

CS5785 - Applied Machine Learning - Lecture 13

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October 11, 2018

1 Recap - Unsupervised learning/Dimensionality Reduction

We can use dimensionality reduction when we have $X \in \mathbb{R}^{n \times p}$ that is high dimensional data and want to transform it to a matrix $Z \in \mathbb{R}^{n \times q}$ where $q < p$ that preserves the most information.

We can use PCA:

$$z = e(x)$$

$$\hat{x} = d(z)$$

$$e(x) = Ax$$

$$d(Z) = A^T Z$$

$$A^T A = I$$

The best A is $A = Vq$: first q columns from V in SVD of X

2 Multidimensional Scaling

We have the following dissimilarity measures and their usual uses in practice:

Dissimilarity measures	Usual Uses
Euclidean Distance	Vectors
Chi-squared Distance	Distributions
Cosine Similarity	Distance between bag of words, patterns
Edit Distance	Distance between strings

Multidimensional Scaling process: we start with the distance/dissimilarity matrix $D \in \mathbb{R}^{n \times n}$.

What vectors $D \in \mathbb{R}^{n \times p}$ recover D in the closest way via $D_{ij} \approx \|X_i - X_j\|_2^2$ i.e. X is an embedding of the n abstract data points in p dimensions that maintains the dissimilarities as vector distances.

If indeed $D_{ij} = \|X_i - X_j\|_2^2$ then

$$D_{ij} = \|X_i\|_2^2 + \|X_j\|_2^2 - 2X_i^T X_j \quad D_{ij} = T_{ii} + T_{jj} - 2T_{ij}$$

such that $T = XX^T \in \mathbb{R}^{n \times n}$

$$T = \frac{1}{2} \left(D - \frac{Dee^T}{n} - \frac{ee^T D}{n} + \frac{e^T D e}{n^2} \right)$$

In MDS, we eigendecompose T to get X such that $T = XX^T$, $T = UVU^T$ and $X = U_p V_p^{1/2}$, a reconstruction up to rotation and translation.

Fact: if D comes from euclidean distances then MDS is just PCA.

3 Clustering

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The goal here is to assign data points to finitely many clusters. One natural perspective would be to find clusters in data. Another perspective is dimensionality reduction where $e(x) = 1, \dots, k$

Encoder = cluster membership

Decoder = mean of that cluster = $\mu_{e(x)}$, $e(x) \in 1, \dots, k$ and $d(z) = \hat{\mu}_z$

3.1 K-means Algorithm

The goal with k-means clustering is to assign data points to clusters such that the within cluster distance is small:

Let $c(i) = 1, \dots, K$ indicate the assignment of point i .

The quality of c is defined in terms of the within-cluster distance:

$$W(c) = \sum_{j=1}^k \sum_{i:c(i)=j} \|X_i - \mu_j\|_2^2 \text{ where:}$$

$$\mu_j = \frac{1}{n_j} \sum_{i:c(i)=j} X_i$$

$$n_j = \sum_{i:c(i)=j} 1$$

$$W(c) = \frac{1}{2} \sum_{j=1}^k \frac{1}{n_k} \sum_{i:c(i)=j} \sum_{i':c(i')=j} \|X_i - X_{i'}\|_2^2$$

What is the best C (min $W(c)$)? How many C 's are there? The number of ways of assigning n things to k buckets is far too big for us to be able to compute so we use a greedy algorithm (k-means):

- 1) Start with some C_0
- 2) For $t = 1, 2, \dots$:

1. Compute the cluster means for C_{t-1}

$$\mu_j^{(t-1)} = \frac{1}{n_j^{(t-1)}} \sum_{i: C_{t-1}(i)=j} X_i \text{ for every } j = 1, \dots, k$$

2. Reassign n data points to the closest mean:

$$C_t(i) = \operatorname{argmin}_j \|x_i - \mu_j^{(t-1)}\|_2^2 \text{ for every } j = 1, \dots, k$$

Observation 1: different initialization lead to different solutions. Try different (random) starting points. Pick the final clustering with smallest $W(c)$.

Observation 2: k-means will always converge in finite number of steps.

- Might not be to global option.
- Might take a long time so often we terminate after a fixed number of steps.
- Do not choose k by cross-validation.
- Increase k to decrease $\min_c W(c)$
- There do exist information theoretical ways to choose k , and there is no "right" or "wrong" way.

4 Soft Clustering

Hard clustering: k-means assigns each point to exactly 1 cluster

Soft clustering: sometimes it's not clear that there's a single cluster that a point belongs to, instead we want to partially assign points to clusters.

Suppose we have a binomial distribution as follows: $X_0 \sim N(\mu_0, \sigma^2)$

$Y \sim \text{Bernoulli}(\pi)$

$$X = (1 - Y)X_0 + YX_1 = \begin{cases} X_0 & Y = 0 \\ X_1 & Y = 1 \end{cases} \quad (X_1 Y = y) \sim N(\mu_y, \sigma_y^2)$$

Approach : find $\theta = (\pi, \mu_0, \sigma_0^2, \mu_1, \sigma_1^2)$ that maximize the likelihood of the data (MLE)