3. PERCEPTRON

Consider the problem in supervised learning. In this case, the data is given as a set of pairs

$$D = \{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(n)}, y^{(n)})\}\$$

 $x^{(i)} = \text{input} = \text{independent variable} = \text{predictor} = \text{attribute}$

 $y^{(i)}$ = output = dependent variable = response = target

Goal: we want to learn the underlying relationship (hypothesis) between x and y, so that next time we are given an instance of an input x, we can predict its corresponding output y.

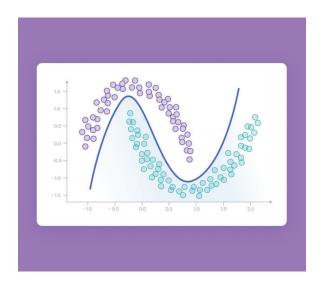
• Typically, $x \in \mathbb{R}^d$ is a vector of d components

$$x = \begin{bmatrix} x_1 \\ x_2 \\ \dots \\ x_d \end{bmatrix}$$

More precisely, x might be a customer, patient, image, text, song, etc., and we need to represent it numerically

$$\varphi(x) = \text{ feature representation } \in \mathbb{R}^d$$

• In this section, we will study a <u>binary classification</u> problem where we are trying to separate two classes of data instances (data points, examples, samples). We assume $y \in \{+1, -1\}$, denoting the labels for data instances.



 $\rm https://www.v7labs.com/blog/semi-supervised-learning-guide$

• To create a learning algorithm, we need a hypothesis class \mathcal{H} which is a set of all possible functions (hypotheses) $x \mapsto y$. More precisely,

$$y = h(x; \theta)$$

where θ denotes parameters.

$$x \to h \to y$$

The goal is to find h that agrees with the given data in D and we hope that h will perform well in the future.

• Next, we define what makes one hypothesis better than another. We define the loss (cost, error) function for a single data instance $(x^{(i)}, y^{(i)})$ that measures the error of predicting the class label $\hat{y}^{(i)} := h(x^{(i)})$ when the actual true class label is $y^{(i)}$. We denote the loss by

$$L(\hat{y}^{(i)}, y^{(i)})$$

where

$$\hat{y}^{(i)} \in \{+1, -1\}$$
 predicted label $y^{(i)} \in \{+1, -1\}$ actual true label

Objective: have a small loss on the \underline{new} data

Proxy: have a small loss on the given training data $D = \{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(n)}, y^{(n)})\}$ We minimize the loss on the training data

$$\mathcal{E}_{train}(h) = \frac{1}{n} \sum_{i=1}^{n} L(h(x^{(i)}), y^{(i)})$$
 training set error

and we hope that the loss on the test data (additional data that the algorithm has not seen) $\{(x^{(n+1)}, y^{(n+1)}), (x^{(n+2)}, y^{(n+2)}), \dots, (x^{(n+n')}, y^{(n+n')})\}$ will also be small

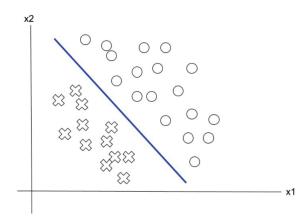
$$\mathcal{E}_{test}(h) = \frac{1}{n'} \sum_{i=n+1}^{n+n'} L(h(x^{(i)}), y^{(i)})$$
 test error

• How do we come up with a learning algorithm?

$$D \to \boxed{\text{Learning Algorithm}(\mathcal{H})} \to h$$

- be a clever human
- use optimization

• In this section, we consider *linear classifiers*, i.e., we assume that \mathcal{H} is a class of all possible linear separators.



https://automaticad dison.com/linear-separability-and-the-xor-problem/

In particular, we consider a hypothesis of the form

$$h(x; w, b) = sign(w^{T}x + b) = \begin{cases} +1, & w^{T}x + b > 0 \\ -1, & w^{T}x + b < 0 \end{cases}$$

where $w \in \mathbb{R}^d$ and $b \in \mathbb{R}$.

Recall

$$w^{T}x = [w_{1} \ w_{2} \ \dots \ w_{d}] \begin{bmatrix} x_{1} \\ x_{2} \\ \dots \\ x_{d} \end{bmatrix} = w_{1}x_{1} + w_{2}x_{2} + \dots + w_{d}x_{d} = \sum_{j=1}^{d} w_{j}x_{j}$$

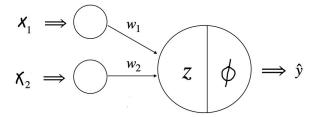
The values of vector w are called weights and b is called a bias.

- The algorithm computes the weighted sum of the inputs and if this weighted sum exceeds some threshold (specifically, -b), then the predicted label is positive, and if the weighted sum is less than this threshold, the predicted label is negative.
- For simplicity, from now on, we consider the case d=2 with two inputs $(x_1, x_2) \in \mathbb{R}^2$. We can also visualize the above hypothesis as a single neuron.

Given an input (x_1, x_2) , the predicted output \hat{y} is found using two steps:

1. the quantity z (pre-activation) is calculated by

$$z = w_1 x_1 + w_2 x_2 + b = \sum_{j=1}^{2} w_j x_j + b$$

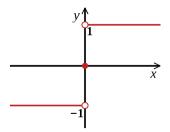


2. the predicted output \hat{y} is calculated by applying the activation function ϕ as

$$\hat{y} = \phi(z).$$

In this case, the activation function is the "sign" function

$$\phi(z) = \begin{cases} +1, & z > 0 \\ 0, & z = 0, \\ -1, & z < 0 \end{cases}$$



https://en.wikipedia.org/wiki/Sign_function

• To simplify the notation we define

$$\bar{x} = (x_1, x_2, 1)$$

 $\bar{w} = (w_1, w_2, b)$

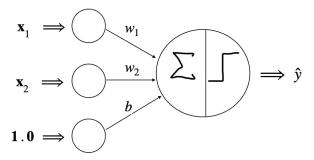
Notice that then

$$z = w^{T}x + b$$

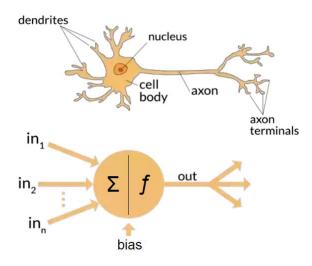
$$= w_{1}x_{1} + w_{2}x_{2} + b$$

$$= \bar{w}^{T}\bar{x}$$

$$= \sum_{i=1}^{3} w_{i}x_{i}$$



• A single artificial neuron is a model of a biological neuron and is a building block for artificial neural networks.



https://towards datascience.com/the-differences-between-artificial-and-biological-neural-networks-a8b46db828b7, which is a substantial of the control of t

• Note that the separator (decision boundary) is of the form

$$w_1x_1 + w_2x_2 + b = 0$$
$$x_2 = -\frac{w_1}{w_2}x_1 - \frac{b}{w_2}$$

which is the equation of the line in the (x_1, x_2) -plane.

• Perceptron

Given the data set

$$D = \{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(n)}, y^{(n)})\}\$$

to find the linear classifier we need to find the model parameters, i.e., the weights w_1 , w_2 and the bias b. Perceptron is one of the methods for finding those parameters.

perceptron (D, T, η) $\bar{w} = \bar{0} \qquad \qquad \text{initialize the parameter } \bar{w} = (w_1, w_2, b) \text{ to be the zero vector for t = 1 to T}$ $\qquad \qquad \text{for i = 1 to n}$ $\qquad \qquad \hat{y}^{(i)} = \phi(\bar{w}^T \bar{x}^{(i)})$ $\qquad \qquad \bar{w} := \bar{w} - \eta(\hat{y}^{(i)} - y^{(i)}) \bar{x}^{(i)}$ return \bar{w}

- Here, T is the number of iterations and η is the learning rate of the ML model (typically, a number between 0 and 1). Both T and η are hyperparameters.
- Let us try to understand the update rule for \bar{w} when we are at a data instance $(x^{(i)}, y^{(i)})$

$$\bar{w}_{new} = \bar{w}_{old} - \eta(\hat{y}^{(i)} - y^{(i)})\bar{x}^{(i)}$$

Recall that $y^{(i)}$ is the actual true class label, while $\hat{y}^{(i)} := h(x^{(i)})$ is the predicted class label computed by our model.

- * If our algorithm did not make a mistake, then $y^{(i)} = \hat{y}^{(i)}$. In that case, $\bar{w}_{new} = \bar{w}_{old}$.
- * If our algorithm misclassified $x^{(i)}$, there are two possibilities:

Case 1:
$$y^{(i)} = 1$$
 and $\hat{y}^{(i)} = -1$

This means that $x^{(i)}$ is a positive example and it is classified as negative. In other words, this means that the algorithm claims

$$\bar{w}^T \bar{x}^{(i)} < 0,$$

while actually $\bar{w}^T \bar{x}^{(i)} > 0$. The rule says to modify \bar{w} so that

$$\bar{w} = \bar{w} + 2\eta \bar{x}^{(i)}.$$

Note that since our algorithm claims $\bar{w}^T \bar{x}^{(i)} = ||\bar{w}|| ||\bar{x}^{(i)}|| \cos \theta < 0$, it means that the angle between \bar{w} and $\bar{x}^{(i)}$ is more than 90. By modifying \bar{w} , we ensure that the angle between updated \bar{w} and $\bar{x}^{(i)}$ is less than 90 as it should be.

Case 2:
$$y^{(i)} = -1$$
 and $\hat{y}^{(i)} = 1$

This means that $x^{(i)}$ is a negative example and it is classified as positive. This means that the algorithm claims

$$\bar{w}^T \bar{x}^{(i)} > 0,$$

while actually $\bar{w}^T \bar{x}^{(i)} < 0$. The rule says to modify \bar{w} so that

$$\bar{w} = \bar{w} - 2\eta \bar{x}^{(i)}.$$

Note that since our algorithm claims $\bar{w}^T \bar{x}^{(i)} = ||\bar{w}|| ||\bar{x}^{(i)}|| \cos \theta > 0$, it means that the angle between \bar{w} and $\bar{x}^{(i)}$ is less than 90. By modifying \bar{w} , we ensure that the angle between updated \bar{w} and $\bar{x}^{(i)}$ is more than 90 as it should be.

- If D is linearly separable, the percentron is guaranteed to converge and to produce a classifier. If D is not linearly separable, the perceptron will not converge (see the proof in [2], pages 18-20).
- Note that the update rule states

$$\begin{bmatrix} w_1 \\ w_2 \\ b \end{bmatrix} = \begin{bmatrix} w_1 \\ w_2 \\ b \end{bmatrix} - \eta(\hat{y}^{(i)} - y^{(i)}) \begin{bmatrix} x_1^{(i)} \\ x_2^{(i)} \\ 1 \end{bmatrix}$$

or, in expanded form,

$$w_1 = w_1 - \eta(\hat{y}^{(i)} - y^{(i)}) x_1^{(i)}$$

$$w_2 = w_2 - \eta(\hat{y}^{(i)} - y^{(i)}) x_2^{(i)}$$

$$b = b - \eta(\hat{y}^{(i)} - y^{(i)})$$

Python code: Lecture_3_Perceptron.ipynb

Homework 1:

- We mentioned that perceptron converges if the data is linearly separable. Try sklearn perceptron model for versicolor and virginica, with sepal length and petal length. What do you observe?
- We created My_Perceptron class for only 2 inputs. Extend this code for 3 inputs. Investigate the iris data set and choose 3 features to classify setosa and versicolor using your code. Notice that you cannot easily plot the decision boundary now since the data is 3-dimensional, but you can still compare the actual and the predicted labels to see how your algorithm is performing.
- Try to generalize My_Perceptron code so it could be used for any number of inputs. (Hint: Recall, that for a list w we can use w[-1] and w[:-1] to access the last value in the list and all the values expect the very last value. Also, use np.dot, NumPy dot product, to compute the pre-activation value of z.)

References and Reading Material:

- [1] Hands-On Machine Learning with Scikit Learn, Keras & TensorFlow, Geron (pages 279-288)
- [2] MIT notes:
 - 2-LinearClassifiers.pdf
 - 3-Perceptron.pdf (the proof of Perceptron convergence on pages 18-20 is optional)

https://openlearninglibrary.mit.edu/courses/course-v1:MITx+6.036+1T2019/course/