Clustering and The Expectation-Maximization Algorithm Unsupervised Learning

Marek Petrik

3/7

Some of the figures in this presentation are taken from "An Introduction to Statistical Learning, with applications in R" (Springer, 2013) with permission from the authors: G. James, D. Witten, T. Hastie and R. Tibshirani

Learning Methods

1. **Supervised Learning**: Learning a function *f*:

 $Y = f(X) + \epsilon$

- 1.1 Regression
- 1.2 Classification
- 2. Unsupervised learning: Discover interesting properties of data (no labels)

 X_1, X_2, \ldots

- 2.1 Dimensionality reduction or embedding
- 2.2 Clustering

Principal Components Analysis

- Reduce dimensionality
- Start with features $X_1 \dots X_n$
- Construct *fewer* features $Z_1 \dots Z_M$

$$Z_1 = \phi_{11}X_1 + \phi_{21}X_2 + \ldots + \phi_{p1}X_p$$

• Weights are usually normalized (using ℓ_2 norm)

$$\sum_{j=1}^{p} \phi_{j1}^2 = 1$$

Data has greatest variance along Z₁

1st Principal Component



► 1st Principal Component: Direction with the largest variance

$$Z_1 = 0.839 \times (pop - \overline{pop}) + 0.544 \times (ad - \overline{ad})$$

More Unsupervised Learning: Discovering Structure of Data

- 1. K-Means Clustering
- 2. Hierarchical Clustering
- 3. Expectation-Maximization Method (Not Covered in ISL, see ESL 8.5)

Clustering

Simplify data in a different way than PCA.

PCA finds a low-dimensional representation of data

 Clustering finds homogeneous subgroups among the observations

Clustering: Assumptions and Goals

- Exists a method for measuring similarity between data points
- Some points are more similar than others

 Discover latent patterns that exist but may not be observed/observable

Clustering: Assumptions and Goals

- Exists a method for measuring similarity between data points
- Some points are more similar than others

Want to identify similarity patterns

- 1. Discover the different types of disease
- 2. Market segmentation: Types of users that visit a website
- 3. Discover movie or book genres
- 4. Discover types of topics in documents
- Discover latent patterns that exist but may not be observed/observable

Clustering Algorithms

- K-Means: simple and effective
- Hierarchical clustering: Many complex clusters
- Many other clustering methods, most heuristics
- EM: General algorithm for dealing with latent variables by maximizing likelihood

K-Means Clustering

- Cluster data into complete and non-overlapping sets
- Example:



- k-th cluster: C_k
- *i*-th observation in cluster $k: i \in C_k$

- k-th cluster: C_k
- *i*-th observation in cluster $k: i \in C_k$
- ► Find clusters that are homogeneous: W(C_k) homogeneity of clusters

$$\min_{C_1,\dots,C_K} \sum_{k=1}^K W(C_k)$$

• k-th cluster: C_k

- *i*-th observation in cluster $k: i \in C_k$
- ► Find clusters that are homogeneous: *W*(*C_k*) homogeneity of clusters

$$\min_{C_1,\dots,C_K} \sum_{k=1}^K W(C_k)$$

Define homogeneity as in-cluster variance

$$\min_{C_1,\dots,C_K} \sum_{k=1}^K \left(\frac{1}{|C_k|} \sum_{i,i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2 \right)$$

• k-th cluster: C_k

- *i*-th observation in cluster $k: i \in C_k$
- ► Find clusters that are homogeneous: W(C_k) homogeneity of clusters

$$\min_{C_1,\dots,C_K} \sum_{k=1}^K W(C_k)$$

Define homogeneity as in-cluster variance

$$\min_{C_1,\dots,C_K} \sum_{k=1}^K \left(\frac{1}{|C_k|} \sum_{i,i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2 \right)$$

This is an NP hard problem

K-Means Algorithm

Heuristic solution to the minimization problem

- 1. Randomly assign cluster numbers to observations
- 2. Iterate while clusters change
 - 2.1 For each cluster, compute the centroid
 - 2.2 Assign each observation to the closest cluster

Note that:

$$\frac{1}{|C_k|} \sum_{i,i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2 = 2 \sum_{i,i' \in C_k} \sum_{j=1}^p (x_{ij} - \bar{x}_{kj})^2$$

K-Means Illustration



Properties of K-Means

- Local minimum: Does not necessarily find the optimal solution
- Multiple runs can result in different solutions
- Choose the result of the run with minimal objective
- Cluster labels do not matter

Multiple Runs of K-Means



235.8



310.9



Hierarchical Clustering

- Multiple levels of similarity needed in complex domains
- Build a similarity tree



Dendrogram: Similarity Tree



Hierarchical Clustering Algorithm

- 1. Begin with n observations and compute $\binom{n}{2}$ dissimilarity measures
- 2. For i = n, n 1, ..., 2
 - 2.1 Fuse 2 most similar clusters
 - 2.2 Update i 1 dissimilarities

Hierarchical Clustering Algorithm: Illustration



Dissimilarity Measure: Linkage

- 1. Complete
- 2. Single
- 3. Average
- 4. Centroid

Impact of Dissimilarity Measure



Clustering in Practice

- Fraught with problems: no clear measure of quality (like MSE)
- ▶ How to choose *k*? Problem dependent
- Standardize features, center them?
- What dissimilarity to use?

Clustering in Practice

- Fraught with problems: no clear measure of quality (like MSE)
- ► How to choose *k*? Problem dependent
- Standardize features, center them?
- What dissimilarity to use?
- Careful over-explaining clustering results: source: http://miriamposner.com

List of Topics

Meta-Humanist? >

- 1. digitalhumanities humanist org http lists interface listmember php list computing
- 2. humanities digital research arts department scholars sciences projects director academic
- 3. digital http gmail dho dublin day project susan subject www
- 4. humanist www kessler ubiquity jascha org ucla subject professor acm
- 5. text texts project archive images edition editions tools textual editing -
- 6. time people back thing things willard good point mind read
- 7. social art university systems networks computing visual uk symposium analysis
- 8. text markup humanist tool xml subject don schmidt wendell desmond
- 9. uk ac london kcl www king http college research centre Digital archaeology?
- 10. humanities digital archaeology words university spatial work gis session november
- 11. lachance utoronto tis chass http ca quote island francois dec

Textual editing?

Expectation-Maximization

- Maximum likelihood approach to clustering
- General method for dealing with latent features / labels
- Especially useful with generative models
- A heuristic method used to solve complex optimization problems
- Generalization of the idea: Minorization-Maximization
- Gentle introduction: https://www.cs.utah.edu/ ~piyush/teaching/EM_algorithm.pdf

Recall LDA

 Generative model: capture probability of predictors for each label



Predict:

 Generative model: capture probability of predictors for each label



Predict:

1. $\Pr[\text{balance} \mid \text{default} = \text{yes}] \text{ and } \Pr[\text{default} = \text{yes}]$

 Generative model: capture probability of predictors for each label



Predict:

- 1. $\Pr[\text{balance} \mid \text{default} = yes] \text{ and } \Pr[\text{default} = yes]$
- 2. $\Pr[\text{balance} \mid \text{default} = no] \text{ and } \Pr[\text{default} = no]$

 Generative model: capture probability of predictors for each label



Predict:

- 1. $\Pr[\text{balance} \mid \text{default} = yes] \text{ and } \Pr[\text{default} = yes]$
- 2. $\Pr[\text{balance} \mid \text{default} = no] \text{ and } \Pr[\text{default} = no]$
- Classes are normal: Pr[balance | default = yes]

LDA vs Logistic Regression

Logistic regressions:

 $\Pr[\mathsf{default} = \operatorname{yes} | \mathsf{balance}]$

Linear discriminant analysis:

 $\Pr[\text{balance} \mid \text{default} = \text{yes}] \text{ and } \Pr[\text{default} = \text{yes}]$ $\Pr[\text{balance} \mid \text{default} = \text{no}] \text{ and } \Pr[\text{default} = \text{no}]$

LDA with 1 Feature

• Classes are normal and class probabilities π_k are scalars

$$f_k(x) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left(-\frac{1}{2\sigma^2}(x-\mu_k)^2\right)$$

• Key Assumption: Class variances σ_k^2 are the same.



EM For LDA

- Labels are missing, guess them
- Find the most likely model <u>and</u> latent observations:

$$\max_{\text{model}} \log \ell(\text{model}) = \max_{\substack{\text{model}\\ \text{latent}}} \log \sum_{\text{latent}} \Pr[\text{data}, \text{latent} \mid \text{model}] =$$

EM For LDA

- Labels are missing, guess them
- Find the most likely model <u>and</u> latent observations:

$$\begin{split} \max_{model} \log \ell(model) &= \max_{\substack{model \\ latent}} \log \sum_{latent} \Pr[data, latent \mid model] = \\ &= \max_{\substack{model \\ latent}} \log \sum_{latent} \Pr[data \mid latent, model] \Pr[latent \mid model] \end{split}$$

EM For LDA

- Labels are missing, guess them
- Find the most likely model <u>and</u> latent observations:

$$\begin{split} \max_{\text{model}} \log \ell(\text{model}) &= \max_{\substack{\text{model} \\ \text{latent}}} \log \sum_{\text{latent}} \Pr[\text{data}, \text{latent} \mid \text{model}] = \\ &= \max_{\substack{\text{model} \\ \text{latent}}} \log \sum_{\text{latent}} \Pr[\text{data} \mid \text{latent}, \text{model}] \Pr[\text{latent} \mid \text{model}] \end{split}$$

• Difficult and non-convex optimization problem ($\log \sum$)

EM Derivation

- Iteratively approximate and optimize the log-likelihood function
 - 1. Construct a concave lower bound
 - 2. Maximize the lower bound
 - 3. Repeat
- Notation:
 - Model: θ
 - Data: x
 - Latent variables: z

$$\max_{\theta, z} \log \ell(\theta, z) = \max_{\theta, z} \log \Pr[x \mid \theta] =$$
$$= \max_{\theta, z} \log \sum_{z} \Pr[x \mid z, \theta] \Pr[z \mid \theta]$$

EM Derivation

- Suppose we have an estimate of the model θ_n
- How to compute θ_{n+1} that improves on it?

$$\begin{aligned} \theta_{n+1}, z_{n+1} &= \\ \arg\max_{\theta, z} \log \sum_{z} \Pr[x, z \mid \theta] = \arg\max_{\theta, z} \log \sum_{z} \Pr[z \mid \theta] \Pr[z \mid x, \theta] = \\ &= \arg\max_{\theta, z} \log \sum_{z} \Pr[z \mid \theta] \Pr[z \mid x, \theta] \frac{\Pr[z \mid x, \theta_n]}{\Pr[z \mid x, \theta_n]} = \\ &= \arg\max_{\theta, z} \log \sum_{z} \Pr[z \mid x, \theta_n] \frac{\Pr[z \mid \theta] \Pr[z \mid x, \theta]}{\Pr[z \mid x, \theta_n]} \leq \\ &\stackrel{\text{jensen's}}{\leq} \arg\max_{\theta, z} \sum_{z} \Pr[z \mid x, \theta_n] \log \frac{\Pr[z \mid \theta] \Pr[z \mid x, \theta]}{\Pr[z \mid x, \theta_n]} = \\ &= \arg\max_{\theta, z} \sum_{z} \Pr[z \mid x, \theta_n] \log \frac{\Pr[z \mid \theta] \Pr[z \mid x, \theta]}{\Pr[z \mid x, \theta_n]} = \end{aligned}$$

EM Algorithm

- 1. **E Step**: Estimate $\Pr[z \mid x, \theta_n]$ for all values of z. (Construct the lower bound)
- 2. M-Step: Maximize the lower bound:

$$\theta_{n+1} = \arg \max_{\theta} \sum_{z} \Pr[z \mid x, \theta_n] \log \Pr[x, z \mid \theta]$$

This can be solved using traditional MLE methods with weighted samples

EM for Mixture of Gaussians

Rough sketch

- 1. Randomly assign cluster weights to observations
- 2. Iterate while clusters change
 - 2.1 For each cluster, compute the centroid based on observation **weights** of observations
 - 2.2 Assign each observation new cluster **weights** based on the distances from centroids

Other Applications of EM

- Very powerful and general idea!
- Training with missing data for many model types
- Hidden variables in Bayesian nets
- Identifying confounding variables
- Solving difficult (complex) optimization problem: MM