Monte Carlo project

Stratification

Marina BLAZEVIC, Mehdi EL KACEMY, Samy KOBBITE

ENSAE Paris

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Introduction

Estimation by Monte Carlo and Quasi Monte Carlo methods

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4 Approach by Importance Sampling

Bibliography

Project Framework

• We want to estimate the integral of the following function :

$$u \in [0,1]^d$$
, pour $d \ge 1$: $f(u) = 1 + \sin\left(2\pi\left(\frac{1}{d}\sum_{i=1}^d u_i - \frac{1}{2}\right)\right)$

For this, we will use several stratification methods..

- Idea: Integrate numerically by subdividing the integration interval, and performing a Monte Carlo simulation in each subinterval with a different number of random points.
- Advantage: Reduce the variance of the estimate of the integral compared to a simple Monte Carlo simulation
- We estimate the integral I of a function f(x) on the interval [a, b] by estimating I:

$$I \approx \frac{b-a}{N} \sum_{i=1}^{N_s} \frac{1}{n_i} \sum_{j=1}^{n_i} f(x_{ij}^{(i)})$$

 N_s : number of sub-intervals

 n_i : number of random points in the subinterval i

 $x_{ii}^{(i)}$: j-th random point in subinterval i

Monte Carlo method

Principle

Let n be the number of random points generated in the domain $[0,1]^d$. We can estimate the integral of the function f(u) using the following formula:

$$I \approx \frac{1}{n} \sum_{i=1}^{n} f(u_i)$$

where u_i is the *i*-th random point in the domain $[0,1]^d$.

• Quasi Monte Carlo method:

The generation of random numbers is replaced by a deterministic sequence of points such as the Halton sequence or the quasi-random sequence of Sobol.

Principle

```
def monte_carlo_integration(d,Ns):
    samples = np.random.uniform(0, 1, size=(Ns,d))
    values = f(samples)
    return np.mean(values)
```

Figure: Code for MC

```
def quasi_monte_carlo1(d, Ns):
    samples = stats.qmc.Sobol(d).random(Ns)
    values = f(samples)
    return np.mean(values)
```

Figure: Code for QMC with Sobol

Construction of QMC according to Sobol or Halton

Generation according to Sobol

Method based on a family of sequences of binary numbers constructed from irreducible polynomials of high degree.

Polynomials used to generate sequences of binary numbers which are then converted into reals via inverse transformation.

• Generation according to Halton:

Choice of a numeric base b and generation of a sequence of numbers x_i which are fractions in base b

• In practice:

Sobol is more efficient for large dimensions Halton easier to generate

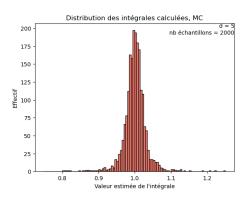
Construction of our comparison metrics

 Mean Squared Error: mean of the squares of the differences between the estimated values and the true values

$$\mathsf{MSE} = \frac{1}{N} \sum_{i=1}^{N} (\hat{I}_i - I)^2$$

• **Confidence intervals**: intervals of probable values for the value of the integral.

Comparison Between MC and QMC Sobol



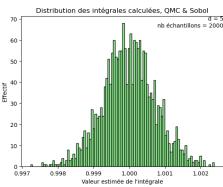


Figure: Distribution MC

Figure: Distribution QMC Sobol

For d = 5, nb samples = 2000 and nb estimates = 1000: MSE.MC = 0.00018 and MSE.QMC = 1.8955e-07 CI.MC = (0.97438, 1.0497) and CI.QMC = (0.9617, 1.0384)

Comparison between MC and QMC Sobol

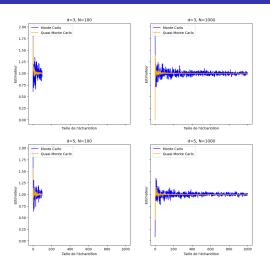


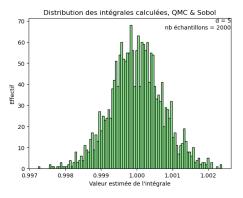
Figure: Comparison of convergences between MC and QMC for different parameter values

Comparison Between MC and QMC Sobol

C	ł	nb_echantillons	nb_estimations	MSE_MC	MSE_QMC
	1	10000	1000	0.000324	2.2554e-09
	2	10000	1000	0.000612	4.4849e-09
	3	10000	1000	0.000971	6.1972e-09
	4	10000	1000	0.001279	9.1086e-09

Figure: Comparison between MC and QMC Sobol for different values of d

Comparison between Sobol and Halton for QMC



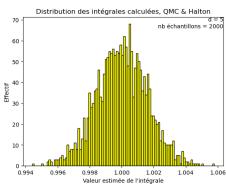


Figure: Distribution QMC Sobol

Figure: Distribution QMC Halton

For d = 5, nb samples = 2000 and nb estimates = 2000:

 $\mathsf{MSE.S} = 1.8728\text{e-}07 \text{ and } \mathsf{MSE.H} = 7.1403\text{e-}07$

CI.QMCS = (0.9862, 1.0139) and CI.QMCH = (0.9860, 1.0137)

Comparison between Sobol and Halton for QMC : numerical errors

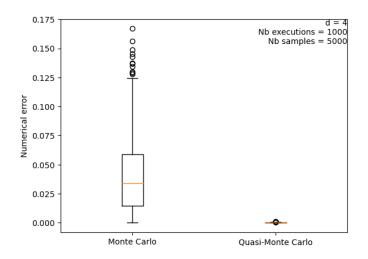


Figure: Numerical errors for MC and QMC

Article frame

- Article by Nicolas Chopin and Mathieu Gerber.
- Proposition of 2 unbiased estimators of the integral of the function f over $[0,1]^s$ depending on a regularity parameter $r \in \mathbb{N}$
- Based on cubic stratification
- Here, the goal is to implement the Haber estimators of order 1 and 2:

$$\begin{split} I_{1,k}(f) &:= \tfrac{1}{k^s} \sum_{c \in C_k} f(c + U_c), \quad U_c \sim U\left(-\tfrac{1}{2k}, \tfrac{1}{2k}\right)^s \\ &\text{et } I_{2,k}(f) := \tfrac{1}{k^s} \sum_{c \in C_k} g_c(Uc), \ U_c \sim U\left[-\tfrac{1}{2k}, \tfrac{1}{2k}\right]^s \end{split}$$

with

$$C_k = C_{0,k}$$
 and $C_{m,k} = \left\{ \frac{2j_1+1}{2k}, \dots, \frac{2j_s+1}{2k} \right\} \mid (j_1, \dots, j_s) \in \{-m, \dots, k+m-1^s\}$ and $g_c(u) := \frac{f(c+u)+f(c-u)}{2}$, where $n=2k$

Haber of order 1

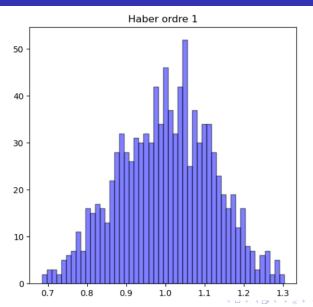
```
def haber_ordre1(N):
    n = k**s
    estimates = []
    for _ in range(N):
        Uc = [random.uniform(-1/(2*k), 1/(2*k)) for i in range(s)]
        I = 0
        for c in C(k, s):
        | I += f(tuple(ci + ui for ci, ui in zip(c, Uc)))
        I /= n
        estimates.append(I)
    return estimates
```

Figure: Our code for Haber 1

Haber of order 2

Figure: Our code for Haber 2

Result for Haber of order 1



Results for Haber of order 1 and 2

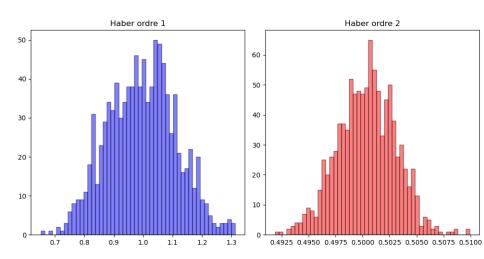


Figure: Distribution of estimators for s = 4, k = 5, N = 1000

Comparison in terms of speed and MSE

L'algorithme monte_carlo_integration a pris 0.0009996891021728516 secondes pour s'exécuter. L'algorithme quasi_monte_carlo a pris 0.0020079612731933594 secondes pour s'exécuter. L'algorithme haber_ordre1 a pris 55.08616590499878 secondes pour s'exécuter.

Figure: Time for d = 10, s=4, k=5, number of samples = 10000

L'algorithme monte_carlo_integration a pris 0.001999378204345703 secondes pour s'exécuter. L'algorithme quasi_monte_carlo a pris 0.003002166748046875 secondes pour s'exécuter. L'algorithme haber_ordre1 a pris 0.4043452739715576 secondes pour s'exécuter.

Figure: Time for d=4, s=2, k=2 and number of samples =10000

 $\label{eq:mc_mc} \mbox{'MSE_MC} = 0.006731834267663948 \mbox{ et } \mbox{MSE_QMC} = 9.053440901287969e-07 \mbox{ et } \mbox{MSE_haber1} = 7.284517252308317e-05\mbox{'MSE_MC} = 0.006731834267663948 \mbox{ et } \mbox{MSE_QMC} = 9.053440901287969e-07 \mbox{ et } \mbox{MSE_haber1} = 7.284517252308317e-05\mbox{'MSE_MC} = 0.006731834267663948 \mbox{ et } \mbox{MSE_NC} = 0.0067318342676394 \mbox{ et } \mbox{ et } \mbox{MSE_NC} = 0.0067318342676394 \mbox{ et } \mbox{MSE_NC} = 0.0067318342676394 \mbox{ et } \$

Figure: MSE for d = 4, s = 2, k = 2 and number of samples = 1000

Principle

• Importance sampling is a numerical method for estimating the integral of a function h(u) over the interval $[0,1]^d$:

$$\int_{[0,1]^d} h(u) du$$

- Use a probability distribution different from the uniform distribution over [0,1]^d to generate samples, and weight each sample according to the probability of being chosen from this distribution.
- Here, generation of the samples u_1, u_2, \ldots, u_N from a probability distribution p(u) different from the uniform distribution, weighting of each sample according to the ratio between the function h(u) and the weighting function g(u).
- The estimator of the integral is then given by:

$$\hat{I} = \frac{1}{N} \sum_{i=1}^{N} \frac{h(u_i)}{g(u_i)}$$



Our function

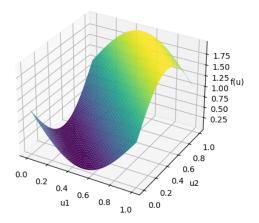


Figure: Studied function in multidimensionnal view

Our method

We chose g(u) such as that when

$$||\mathbf{u}||_1 = \sum_{i=1}^n |u_i|$$

is extreme the weight is more important in the distribution.

Hence, when we have important values that strongly affects the value of the integral, we are sure that they are taken into account.

Once we are sure that they are taken into account, we downweight them

Our function

```
def g(u, n, value1, value2):
   norm1 = np.sum(np.abs(u))
   if norm1 > 3 * d / 4 or norm1 < d / 4:
        return value1
   else:
        return value2</pre>
```

Figure: Our g function to create weights

```
The best estimated value of the integral is 1.0045025285020868
The optimal value1 for function g is 0.0
The optimal value2 for function g is 1.0
```

Figure: Estimated best values

Bibliography

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Bibliography

The link to our Git: https://github.com/marinablaz/ProjetMonteCarlo