

Statistical testing of the Sloth function

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1 Introduction

Sloth is a slow-timed hash function where the resulting hash can be verified quickly.

Sloth takes as input a *seed* and a large *prime number*, and is used for uncontrollable random number generation, where the resulting random number is referred to as a *beacon*.

This project consists in testing the existence of a correlation between **the random value produced with a given seed** and **the random value produced with this same seed with one bit flipped**, with all other parameters unchanged.

For this, we realise a Chi-square test of independence based on sample data.

2 Definitions

2.1 The Chi-square distribution

The Chi-square distribution with d degrees of freedom, denoted by χ_d^2 , is the distribution of a sum of the squares of d mutually independent standard normal random variables.

We write :

$$\chi_d^2 = Z_1^2 + Z_2^2 + Z_3^2 + \dots + Z_d^2$$

where $Z_1, \dots, Z_d \stackrel{iid}{\sim} N(0, 1)$ and $d \in \mathbb{N}_{>0}$.

We denote by μ and σ^2 , the mean and the variance respectively. They are expressed as,

$$\mu = E(\chi_d^2) = E\left(\sum_{i=1}^d Z_i^2\right) = \sum_{i=1}^d E(Z_i^2) = d,$$

by linearity of expectation.

$$\sigma^2 = \text{var}\left(\sum_{i=1}^d Z_i^2\right) = \sum_{i=1}^d \text{var}(Z_i^2) = d\text{var}(Z_i^2) = d(E(Z_i^4) - E(Z_i^2)^2) = d(3 - 1) = 2d,$$

using that all Z_i are independent and identically distributed and that $E(Z_i^4) = 3$.

The chi-square distribution is widely used in inferential statistics, and in particular, in hypothesis testing. A test is described as Chi-square if the resulting test statistic obtained follows a Chi-square distribution with d degrees of freedom under the null hypothesis. This Chi-square test statistic is also known as the Pearson statistic and is defined as follows.

2.2 The Chi-square statistic

Let O_1, \dots, O_d be the number of observations of a random sample of size $N = n_1 + \dots + n_d$, falling into the disjoint categories $1, \dots, d$ whose expected values are E_1, \dots, E_d where $E_i > 0$.

Then the **Chi-square statistic** is

$$T = \sum_{i=1}^d \frac{(O_i - E_i)^2}{E_i}.$$

3 Chi-square test of independence applied to Sloth

Let $g(s)$ be the output of *sloth* for the seed s .

We first randomly select the public prime number p of 2048 bits.

For each observation, we randomly select an input $s \in \{0, \dots, p - 1\}$ and define s' obtained by flipping one bit in the sequence s and keeping all other digits identical.

We obtain a pair (b_0, b_1) , where $b_0 = g(s)$ and $b_1 = g(s')$, as shown in Figure 1 below.

For instance, suppose $s=10100$, for which the generator outputs $g(s) = 13 = b_0$.

Now, take $s'=10110$ (2nd bit is flipped) for which the generator outputs $g(s') = 2 = b_1$

We obtain the pair $(13, 2)$.

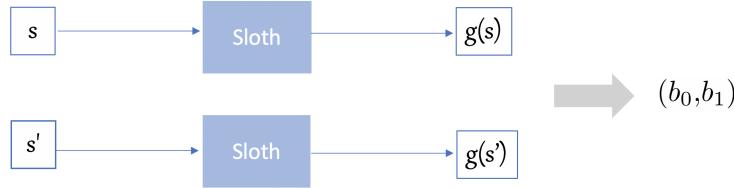


Figure 1: Illustration of 1 observation

3.1 Defining random variables and reporting data

Consider the congruence classes of integers modulo $n : [0]_n, [1]_n, \dots, [n - 1]_n$.

These are all the possible remainders of the division by n .

→ For inputs s and s' , we define the **random variables**:

X : $[i]_n$ such that $b_0 \in [i]_n$.

Y : $[j]_n$ such that $b_1 \in [j]_n$.

$X, Y \in \{[0]_n, [1]_n, \dots, [n - 1]_n\}$.

After gathering the observations, data is expressed in the following contingency table.

We examine the variables X and Y . The variable X represents the rows of the table, so that the rows are labelled: $[0]_n, \dots, [n - 1]_n$. The variable Y represents the columns of the table so that the columns are labelled: $[0]_n, \dots, [n - 1]_n$. Let the observed value be O_{ij} that represents the observed number of cases in row i and column j , that is, observed number of cases where X took value $[i]_n$ and Y took the value $[j]_n$, we denote this event by $[i]_n[j]_n$.

(i.e. $b_0 \in [i]_n$ and $b_1 \in [j]_n$ when considering the pair (b_0, b_1)).

Let the row totals be R_i for row i and the column totals C_j for column j , and N , the sample size.

Variable X	Variable Y					Row Total (R_i)
	$[0]_n$	$[1]_n$	$[2]_n$	\dots	$[n - 1]_n$	
$[0]_n$	O_{11}	O_{12}	O_{13}	\dots	O_{1n}	R_1
$[1]_n$	O_{21}	O_{22}	O_{23}	\dots	O_{2n}	R_2
\cdot	\cdot	\cdot	\cdot	\cdot	\cdot	\cdot
\cdot	\cdot	\cdot	\cdot	\cdot	\cdot	\cdot
\cdot	\cdot	\cdot	\cdot	\cdot	\cdot	\cdot
$[n - 1]_n$	O_{n1}	O_{n2}	O_{n3}	\dots	O_{nn}	R_n
Column Total (C_j)	C_1	C_2	C_3	\dots	C_n	N

3.2 Hypothesis testing

After defining the random variables, we are able to state the null and alternative hypotheses. The null hypothesis is that variables X and Y are independent of each other. The alternative hypothesis is that X and Y are dependent variables.

We state it as :

H_0 : X and Y are independent.

H_1 : X and Y are dependent.

If the null hypothesis H_0 is true, then by **Pearson Theorem**, the Pearson statistic converges in distribution to the Chi square distribution with d degrees of freedom :

$$T = \sum_{i,j=1}^n \frac{(O_{ij} - E_{ij})^2}{E_{ij}} \rightarrow \chi_d^2,$$

where O_{ij} is the observed number of cases where X took value $[i]_n$ and Y took the value $[j]_n$. The degree of freedom d is $(r - 1) \times (c - 1)$, where r and c denote the number of rows and columns respectively, in this case they are both equal to n .

E_{ij} is the expected number of observations in row i and column j . The expected values are computed on the basis of the null hypothesis. If variables X and Y are independent of each other, let the probability that $X = X_i = [i]_n$ and $Y = Y_j = [j]_n$ be P_{ij} ,

$$P_{ij} = P(X_i \cap Y_j) = P(X_i)P(Y_j) = \frac{R_i}{N} \times \frac{C_j}{N}.$$

That is, P_{ij} is the probability of falling in a case which is in row i and column j , under the independence assumption. Thus the expected number of cases in row i and column j is:

$$E_{ij} = P_{ij} \times N = \frac{R_i \times C_j}{N}.$$

3.3 Proof of Pearson's theorem

Each O_{ij} represents the number of time the event $[i]_n[j]_n$ occurred. So, if we consider that the joint distribution O_{11}, \dots, O_{nn} is multinomial with denominator N and probabilities $p_{ij} = E_{ij}/N$, then each O_{ij} follows a Binomial distribution $B(N, p_{ij})$, and thus

$$E(O_{ij}) = Np_{ij} = E_{ij}, \quad \text{var}(O_{ij}) = Np_{ij}(1 - p_{ij}) = E_{ij}(1 - E_{ij}/N) \approx E_{ij}.$$

Thus, by the Central Limit Theorem, $Z_{ij} = \frac{(O_{ij} - E_{ij})}{\sqrt{E_{ij}}} \sim N(0, 1)$ assuming large N , and we have that

$$T = \sum_{i,j=1}^n \frac{(O_{ij} - E_{ij})^2}{E_{ij}} = \sum_{i,j=1}^n Z_{ij}^2 \sim \chi_d^2.$$

The constraint $\sum_{i,j=1}^n O_{ij} = N$ means that only $cr - c - r + 1$ of the Z_{ij} can vary independently, thus reducing the degrees of freedom to $(r - 1)(c - 1)$.

3.4 Sample data analysis

3.4.1 Decision procedure

In this part, we choose a level α at which we want to test H_0 . The most common alpha value is $\alpha = 0.05$, establishing a 95% confidence level.

α is the *probability* of rejecting H_0 when H_0 is actually true.

After setting the alpha value, we compute the critical value cv such that $P(\chi_d^2 \geq cv) = 0.05$ (Figure 2), that is the probability that a random variable following a χ_d^2 distribution takes a value greater than cv .

If the test statistic T takes a value that is greater than cv , it indicates a strong evidence against the null hypothesis, as there is less than 5% chance the null hypothesis is correct. Therefore, we reject the null hypothesis, and accept the alternative hypothesis.

We set the **decision rule**:

Reject H_0 if $T > cv$, otherwise, accept.

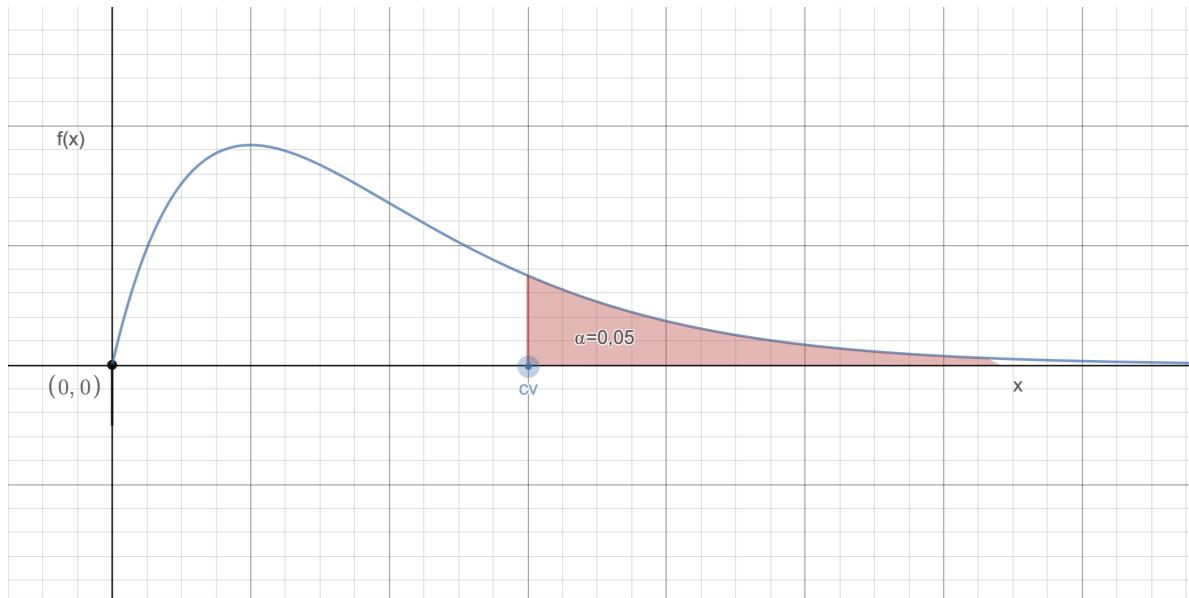


Figure 2: Plot of the Chi-square distribution

3.4.2 Measure of evidence using p-value

The decision procedure as seen above solely consists in either “reject” or “not reject” the null hypothesis. The test statistic T is used to derive the *p-value* that gives more information about the sample data and also measures the evidence against H_0 .

The *p-value* is the probability under H_0 of observing a value of the test statistic more extreme than what was actually observed.

Hence, $P(\chi_d^2 \geq T) = p\text{-value}$ (Figure 3), where T is the Chi-square statistic.

The smaller the *p-value*, the stronger the evidence against H_0 . For this test, we will refer to the commonly used evidence scale shown in Figure 4.

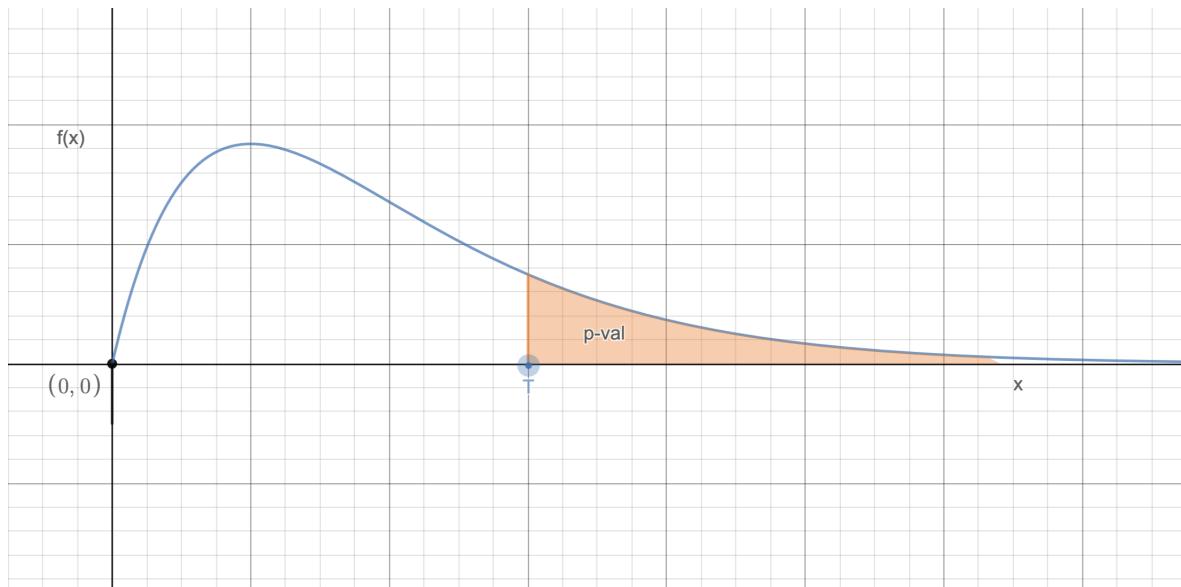


Figure 3: Plot of the Chi-square distribution

p-value	evidence
< .01	very strong evidence against H_0
.01 - .05	strong evidence against H_0
.05 - .10	weak evidence against H_0
> .1	little or no evidence against H_0

Figure 4: The evidence scale

4 Test results

In this section we will present the test results.

We realised 4 tests: by keeping the same prime number p of 2048 bits generated randomly and the same number of iterations, we modified the modulo value n in order to vary the categories. For each n , we also test *sloth* with and without the *xor* operation that is made on the resulted *random number* just before outputting it, to see how this operation affects the result.

- p is generated randomly using the function `mpz_urandomb` provided by GMP library.
- The number of iterations is set to 1. (For 20000 observations, the test took roughly 2 minutes.)
- We generate a different sample for each test.
- In the reported results, we chose to flip the 93rd bit of the binary representation of the seed s for each observation.

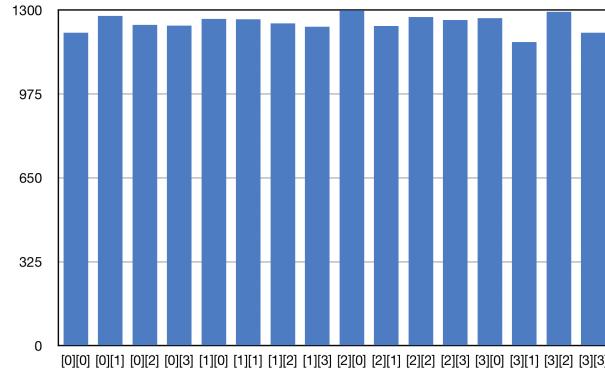
The implementation of the test can be found in <https://github.com/Emna-FENDRI/SlothTest.git>.

After each test, we build the two corresponding **histograms** (with and without the *xor* operation), to visualize the number of observations falling into the n^2 possible categories that we denoted $[i][j]$.

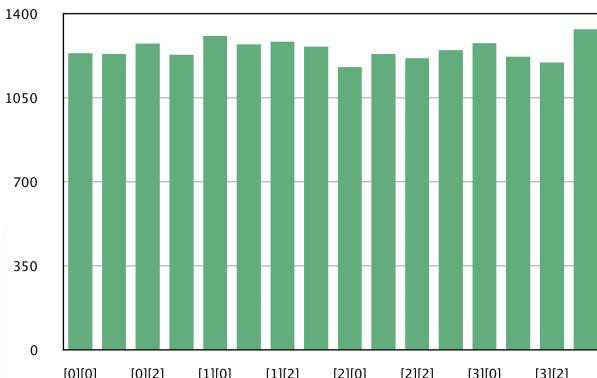
We also **plot** the corresponding χ^2 distribution, its critical value for a 0.05 significance level and the computed test statistics.

4.1 Modulo 4

This first test is done by considering the congruence classes of the generated *beacons* modulo 4, and the corresponding critical value is **cv = 16.919**.



(a) Without *xor*, for 20000 observations



(b) With *xor*, for 20000 observations

	Test Statistic	p-val	Result
Without xor	8.745	0.461	No evidence against H_0
With xor	11.651	0.233	No evidence against H_0

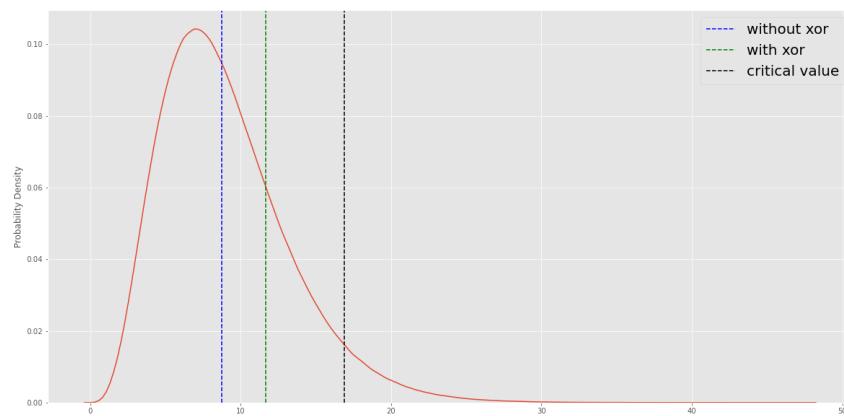
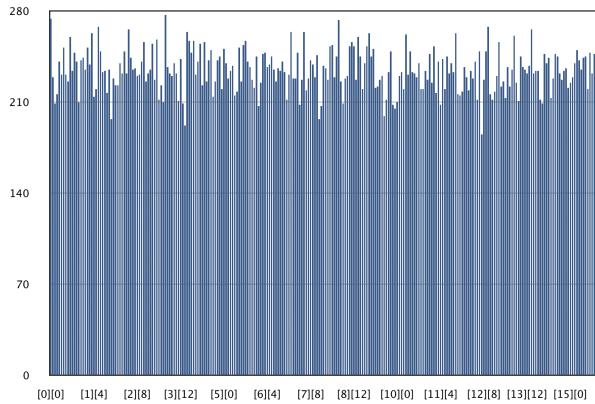


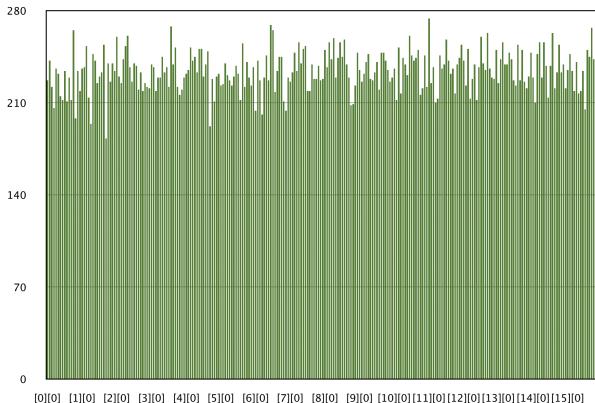
Figure 6: The χ^2_9 probability distribution

4.2 Modulo 16

This second test is done by considering the congruence classes of the generated *beacons* modulo 16, and the corresponding critical value is **cv= 260.992**.



(a) Without xor, for 60000 observations



(b) With xor, for 60000 observations

	Test Statistic	p-val	Result
Without xor	237.940	0,264	No evidence against H_0
With xor	210.511	0,747	No evidence against H_0

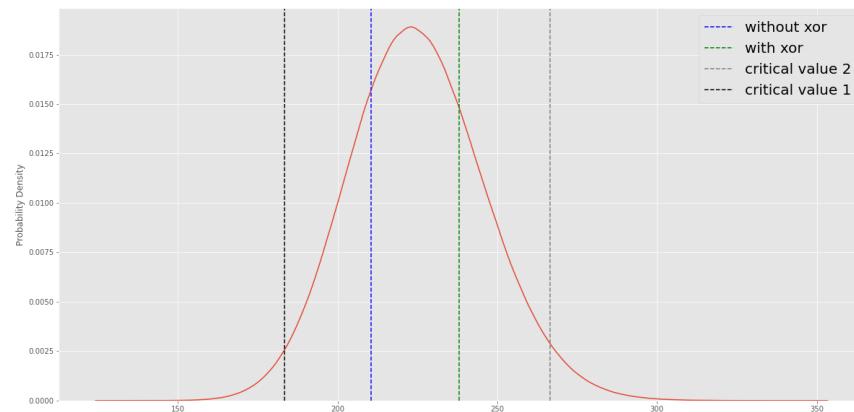
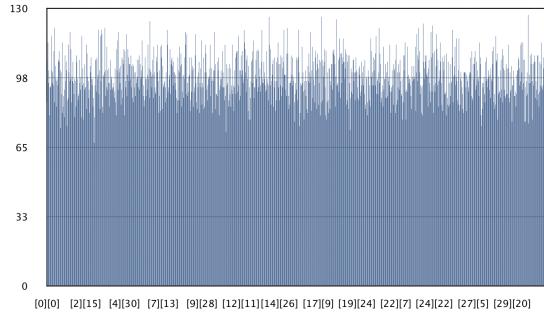


Figure 8: The χ^2_{225} probability distribution

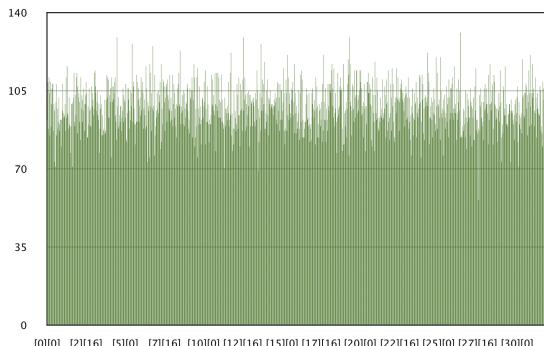
4.3 Modulo 32

This third test is done by considering the congruence classes of the generated *beacons* modulo 32, and the corresponding critical value is $cv = 1034, 2$.

Additionally, due to large degrees of freedom, the χ^2 distribution tends to have a more symmetric shape. So it is interesting to consider a two-tailed test, where each tail covers a probability of 0.025, hence obtaining two critical values: **$cv_1 = 875.072$** and **$cv_2 = 1046.93$** , as shown in Figure 10.



(a) Without *xor*, for 100000 observations



(b) With *xor*, for 100000 observations

	Test Statistic	p-val	Result
Without xor	955.616	0,542	No evidence against H_0
With xor	948.131	0,610	No evidence against H_0

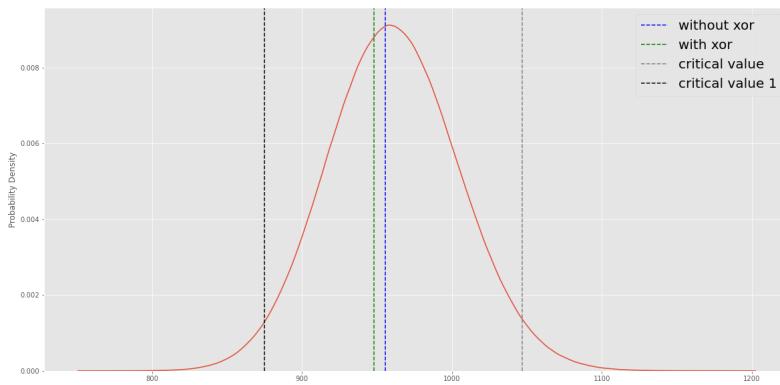
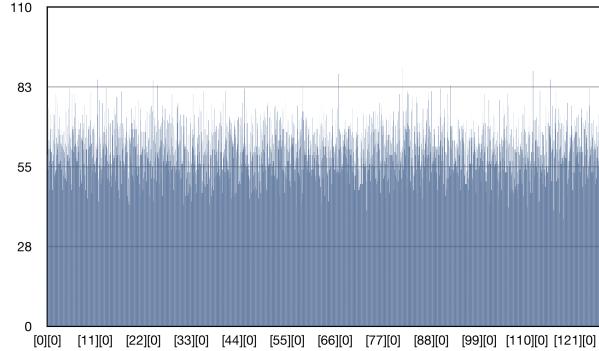


Figure 10: The χ^2_{961} probability distribution

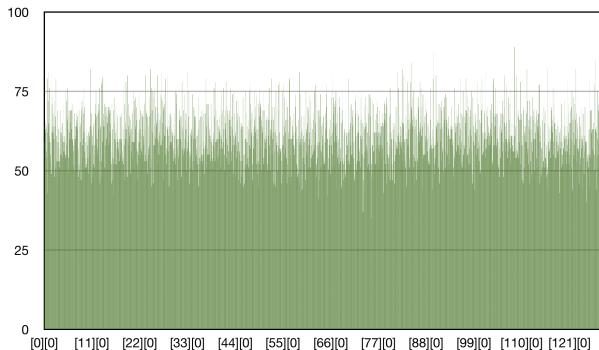
4.4 Modulo 128

This last test is done by considering the congruence classes of the generated *beacons* modulo 128, and the corresponding critical value is $cv = 16425.3$.

Similarly as above, due to large degrees of freedom, we also consider a two-tailed test, where each tail covers a probability of 0.025, hence obtaining two critical values: $\mathbf{cv_1 = 15777}$ and $\mathbf{cv_2 = 16481}$.



(a) Without *xor*, for 1000000 observations



(b) With *xor*, for 1000000 observations

	Test Statistic	p-val	Result
Without xor	16043.631	0,737	No evidence against H_0
With xor	16221.472	0,3024	No evidence against H_0

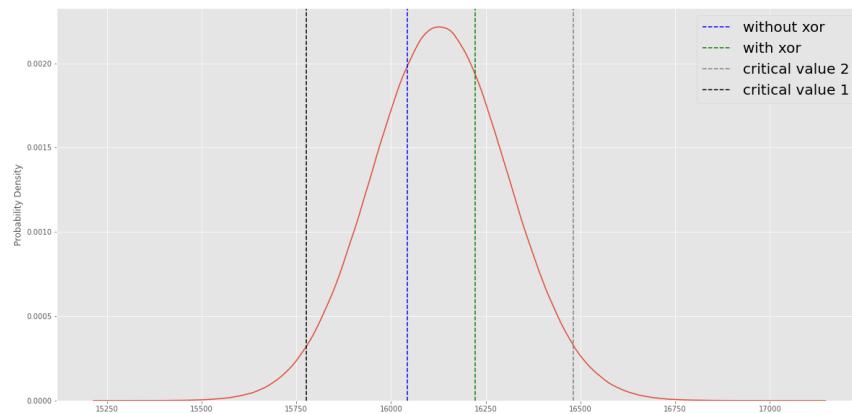


Figure 12: The χ^2_{16384} probability distribution

5 Conclusion

The results obtained in the previous section are very convincing. As we can see from the plots, the test statistics with and without the *xor* clearly fit in the χ^2 distribution, which proves that there is no evidence against H_0 , as expected.

Finally, we can conclude that variables X and Y are not correlated. In fact, the *beacon* resulting from a given seed s will not give any information about the *beacon* obtained from s' .

6 References

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