distributions with unknown parameters
- Expectation Maximization (EM)
 - Similar to k-
 - Estimates not only the center but also size, shape, orientation and relative weight with soft assignments
- To reduce computational complexity adjust covariance_type: "spherical", "diag", "tied" and "full" (default)

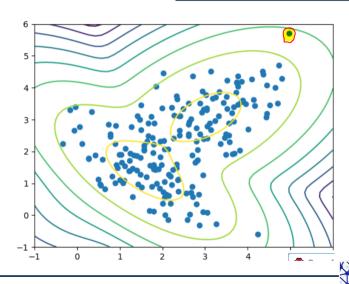


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- Number of clusters *k* is a hyperparameter (similar to K-means)
- Inertia or silhouette not well-defined
- Bayesian Information Criterion (BIC) and Akaike Information Criterion (AIC)
- Both BIC and AIC penalize models that have more parameters to learn (== more clusters) and reward models that fit the data well
- Plot BIC/AIC for an elbow plot







- Principal Components Analysis (PCA)
- Fast-MCD (minimum covariance determinant): variant of Gaussian mixture with a single distribution
- Isolation forest: Random Forest where each decision tree is grown randomly. At each node a random feature is used to split using a random threshold. Outliers tend to be split of relatively fast.
- Local Outlier Factor (LOF): compares density with its neighbors' density
- One-class SVM: can we split observations from origin? Does not scale

Outline

ML5 – unsupervised learning and explainability

Unsupervised Learning

k-Means clustering
t-SNE

Hierarchical Clustering
DBSCAN
Gaussian mixtures

Transfer Learning



- Knowledge from learning one task can be used for a different but related task
- E.g. by learning how to recognize trucks, a model could use this knowledge to recognize a passenger car





- We define a domain *D* and a task *T*, both for a source and a target.
- Domain *D* contains a feature space χ and a marginal probability distribution P(X), where $X = x_1, ..., x_n \in \chi : D = \{\chi, P(X)\}$. (Pan & Yang (2010))
- Task T consists of label space Y and an objective predictive function f(.). The task cannot be observed, but can only be learned from the training pairs x_i and y_i . Hereby, f(.) can be used to predict the label of a new instance x, so $T = \{Y, P(Y|X)\}$.
- Using the knowledge of D_S and T_S , Transfer Learning tries to improve the learning of the predictive function f(.) in D_T , where $D_S \neq D_T$, or $T_S \neq T_T$.

- Image recognition, cancer discovery and natural language processing.
- Useful in cases of data scarcity:
 - Extreme Out-of-the-Money options
 - Different stocks being affected by the same market
 - Credit data on MSMEs (Suryanto et al. 2022)
- Shows increased performance over regular learning. (He et al. 2019)

- Look at when to transfer, how to transfer, and what to transfer. (Lin and Jung 2017)
- When: in classification problems, brute-force transfer might cause a negative transfer, so do not random-guess a transfer, but consider if the performance is above chance level
- How: check if the training and test data are drawn from similar distributions, by checking the Pearson's correlation coefficient, for example
- What: apply feature selection prior to transfer, so irrelevant information isn't transferred



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Explainable AI



- XAI methods can be of substantial help in revealing the decision rules behind, for instance, a complex Random Forest (RDF) model to forecast corporate default Cascarino et al. (2021).
 - default forecasts based on the RDF model exploit more of the available information set + greater importance to indicators with non-linear/non-monotonic relationships with the outcome variable
 - RDF forecasts display stability over time in the importance assigned to key predictors
 - In stress, liquidity ratios also give a relevant contribution to predictions

Robustness



- Robustness
- Causality





- Robustness
- Causality



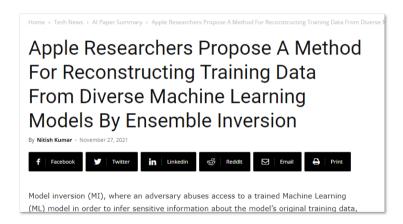


- Robustness
- Causality
- Fairness



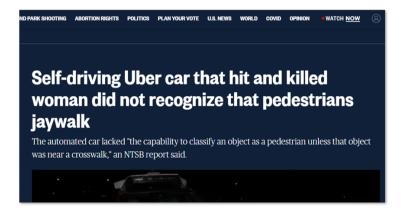


- Robustness
- Causality
- Fairness
- Privacy





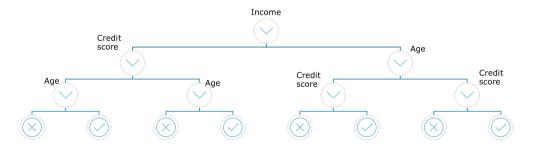
- Robustness
- Causality
- Fairness
- Privacy
- Trust





- 1. White box models
- 2. Global explainability
- 3. Local explainability

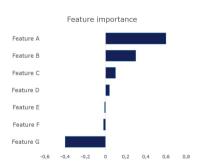
- Models which are familiar or easy to understand
 - e.g. linear regression or decision trees



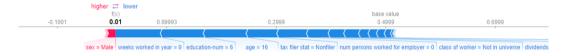


2. Global explainability

- What variables are most important in our model?
- e.g. What factors are important for predicting a higher default rate?
- Pertubation (Explain it like I'm 5 (ELI5)), Partial dependence plots



- Why is a certain decision made?
- Why does loan X have a higher default rate?
 - LIME, Shapley values (SHAP), Counterfactuals





LIME 33

LIME ((Ribeiro et al. 2016), implemented in ELI5 is an algorithm to explain predictions of black-box estimators:

- 1. Generate a fake dataset based on the data analysed
- 2. Use black-box estimator to get target values in a generated dataset (e.g. class probabilities)
- 3. Train a new white-box estimator, using generated data + labels as training data
- 4. Explain the original example through weights of this white-box estimator instead
- 5. Prediction quality of a white-box classifer shows how well it approximates the black-box classifier. If the quality is low then don't trust explanation



- Pretend the AI model is a cooperative game
 - Payout is the prediction accuracy
 - Players are the parameters
- Calculate a fair way to distribute the payout over the players

$$\phi(\nu) = \frac{1}{|N|} \sum_{S \subseteq N\{i\}} \frac{n-1}{|S|}^{-1} [v(S \cup \{i\}) - \nu(S)]$$

1. Relative importance player i 2. Number of players 3. Each coalition without player i 4. Number of coalitions of this size without player i 5. Additional payout of including i to the coalition







Summary 36

In this lecture we covered:

- 1. Some unsupervised learners
 - k-Means clustering, t-SNE, DBSCAN, Gaussian mixtures
- 2. The idea that you can transfer knowledge from one application to another
- 3. A first look at explainable AI
 - white box, global and local explainability



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