

STATISTICAL PATTERN RECOGNITION

ASSIGNMENT 6

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Abstract

Keywords.

1 LDF Based on Perceptron Learning Rule

Recall that a perceptron learns a linear classifier with weight vector W . It predicts:

$$\hat{y} = \text{sign}(W^T x_t)$$

assuming here that $\hat{y} \in \{+1, -1\}$. When the perceptron makes a mistake, it updates the weights using the formula:

$$W^{t+1} = W^t + y_t x_t$$

Imagine that we have $x_t \in I^2$, and we encounter the following data points:

X[1]	X[2]	y
1	1	1
2	-1	-1
-3	-1	-1
-3	1	1

- Starting with $W = [0 \ 0]^T$, use the Perceptron Algorithm to learn on the data points in the order from top to bottom. Show the Perceptron's linear decision boundary after observing each data point. Be sure to show which side is classified as positive.
- Does our learned perceptron maximize the margin between the training data and the decision boundary? If not, draw the maximum-margin decision boundary.

Solution

We'll start by augmenting the feature vectors given in the table. We'll add a new feature x_0 (bias term) and set it to 1 for class 1, and -1 for class -1 . This results in the following data points:

Y[0]	Y[1]	Y[2]	Class
1	1	1	1
-1	-2	1	-1
-1	3	1	-1
1	-3	1	1

we'll then augment the weight vector for a consistent definition of the decision boundary. This yields in

$$W = \begin{bmatrix} 0 \\ 0 \end{bmatrix} \rightarrow a^0 = \begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix}$$

Note that according to the Perceptron Learning Rule, we're looking for a line to meet the following constraint:

$$z_i = a^T y_i > 0 \quad \forall y_i \quad (1.1)$$

weights $a = \begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix}$ cannot correctly classify the first point since $z_1^0 = 0$ which is a sign of misclassification. We can correctly classify this point conducting a *weight update* using Perceptron Learning Rule:

$$a^1 = \begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix} + \begin{bmatrix} 1 \\ 1 \\ 1 \end{bmatrix} = \begin{bmatrix} 1 \\ 1 \\ 1 \end{bmatrix}$$

The resulting linear boundary will incorrectly classify the second point:

$$z_2^1 = [1 \ 1 \ 1] \cdot \begin{bmatrix} -1 \\ -2 \\ 1 \end{bmatrix} \leq 0$$

Thus we have to update the weights using Perceptron Rule again:

$$a^2 = \begin{bmatrix} 1 \\ 1 \\ 1 \end{bmatrix} + \begin{bmatrix} -1 \\ -2 \\ 1 \end{bmatrix} = \begin{bmatrix} 0 \\ -1 \\ 2 \end{bmatrix}$$

The resulting line will poorly perform on the third data point:

$$z_3^2 = [-1 \ 3 \ 1] \cdot \begin{bmatrix} 0 \\ -1 \\ 2 \end{bmatrix} \leq 0$$

Updated weights for the third iteration is provided below:

$$a^3 = \begin{bmatrix} 0 \\ -1 \\ 2 \end{bmatrix} + \begin{bmatrix} -1 \\ 3 \\ 1 \end{bmatrix} = \begin{bmatrix} -1 \\ 2 \\ 3 \end{bmatrix}$$

following the same pattern for the final data point results in the final weight update:

$$a^4 = \begin{bmatrix} 0 \\ -1 \\ 2 \end{bmatrix} + \begin{bmatrix} 1 \\ -3 \\ 1 \end{bmatrix} = \begin{bmatrix} 1 \\ -4 \\ 3 \end{bmatrix}$$

Corresponding decision boundaries for each step is provided in the figure 1.1

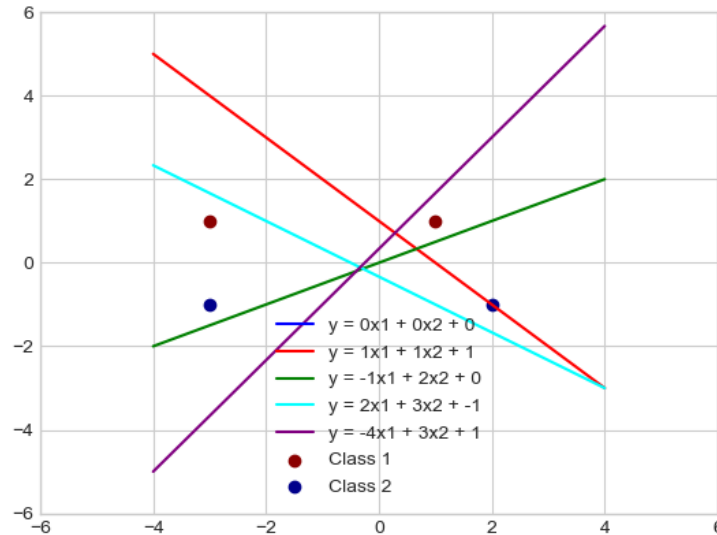


Figure 1.1: Illustration of Perceptron decision boundaries and class points.

The learned Perceptron does not maximize the margin between the training data and the decision boundary. The max-margin linear discriminator is $y = 0$.

2 Perceptron in Practice

Start with the truth tables for boolean *AND* and *OR*. Your goal is to write a small program to train a *two-input one-output* Perceptron using this data. Select some random initial weights in the interval $(0, 1)$. Use a learning parameter 0.1. The stopping condition is 50 iterations or *no change in weights from one epoch to the next*, whichever comes first.

- (a) Encode $FALSE = 0$ and $TRUE = 1$ for the input values. Remember that the output of a Perceptron, by definition is $+1$ or -1 .
- (b) Encode $FALSE = -1$ and $TRUE = +1$ for the input values. Remember that the output of a Perceptron, by definition is $+1$ or -1 .

For each case, please show the evolution in the change of weights by arranging them in a nice tabular format. Remember to include the weight w_0 as a part of the training process and set your x_0 to $+1$ all the time. For each case, please plot the separating hyperplane in the $x_1 - x_2$ plane. Submit your code, result of your run and your comments.

3 Support Vector Machine

Consider a dataset with 2 points in 1D:

$$X_1 = 0, Y_1 = -1 \quad X_2 = \sqrt{2}, Y_2 = 1$$

Consider mapping each point to 3D using the feature vector $\phi(x) = [1, \sqrt{2}x, x^2]^T$. Here, the max margin classifier has the form:

$$\hat{W}, \hat{W}_0 = \operatorname{argmin} ||W||^2 \tag{3.1}$$

so that

$$\begin{aligned} y_1(W^T \phi_1(x) + W_0) &\geq 1 \\ y_2(W^T \phi_2(x) + W_0) &\geq 1 \end{aligned}$$

- (a) Write down a vector that is parallel to the optimal vector \hat{W} . Hint: \hat{W} is perpendicular to the decision boundary between the two points in the 3D feature space.
- (b) What is the value of the margin that is achieved by this \hat{W} ? Hint: The margin is the distance from each support vector to the decision boundary. Think about the geometry of the points in feature space, and the vector between them.
- (c) Solve for \hat{W} , using the fact that the margin is equal to $\frac{1}{||\hat{W}||}$.
- (d) Solve for \hat{W}_0 using your value for \hat{W} and (3.1). Hint: The points will be on the decision boundary, so the inequalities will be tight. A *Tight Inequality* is an equality that is as strict as possible.

Solution

- (a) The initial 1-dimensional dataset is $D = (0 \ \sqrt{2})$. Applying the transformation yields the following points in 3D:

$$D' = \begin{bmatrix} 1 & 1 \\ 0 & 2 \\ 0 & 2 \end{bmatrix}$$

In order to find the *SVM* decision boundary, we'll use the following formula which represents an optimization problem with respect to the constraints given in the problem description:

$$L_D(\alpha) = \sum_{i=1}^2 \alpha_i - \frac{1}{2} \sum_{i=1}^2 \sum_{j=1}^2 \alpha_i \alpha_j z_i z_j x_i \cdot x_j \quad (3.2)$$

This can be written as

$$L_D(\alpha) = \alpha_1 + \alpha_2 - \frac{1}{2} \left(\alpha_1 \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix} - \alpha_2 \begin{bmatrix} 1 \\ 2 \\ 2 \end{bmatrix} \right) \cdot \left(\alpha_1 \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix} - \alpha_2 \begin{bmatrix} 1 \\ 2 \\ 2 \end{bmatrix} \right)$$

simplifying the results yields in

$$L_D(\alpha) = \frac{-9}{2} \alpha_2^2 - \frac{1}{2} \alpha_1^2 + \alpha_1 \alpha_2 + \alpha_1 + \alpha_2 \quad (3.3)$$

To solve (3.3), we should find the derivatives with respect to α_1 and α_2 .

$$\frac{\partial L_D(\alpha)}{\partial \alpha_1} = -\alpha_1 + \alpha_2 + 1 = 0$$

$$\frac{\partial L_D(\alpha)}{\partial \alpha_2} = -9\alpha_2 + \alpha_1 + 1 = 0$$

Thus $\alpha_1 = \frac{5}{4}$ and $\alpha_2 = \frac{1}{4}$. Weights can be obtained using

$$W = \sum_{i=1}^2 z_i x_i \alpha_i \quad (3.4)$$

Thus

$$W = \frac{5}{4} \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix} - \frac{1}{4} \begin{bmatrix} 1 \\ 2 \\ 2 \end{bmatrix} = \begin{bmatrix} 1 \\ -1/2 \\ -1/2 \end{bmatrix}$$

An obvious vector parallel with the weight vector is the normalized vector of W :

$$V = \begin{bmatrix} \frac{1}{\sqrt{1+\frac{1}{4}+\frac{1}{4}}} \\ \frac{-1/2}{\sqrt{1+\frac{1}{4}+\frac{1}{4}}} \\ \frac{-1/2}{\sqrt{1+\frac{1}{4}+\frac{1}{4}}} \end{bmatrix} = \begin{bmatrix} 0.81 \\ -0.40 \\ -0.40 \end{bmatrix}$$

(b)

4 Hierarchical Clustering

Let $X = \{x_i, i = 1, 2, 3, 4, 5\}$, with $x_1 = [1, 1]^T$, $x_2 = [2, 1]^T$, $x_3 = [5, 4]^T$, $x_4 = [6, 5]^T$ and $x_5 = [6.5, 6]^T$. The pattern matrix of X is

$$D(X) = \begin{bmatrix} 1 & 1 \\ 2 & 1 \\ 5 & 4 \\ 6 & 5 \\ 6.5 & 6 \end{bmatrix}$$

Use the following methods to cluster the pattern matrix and draw the dendrogram for hierarchical models:

- (a) Single Linkage
- (b) Complete Linkage
- (c) Average Linkage
- (d) Mean Method

Solution

(a) Note that in we've used *Manhattan* distance in all sections for simplicity. We'll fill our distance table with proper distances:

	x_1	x_2	x_3	x_4	x_5
x_1	0	1	7	9	10.5
x_2	1	0	6	8	9.5
x_3	7	6	0	2	3.5
x_4	9	8	2	0	1.5
x_5	10.5	9.5	3.5	1.5	0

The *Single Linkage* method initializes each of the points in the dataset to a cluster respectively (a bottom-up approach).

Step 1 ($d = 1$)

At the very first beginning, the lowest distance between two points is selected from the table. The cluster $\{x_1, x_2\}$ is made in this step.

Step 2 ($d = 1.5$)

	$\{x_1, x_2\}$	x_3	x_4	x_5
$\{x_1, x_2\}$	0	6	8	9.5
x_3	6	0	2	3.5
x_4	8	2	0	1.5
x_5	9.5	3.5	1.5	0

In this step, x_4 and x_5 are merged together with a distance of $d = 1.5$:

	$\{x_1, x_2\}$	x_3	$\{x_4, x_5\}$
$\{x_1, x_2\}$	0	6	8
x_3	6	0	2
$\{x_4, x_5\}$	9.5	2	0

Step 3 ($d = 2$)

In step 3, x_3 , x_4 and x_5 are being merged together.

	$\{x_1, x_2\}$	$\{x_3, x_4, x_5\}$
$\{x_1, x_2\}$	0	6
$\{x_3, x_4, x_5\}$	6	0

Step 4 ($d = 6$)

In the final step, 2 clusters are merged together with $d = 6$. *Dendrogram* for this method is given in figure (4.1). (b) The difference between *Average Linkage* and *Single Linkage*

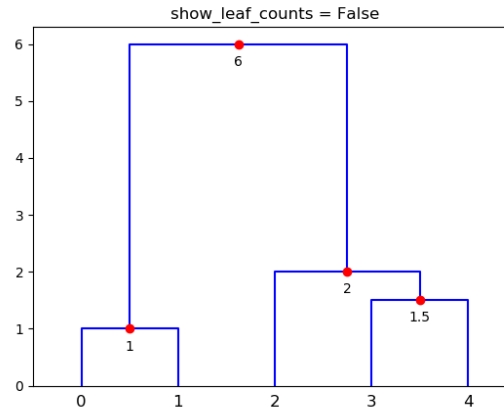


Figure 4.1: Dendrogram illustration of hierarchical clustering with *single link* method.

method is **when** the clustering happens. In the *Complete Linkage* method, the average distance between two clusters are considered while merging two clusters. Following the same pattern for this section yields in the following results:

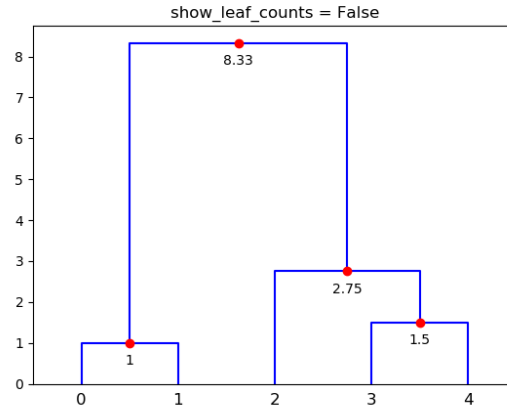


Figure 4.2: Dendrogram illustration of hierarchical clustering with *average link* method.

(c) The difference between *Complete Linkage* and *Single Linkage* method is **when** the clustering happens. In the *Complete Linkage* method, the maximum distance between two clusters are considered while merging two clusters. Following the same pattern for this section yields in the following results: The code for these sections is provided in the *src*

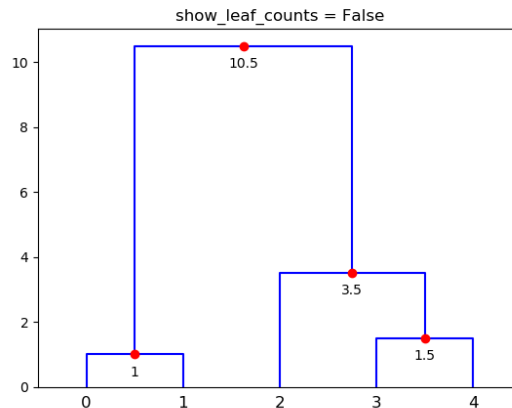


Figure 4.3: Dendrogram illustration of hierarchical clustering with *complete link* method.

folder. (d) In this case, the metric will be changed to *Euclidean*. The mean of each cluster is considered in each step and the distance of two clusters will be equal to the distance of *centroids* of two clusters. The resulting dendrogram is given in figure 4.4.

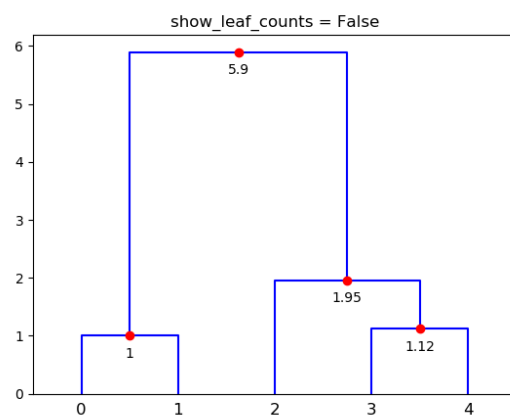


Figure 4.4: Dendrogram illustration of hierarchical clustering with *mean* method.