

# Monte Carlo simulation

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## 1 Introduction

## 2 Statistical theory

### 2.1 Probability space

The **sample space**,  $S$ , is the set of all possible outcomes.

If a sample space contains a finite number of possibilities or an unending sequence with as many elements as there are whole numbers, it is called a **discrete sample space**.

Example: When rolling a standard six-sided die form the discrete sample space, the possible outcomes are  $S = 1, 2, 3, 4, 5, 6$

If a sample space contains an infinite number of possibilities equal to the number of points on a line segment, it is called a **continuous sample space**.

Example: Measuring the heights of people in a population. This is a continuous sample space, because height can take any real value within a given range.

An **event** is a subset,  $A \subseteq S$ , of the sample space. The event is the amount that contains all possible events. An example of a discrete event could be rolling a die and getting an uneven number, this would be the event  $A = 1, 3, 5$ .

For the continuous event, it could be that a person is between 160 cm and 170 cm tall.

The probability of an event  $A$ ,  $P(A)$ , is the sum of the weights of all sample points in  $A$ . The probability of the whole sample space is 1,  $P(S) = 1$  The probability of any event being between 0 and 1,  $0 < P(A) < 1$  The probability of the empty set being 0,  $P(\emptyset) = 0$

Probability of mutually exclusive events

If  $A$  and  $B$  are mutually exclusive,  $A \cap B = \emptyset$ , then

$$P(A \cup B) = P(A) + P(B)$$

Where  $A$  and  $B$  never occur at the same time, so their union is equal to the two events added together.

Probability of union

$$P(A \cup B) = P(A) + P(B) - P(A \cap B)$$

Here, the union of the two events is  $A$  added to  $B$ , but minus their common event, since it otherwise would be added twice.

Two events  $A$  and  $B$  are independent, if

$$P(A|B) = P(A)$$

The equivalent definition to this is:

Two events  $A$  and  $B$  are independent if and only if

$P(A \cap B) = P(A)P(B)$  This says that the probability of both event  $A$  and  $B$  happening, is equal to the product of the two events.

## 2.2 Probability distribution

Data can come in various distributions depending on different parameters such as degrees of freedom. The distribution is the shape of the data and it will have an effect on statistical models. Therefore it is important to have an understanding of distributions.

### 2.2.1 Normal distribution

In the world of statistics, the most common distribution is the normal distribution. It is constructed as a bell shape. The normal distribution is a continuous distribution, with this density function:

$$n(x; \mu, \sigma) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

The distribution is dependent on the mean( $\mu$ ) and the standard deviation( $\sigma$ ), where changes to the mean will result in a change in the positioning of the normal distribution. Whereas a change in the standard deviation will change the spread of the curve. The normal distribution also always contains an area under the curve that is equal to one. This is to ensure that the normal distribution correctly models probability.

There is a special case of the normal distribution, called the standard normal distribution, where the mean is zero and the standard deviation is one. All variations of a normal distribution can be standardized by a transformation of the distribution, using the Z-score formula.

$$Z = \frac{X - \mu}{\sigma}$$

Z in the Z-score represents the amount of standard deviations a given X value, deviates from the mean.

### 2.2.2 The central limit theorem

A very effective theorem in statistics is the central limit theorem. This theorem states that if a random sample  $\bar{X}$ , with the size  $n$ , is taken from a population with a mean and a finite variance, then as  $n$  goes towards infinity, the distribution will resemble a normal distribution. If used with the Z-score formula, the distribution will resemble a standard normal distribution. The formula for the Z-score, when in conjunction with the central limit theorem, looks like this:

$$Z = \frac{\bar{X} - \mu}{\sigma/\sqrt{n}}$$

Where  $\bar{X}$  is a random sample of size  $n$  and  $\mu$  is the mean of the true population. The standard error is represented by  $\sigma/\sqrt{n}$ , where  $\sigma$  is the standard deviation and  $n$  is the sample size. Usually the standard deviation is unknown, for these

situations it's possible to use the estimator  $S^2$ . This estimates the variance of the population from the variance of the sample, by this formula:

$$s^2 = \sum_{i=1}^n \frac{(x_i - \bar{x})^2}{n-1}$$

The square root of the variance is the standard deviation, therefore the square root of the estimator  $S^2$  would be the estimated standard deviation. The problem with using the estimator  $S^2$ , is that with small samples the variance is small and therefore it contains a lot of bias. In this situation the t-distribution would be used instead of the normal distribution, because the t-distribution takes the bias into account the bias of the standard deviation. It does this by having thicker tails, meaning that the probability of more extreme values are higher.

### 2.2.3 The t-distribution

The t-distribution is shaped as the standard normal distribution, in a bell shape and symmetrical around the mean of zero, the difference is that the t-distribution is more variable. This comes from the fact that the t-distribution is dependent on the degrees of freedom. When the degrees of freedom surpasses 30, the rule of thumb is that the distribution will resemble a normal distribution. So before 30 degrees of freedom, the distribution contains more variance. The t-distribution will come to resemble the standard normal distribution, when it surpasses 30 degrees of freedom, this makes sense, since the two distributions have the same formula:

$$T = \frac{\bar{X} - \mu}{S/\sqrt{n}}$$

The only difference is the estimated standard deviation  $S$ .

## 2.3 Statistical methods

### 2.3.1 Confidence intervals

The confidence interval is a good tool to use, when trying to estimate a parameter of a population. Its used to create an interval, where the parameter has a probability to lie inside of. This probability is called the confidence level and it's a chosen value, usually the chosen confidence level is either 95% or 99%. The confidence interval will become bigger with a larger confidence level. A good confidence interval is small with a large confidence level, this will usually occur when the sample size is large. The chosen confidence level relates to an  $\alpha$ -value, where as an example the chosen confidence level is 95%, then the  $\alpha$ -value would be 5% or normally written as 0.05. The  $\alpha$ -value will sometimes be needed to find the critical value, that is used to calculate the margin of error, as an example it's used when trying to find the critical value of the confidence interval, when working with a t-distribution.

To set up a confidence interval, the margin of error needs to be computed and then that will be both added and subtracted from the point estimate. This will give the values of the outer bounds of the interval. The margin of error is calculated from this formula:

$$\text{Margin\_of\_error} = \text{critical\_value} \pm \text{standard\_error}$$

The standard error will change depending on which parameter that the confidence interval is estimating, but the general formula for the standard error is:

$$\frac{\sigma}{\sqrt{n}}$$

An example of computing a confidence interval of the mean while working with a standard normal distribution, then the formula for the confidence interval would be this:

$$P(-z_{\alpha/2} < Z < z_{\alpha/2}) = 1 - \alpha$$

Where  $1 - \alpha$  is the confidence level. As it's the mean that is being estimated, then instead of Z-score, then  $\mu$  must be isolated and that is done by multiplying  $\frac{\sigma}{\sqrt{n}}$  and subtracting  $\bar{X}$  on all sides, then multiplying all side by  $-1$  to remove the minus sign. So the formula for a confidence interval of the mean will look like this:

$$P(\bar{X} - z_{\alpha/2} \frac{\sigma}{\sqrt{n}} < \mu < \bar{X} + z_{\alpha/2} \frac{\sigma}{\sqrt{n}}) = 1 - \alpha$$

This formula will give the upper and lower bounds of the confidence interval.

### **The interpretation of a confidence interval**

To interpret a confidence interval, it would be incorrect to interpret the confidence level of some value  $x$ , as the probability of the true parameter being inside of the interval. The reason behind this is that the computed interval is static, so either the value  $x$  is inside the interval or it's not. So the correct way of interpreting the confidence interval is by taking multiple samples and computing the confidence interval for all samples, then the value  $x$  would reside inside 95% of the confidence intervals. **Kilde for fortolkningen af kofidense intervaller:**

[http : //www.drhuang.com/science/mathematics/book/probability\\_and\\_statistics\\_for\\_engineering\\_and\\_the\\_sciences](http://www.drhuang.com/science/mathematics/book/probability_and_statistics_for_engineering_and_the_sciences)

### **2.3.2 Hypothesis testing**

A hypothesis test is used to test an assumption about a population. This is done from a sample of the population, as the information about the population is

usually hard to come by. A hypothesis test is set up, by having a null hypothesis and an alternate hypothesis.

$$H_0 = \text{Null hypothesis}$$

$$H_a = \text{Alternate hypothesis}$$

When working with hypothesis testing, the hypothesis  $H_0$  is usually represented as the status quo, where as the hypothesis  $H_a$  is represented as the opposition. It is also important to note that there is only two outcomes of a hypothesis test, either  $H_0$  is rejected in favor of  $H_a$  or  $H_0$  is failed to be rejected. Therefore in no situation can  $H_0$  be stated to be an absolute truth, as there might be other samples where  $H_0$  will be rejected. Therefore in a hypothesis test  $H_0$  needs to be the thing that can be rejected and if  $H_0$  gets rejected, then  $H_a$  will become the new status quo until proven otherwise.

In a hypothesis test  $H_0$  will be the assumption that a parameter for two populations is the same, where as  $H_a$  can be either one of three assumptions, depending on the intention of the hypothesis test.

$$H_0 : \theta = \theta_0$$

$$1. H_a : \theta \neq \theta_0$$

$$2. H_a : \theta < \theta_0$$

$$3. H_a : \theta > \theta_0$$

When the direction of the rejection is not important and also is unknown, then (1) will be the case. This scenario sets up a two-tailed-test, where the hypothesis test is used to reject  $H_0$  if  $H_a$  is either significantly larger or smaller than  $H_0$ , this means that the critical area is on both sides of the difference of  $\theta$  and  $\theta_0$ . Either (2) or (3) will set up a one-tailed-test, where depending on what is important, either the hypothesis test is used to determine if  $H_a$  is significantly bigger or smaller than  $H_0$ . This means that the critical area only spans one side of the difference between  $\theta$  and  $\theta_0$ .



### 3 Polynomial regression

### 3.1 Assumptions

## 4 Pseudo random number generator

## 5 Pseudo Random Number Generator

To generate the dataset, we use a Pseudo-Random Number Generator (PRNG). PRNGs are algorithms that produce sequences of numbers that appear random but are actually deterministically generated from an initial seed value. While these numbers are not truly random, they are sufficiently unpredictable for many practical applications. Random numbers are widely used in fields such as statistics, game theory, cryptography, and simulations. These applications require numbers that behave as if they were random, yet can be reproduced when needed. This is where PRNGs come in—they allow for repeatable randomness, making them ideal for controlled experiments, testing, and security. This chapter will explore the key concepts behind PRNGs. Before going into the mechanics of these generators, it is important to first understand what 'random' means and the characteristics that define truly random numbers.

### 5.1 Properties of PRNGs

The quality of a PRNG is determined by several key factors that influence its use for different applications. Some of the properties of a good PRNG are: Independence, a large period and reproducibility. The numbers produced by the PRNG should be statistically independent, ensuring that each generated value exhibits no correlation with previous numbers or other sequences. This implies that knowledge of previously generated numbers or sequences provides no advantage in predicting the next output. A PRNG operates within a specific interval before its sequence begins to repeat. A high-quality PRNG has a long interval, delaying repetition and enhancing its unpredictability. Conversely, a PRNG with a shorter period becomes more predictable and less suitable for practical use. A key feature of a PRNG is its ability to reproduce the same sequence of numbers when given a specific seed. This property is particularly useful in testing and simulation scenarios, where it is essential to generate identical sequences multiple times for consistency and reproducibility. In addition, a PRNG must be fast and efficient to prevent it from introducing performance bottlenecks within an application. The speed of number generation directly impacts computational efficiency, especially in applications requiring a large volume of random numbers. An inefficient PRNG can significantly slow down processes, undermining the overall performance of the system. Therefore, balancing randomness and efficiency is essential for practical applications.

### 5.2 Linear Congruential Generator

Linear Congruential Sequence (LCS) is a commonly used approach to generate pseudo-random numbers. LCS generates a sequence of numbers using a linear recurrence relation. LCS is expressed as:

$$X_{n+1} = (aX_n + c) \bmod m.$$

where  $X_0$  is the seed,  $a$  is the multiplier,  $c$  is the increment, and  $m$  is the modulus.

**Example:** Given  $a = 5$ ,  $c = 1$ ,  $m = 16$ , and  $X_0 = 7$ :

$$X_1 = (5 \cdot 7 + 1) \bmod 16 = 4$$

$$X_2 = (5 \cdot 4 + 1) \bmod 16 = 5$$

$$X_3 = (5 \cdot 5 + 1) \bmod 16 = 10$$

$$X_4 = (5 \cdot 10 + 1) \bmod 16 = 3$$

This sequence has a period of 16. In an LCG, the period can be as large as  $m$ , but choosing parameters carefully is crucial to achieving long periods.

## **6 Monte Carlo**

### **6.1 Assumptions**

## 7 Problem statement

## 8 Classical regression

## 9 Monte Carlo regression



## 10 Comparison between regressions

## 11 Discussion

## 12 Conclusion

## 13 Litteratur