





Basic Terminologies

In the previous week, we learnt about dimensionality reduction techniques like PCA and t-SNE, and about clustering algorithms like K-Means clustering, Gaussian Mixture Models, Hierarchical clustering and DBSCAN. These techniques broadly fall into the category of Unsupervised Learning, which is characterized by a lack of "labeled data".

This week we will study **Supervised Learning**, which can broadly be classified into Regression & Classification problems.

Before getting into the content of this week, let us first understand a few basic but helpful terminologies.

Continuous and Categorical variables

A continuous variable can take an infinite number of distinct numerical values, possibly in a given range of numbers. For example, the **Monthly Income** of employees in a certain firm is a continuous variable.

A categorical variable, on the other hand, can take only a limited (finite) number of distinct values. For example, in an image dataset of single handwritten digits, the digit in the image would be a categorical variable because it can only take a finite number of distinct values, in this case from 0 to 9, and nothing beyond that.

Dependent and Independent Variables

In data science, given a set of variables, we need to establish the relationship between one variable and others. The variable to be estimated is **dependent** on the rest of the variables and hence called the **dependent variable**, while the remaining variables that affect the dependent variable are called independent variables.

For example, if we have 4 features Age, Education Level, Work Experience, and Salary, and need to find the relation between the Salary and the rest of the features, the Salary would be the dependent variable while Age, Education Level, and the Work Experience would be independent variables.

Variance and Standard Deviation

In statistics, it is important to understand the magnitude of the spread of the observed data from the Mean.

Variance and Standard Deviation are two quantities that address this concept. To calculate the variance, we take the difference between each number in the dataset and the mean of the data, square this difference to make it positive (independent of sign), and finally divide the sum of the squares by the total number of values in the dataset.

Mathematically, the variance of the population can be given as follows:

$$Var = \frac{1}{n} \sum_{i=1}^{n} (x_i - \mu)^2$$

Where x_i is a data point, μ is the population mean, and n is the total number of data points.

One of the major drawbacks of using variance, to understand the spread of the data, is its interpretability. The unit of variance is the square of the original unit of the data. To overcome this, another quantity is introduced, which is the square root of the variance. This is called the standard deviation of the population.

Mathematically, the standard deviation of the population can be given as follows:

$$Sd = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (x_i - \mu)^2}$$

Being the square root of the variance, the standard deviation is more interpretable, having the same units as the original data points. The standard deviation is able to give a sense for the measure of spread of the dataset around its mean.

Confidence Interval

Inferential statistics is associated with estimating the population **parameters** by extracting samples from the same population. In general, when we make an estimate about some quantity of the population (for example, mean), we come up with a single number. This single number is called a **point estimate**. For example, if we take a sample from a population and the sample mean is 35, then our most reasonable estimate of the population mean is also 35. The drawback of point estimates is that we do not know **how sure** we can be that the population mean is close to 35.

To increase the informativeness of our estimate, we associate it with another concept known as the confidence interval.

A confidence interval is a **range of values around the point estimate**, constructed so that this range will contain the population parameter with a **certain degree of confidence**, expressed in percentage terms.

Let's consider an example: Suppose we are extracting 100 samples (data sets) from a group of students in a university, where each sample has a certain number of records. We have calculated the mean age of students from each sample. When we construct confidence intervals (95% confidence level) using the method that will be covered in the first lecture, 95% of samples (data sets) are expected to result in confidence intervals that contain the true population mean.

The higher the confidence, the greater the width of the confidence interval.

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