# **Monte Carlo** Quasi Monte-carlo (QMC)

Francesco Banterle, Ph.D.

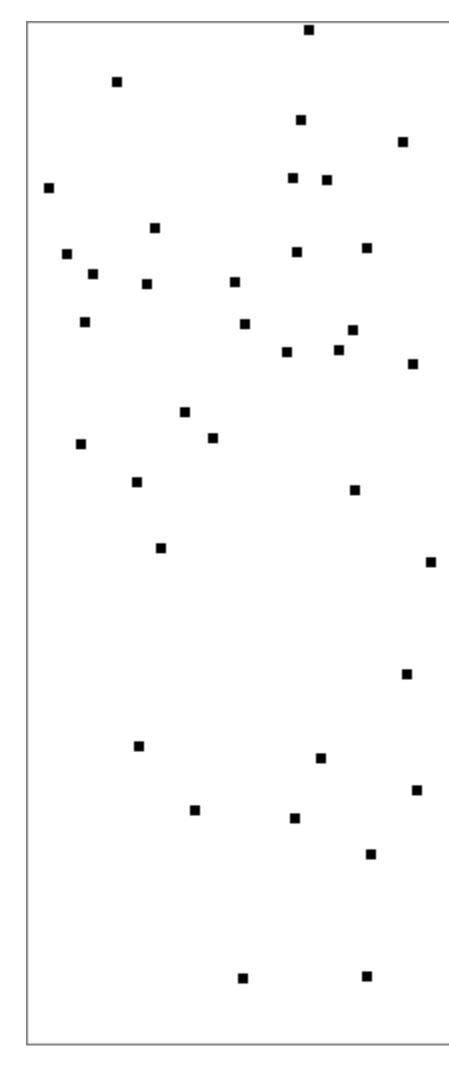
## Quasi Monte-Carlo Introduction

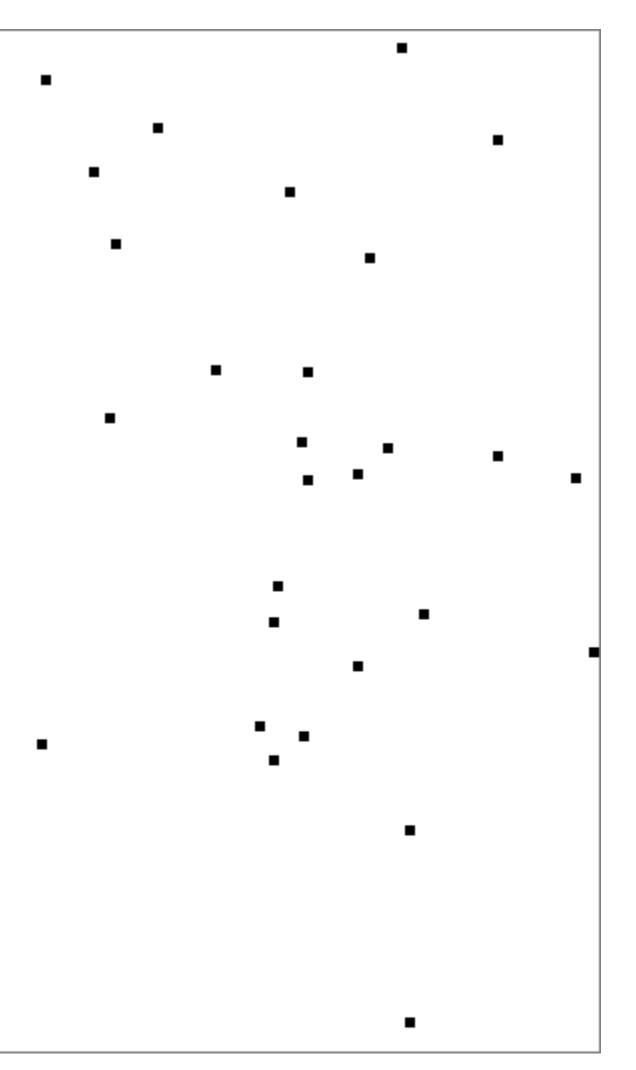
- quantiles, and ratios.
- In Quasi Monte-Carlo or QMC, our goal is to "bend" this law using deterministic samples.
  - We may get better results than the ones of classic Monte-Carlo!

In Monte-Carlo, we have seen that we use randomness to estimate averages,

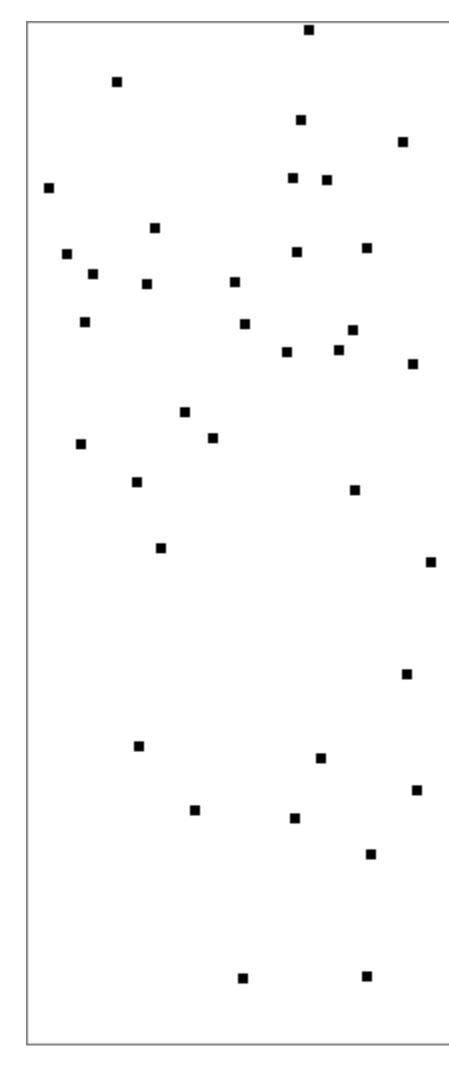
The justification why this works is thanks to the Law of Large Numbers.

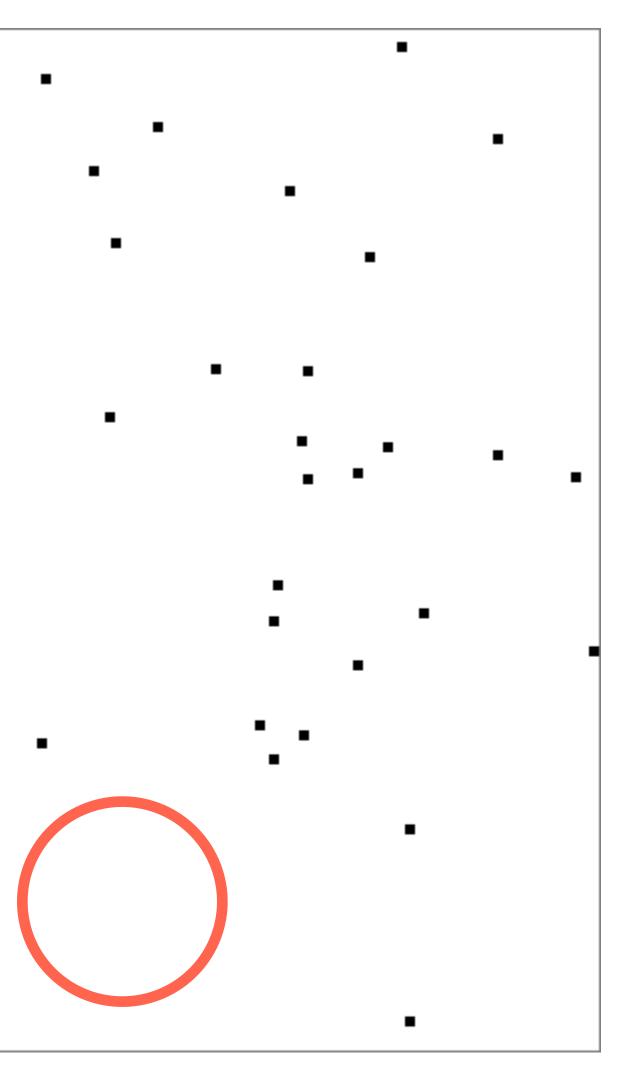
### **Quasi Monte-Carlo** Motivation





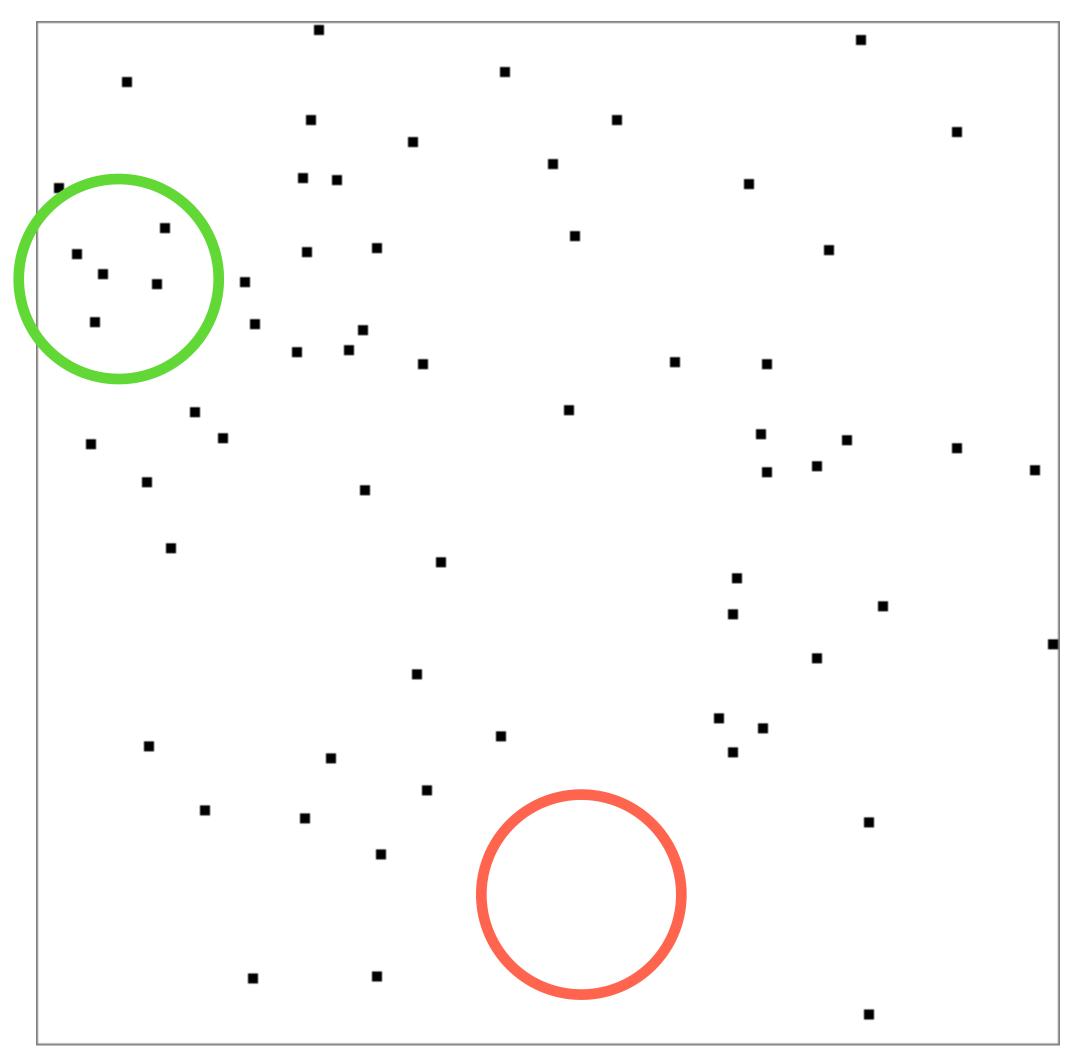
### **Quasi Monte-Carlo** Motivation





#### Gaps

### **Quasi Monte-Carlo** Motivation



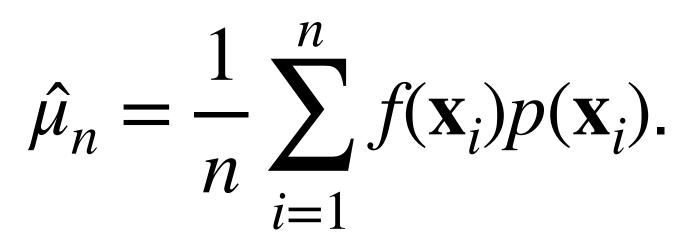
#### Cluster

#### Gaps

#### **Quasi Monte-Carlo** Introduction

• We still estimate:

- - We are half-way between regular grids and Monte-Carlo.
- measures are typically called discrepancies.



• Now, our samples,  $\mathbf{x}_i$ , are deterministic points that fill  $[0,1]^d$  in an even way:

In QMC, how to measure the uniformity of our samples is important, and

• Let's define an interval in d dimension as:

$$\prod_{i=1}^{d} [a_i, b_i] = \left\{ \mathbf{x} \in \mathbb{R}^d \,\middle|\, \forall_{j \in [1,d]} \quad x_j \in [a_i, b_i] \right\}$$

The local discrepancy of n samples x<sub>i</sub> is defined as:

$$\delta(\mathbf{a}) = \frac{1}{n} \sum_{i=1}^{n} \mathbf{1}_{\mathbf{x}_i \in [\mathbf{0}, \mathbf{a})} - \prod_{i=1}^{d} a_i.$$

$$\mathbf{a}, \mathbf{b} \in \mathbb{R}^d \land \forall_i a_i \leq b_i.$$

• Let's define an interval in d dimension as:

$$\prod_{i=1}^{d} [a_i, b_i] = \left\{ \mathbf{x} \in \mathbb{R}^d \, \middle| \, \forall_{j \in [1,d]} \quad x_j \in [a_i, b_i] \right\} \qquad \mathbf{a}, \mathbf{b} \in \mathbb{R}^d \land \forall_i a_i \le b_i.$$

• The local discrepancy of *n* samples **x**<sub>*i*</sub> is defined as:

$$\delta(\mathbf{a}) = \frac{1}{n} \sum_{i=1}^{n} \mathbf{1}_{\mathbf{x}_i \in [\mathbf{0}, \mathbf{a})} - \prod_{i=1}^{d} a_i.$$

The number of points in [0, a)

• Let's define an interval in d dimension as:

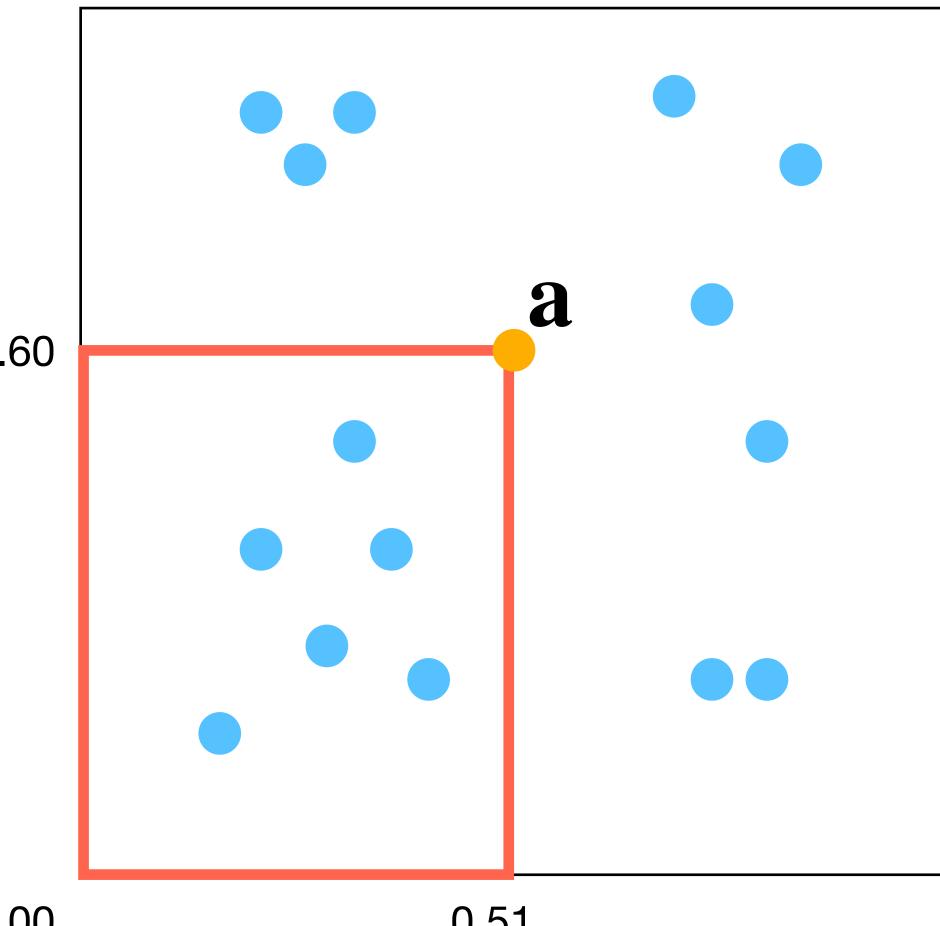
$$\prod_{i=1}^{d} [a_i, b_i] = \left\{ \mathbf{x} \in \mathbb{R}^d \, \middle| \, \forall_{j \in [1, d]} \quad x_j \in [a_i, b_i] \right\} \qquad \mathbf{a}, \mathbf{b} \in \mathbb{R}^d \land \forall_i a_i \le b_i.$$

• The local discrepancy of *n* samples **x**<sub>*i*</sub> is defined as:

$$\delta(\mathbf{a}) = \frac{1}{n} \sum_{i=1}^{n} \mathbf{1}_{\mathbf{x}_i \in [\mathbf{0}, \mathbf{a})} - \prod_{i=1}^{d} a_i.$$
 Volume

The number of points in [0, a)

#### Quasi Monte-Carlo The Start Discrepancy: Example



0.60



$$\delta(\mathbf{a}) = \frac{6}{14} - 0.6 \cdot 0.50 = 0.42 - 0.30 = 0.42$$

0.51

0.12

• When  $\delta(\mathbf{a}) = 0$  there is the perfect balance.

$$D_n^{\star} = D_n^{\star}(\mathbf{x}_1, \dots, \mathbf{x}_n) = \sup_{\mathbf{x} \in [0,1)^d} \left| \delta(\mathbf{x}) \right|$$

• A sequence,  $\mathbf{X}_1, \ldots, \mathbf{X}_n$ , is low discrepancy when:

$$D_n^{\star}(\mathbf{x}_1, \dots, \mathbf{x}_n) = O\left(\frac{(\log n)^d}{n}\right), n \to \infty.$$

### Quasi Monte-Carlo **Trade-offs**

- When we use low discrepancy sequences,  $\mathbf{x}_1, \ldots, \mathbf{x}_n$ , we cannot use the CLT anymore.
- We have Koksma-Hlawka Theorem:

$$\left|\frac{1}{n}\sum_{i=1}^n f(\mathbf{x}_i) - \int_{[0,1)^d} f((x)d\mathbf{x}\right| \le D_n^{\star} \cdot V_{HK}(f),$$

where  $V_{HK}$  is the Hardy and Krause total variation.

- What does this mean?
  - If  $V_{HK}(f) < \infty$  and we approximate  $D_{f}$

 $|\hat{\mu} -$ 

$$P_n^{\star} = o(n^{-1+\epsilon})$$
 with  $\epsilon > 0$ , we have:  
 $\mu \mid = o(n^{-1+\epsilon}).$ 

Low Discrepancy Sequences

#### Low Discrepancy Sequences **Radical Inverse Function**

• The radical inverse function is a simple function defined as:

$$\Phi_b(i) = \sum_{k=0}^{\infty} d_{k,b}(i)b^{-k-1}$$

• This function is based on the fact that we can encode a number i as a sequence of digits:

$$i =$$

•  $\Phi_b$  transforms a positive integer into a floating-point in [0,1) by reversing its digits:

$$\Phi(i)_b$$
 =

• Van Der Corput's sequence is a simple 1D sequence that is based on the radical inverse function using base 2:

$$\mathcal{X}_{i}$$

$$b \ge 2 \land d_{k,b}(i) \in \{0, \dots, b-1\}.$$

$$\sum_{k=0}^{\infty} d_{k,b}(i)b^k.$$

 $\Phi(i)_b = 0.d_{i,0}d_{i,1}...d_n.$ 

 $c_i = \Phi_2(i).$ 

#### Low Discrepancy Sequences **Radical Inverse Function: Example**

 $n = 1 = 1 \times 2^0 + 0 \times 2^2 + ... = (...001)_2$  $\Phi(1)_2 = (0.100...)_2 = 1 \times 2^{-1} = 0.5$ 

 $n = 2 = 0 \times 2^{0} + 1 \times 2^{1} + 0 \times 2^{2} + ... = (...0010)_{2}$  $\Phi(2)_2 = (0.010...)_2 = 0 \times 2^{-1} + 1 \times 2^{-2} = 0.25$ 

#### Low Discrepancy Sequences Radical Inverse Function: Example

i	Binary	Reversed	$\Phi_2(i)$
1	1	0.1	0.5
2	10	0.01	0.25
3	11	0.11	0.125
4	100	0.001	0.0625

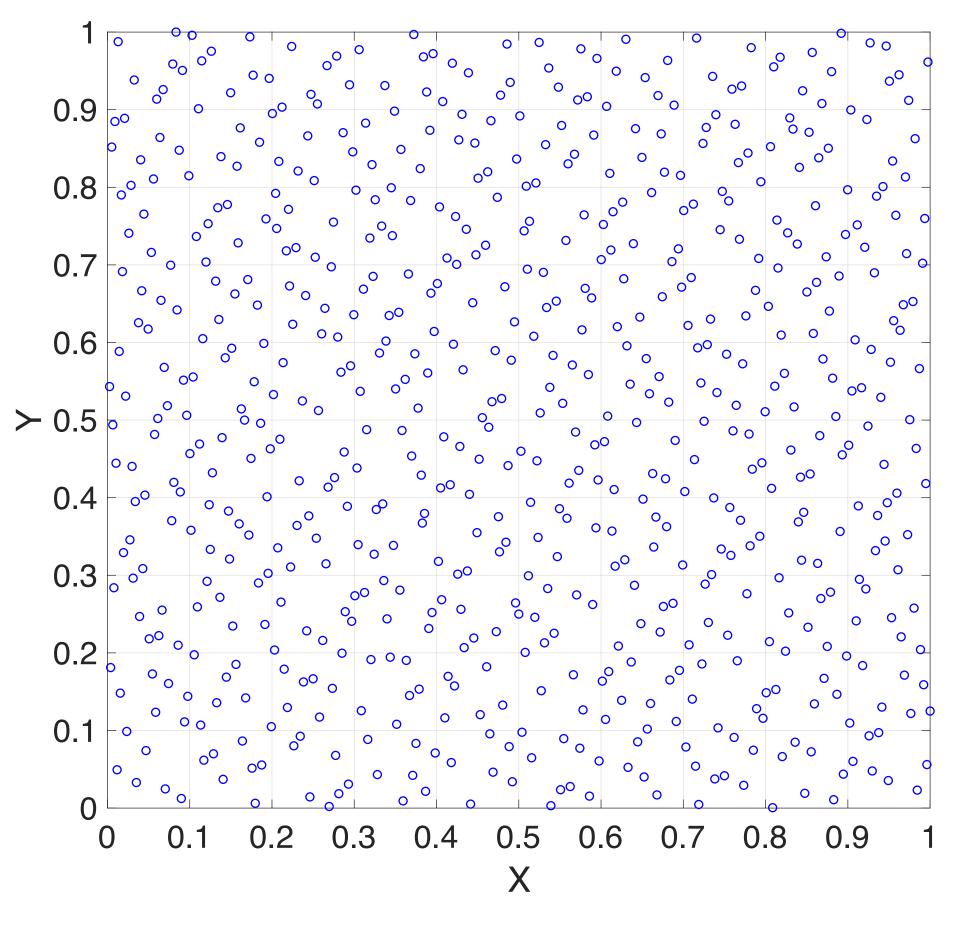
#### Low Discrepancy Sequences Halton Sequence

- The Halton sequence employs the radical inverse base.
- In this case, we use a different base for each dimension:
  - Each base needs to be co-prime with the others!
  - A popular choice is to use the first d-prime for generating a d-dimension vector:

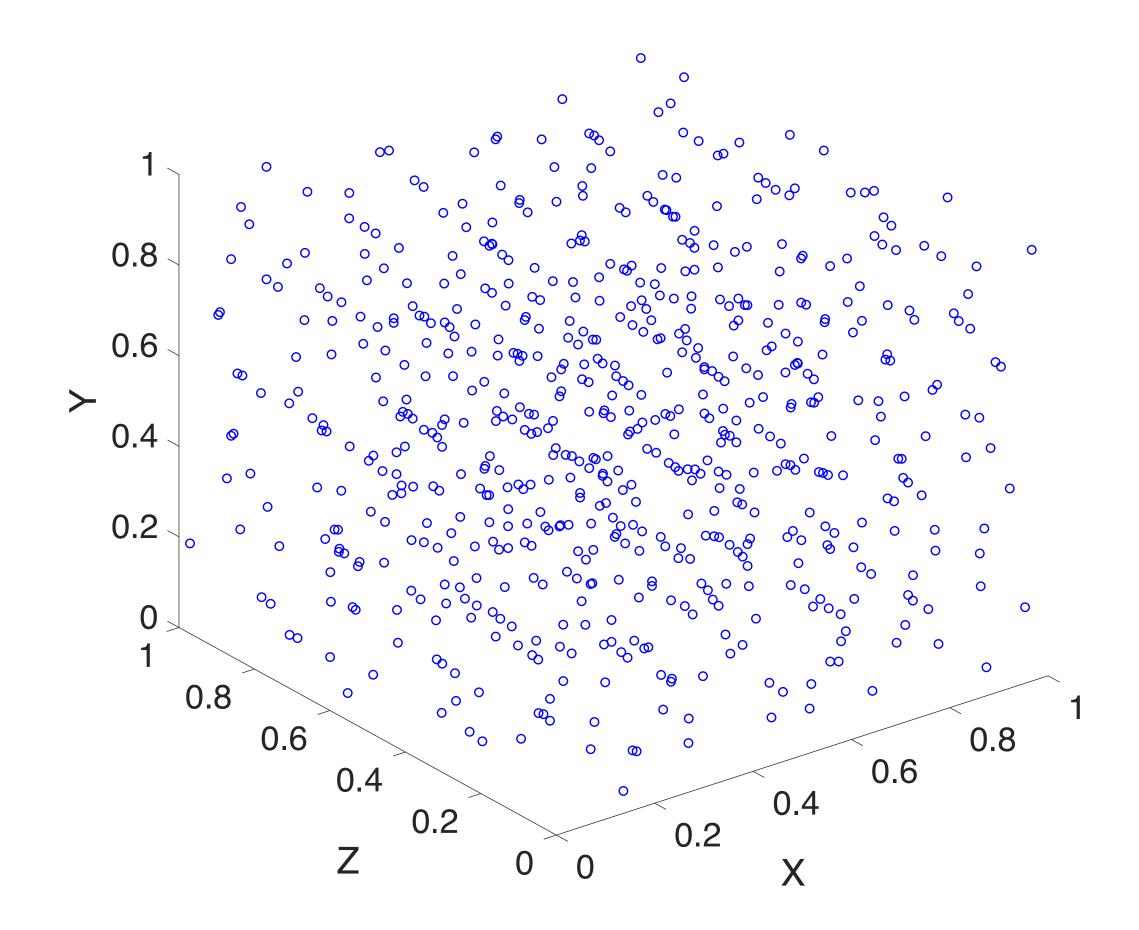
$$\mathbf{x}_{i} = \left(\Phi_{2}(i), \Phi_{3}(i), \dots, \Phi_{p(d)}(i)\right),$$

where p(k) is the k-th prime number.

#### Low Discrepancy Sequences Halton Sequence: Example

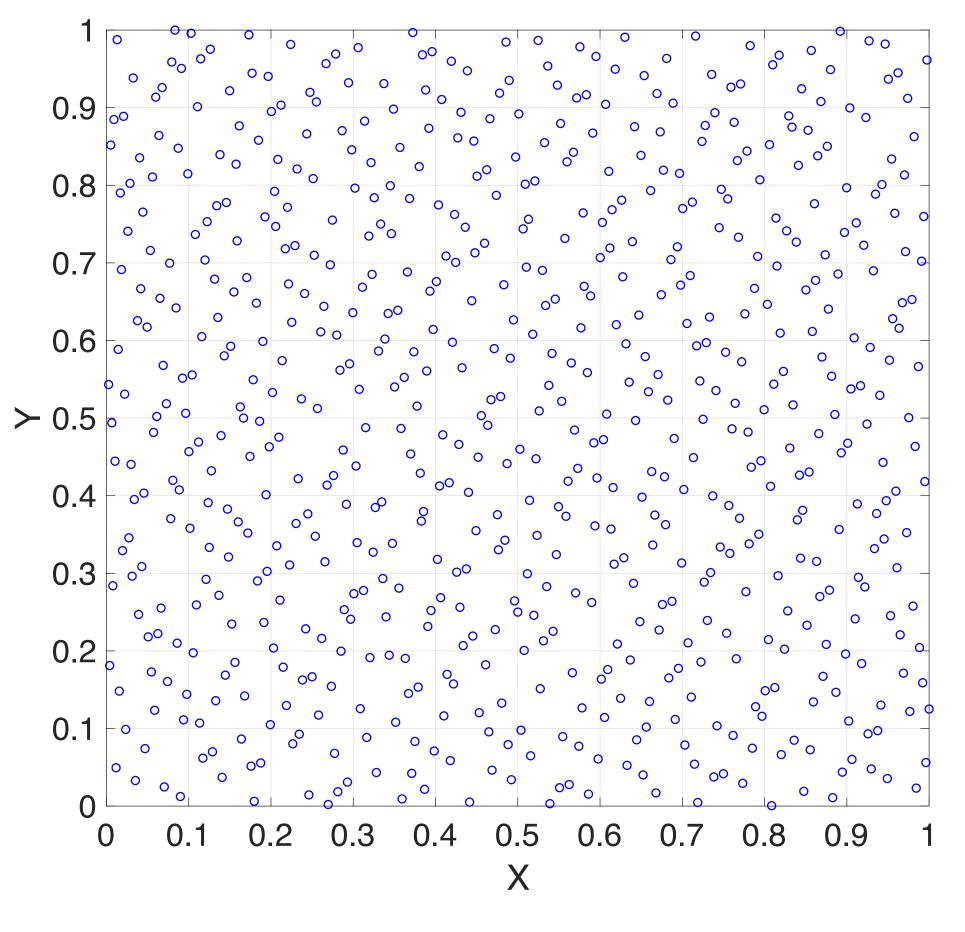


Base X = 2; Base Y = 3

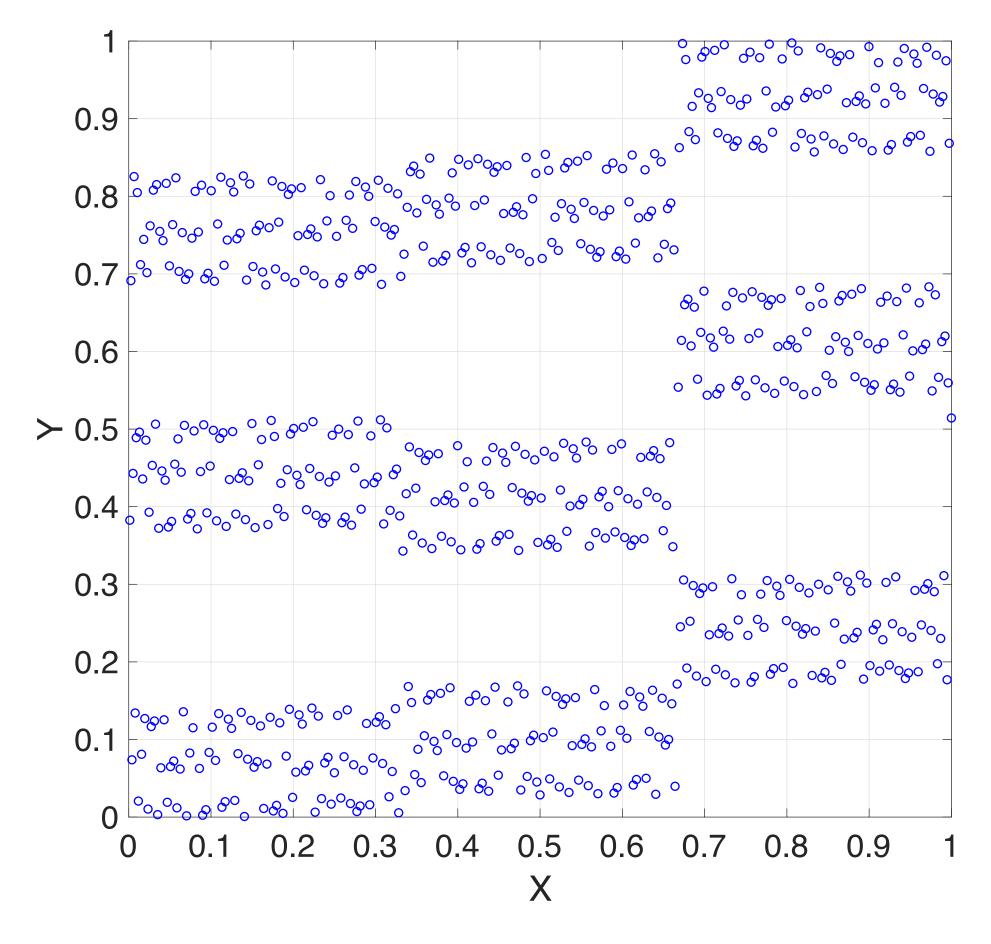


Base X = 2; Base Y = 3; Base Z = 5

#### Low Discrepancy Sequences Halton Sequence: Example



Base X = 2; Base Y = 3



Base X = 2; Base Y = 6

#### Low Discrepancy Sequences Halton Sequence

• The discrepancy when generating a d-dimensional vector is:

where *n* is the number of samples.

 $O\left(\frac{(\log n)^d}{n}\right),$ 

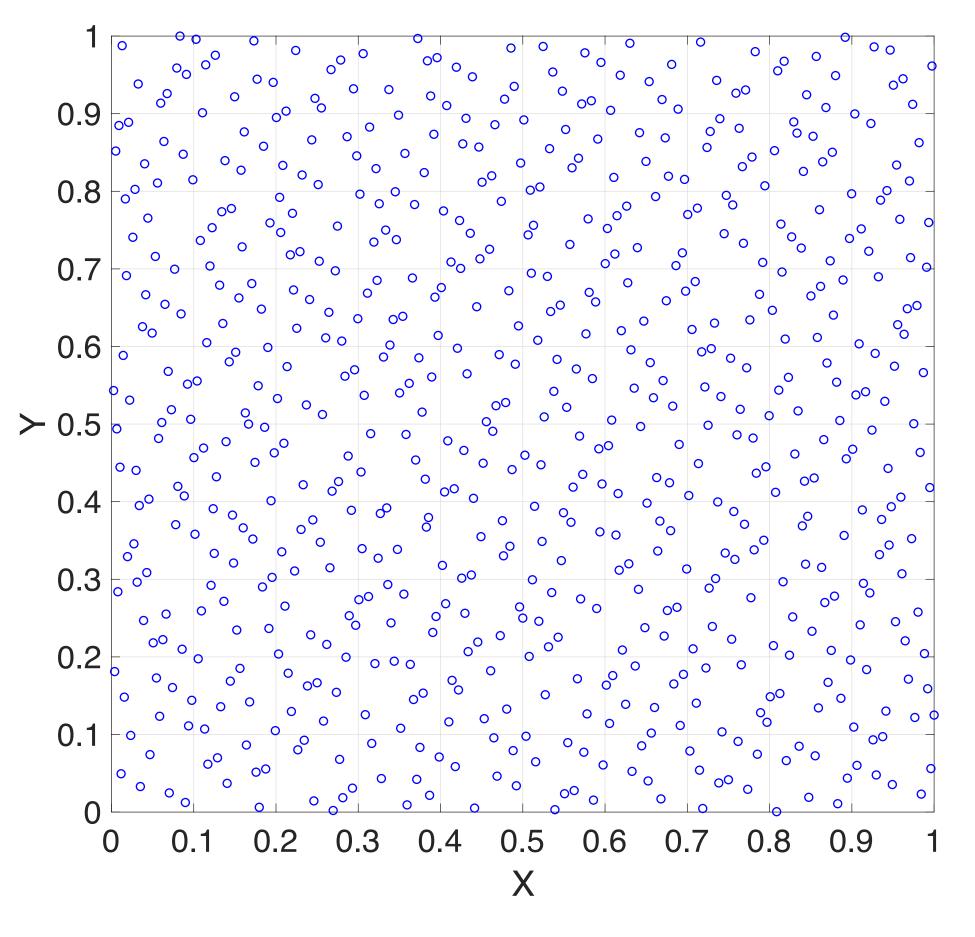
#### Low Discrepancy Sequences Hammersley Sequence

- The Hammersley sequence employs as well the radical inverse base.
- Again, we use a different base for each dimension:
  - Each base needs to be co-prime with the others!
  - As before, we use the first (d 1)-prime for generating a d-dimension vector. The vector, compared to Halton's one, has the following change in the generation:

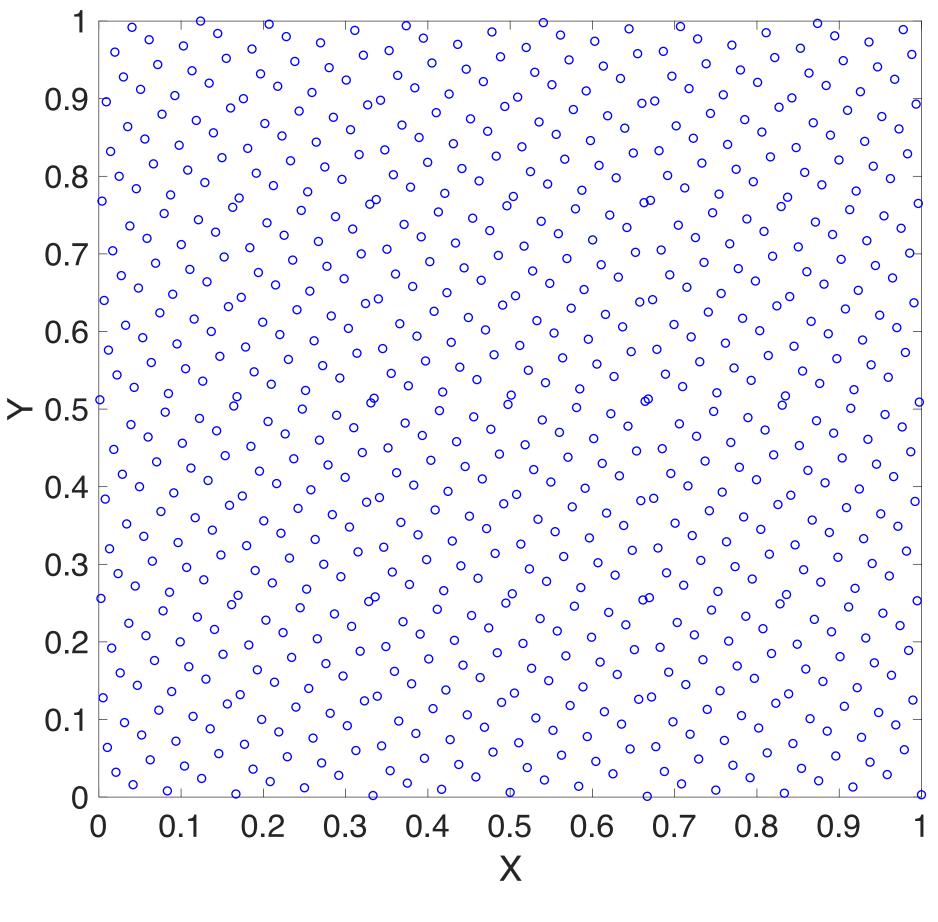
$$\mathbf{x}_{i} = \left(\Phi_{2}(i), \Phi_{3}(i), \dots, \Phi_{p(d-1)}(i), \frac{i}{n}\right)$$

• Note: the number of samples, *n*, has to be known in advance!

#### Low Discrepancy Sequences Hammersley Sequence: Example



Halton Sequence



Hammersley Sequence

#### Low Discrepancy Sequences Halton Sequence

• The discrepancy when generating a d-dimensional vector is:

where *n* is the number of samples.

 $O\left(\frac{(\log n)^{d-1}}{n}\right),$ 

### Low Discrepancy Sequences Limitations

- Both Halton sequence and Hammersley sequence have some issues:
  - We may have regular patterns.
  - They are not ideal for parallel applications:
    - All threads will generate the same sequence!
- A possible solution is to randomize such sequences:
  - We apply a random permutation for the digits of a number.

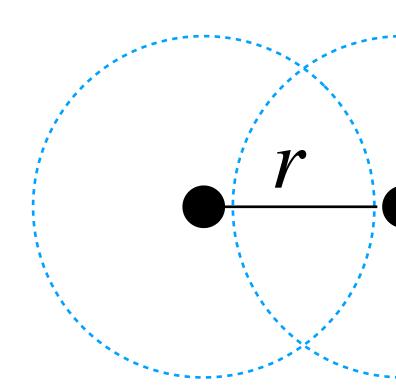
# Low Discrepancy Sequences Other Sequences

- Faure: is based on Van der Corput's sequences, but there is only a base for different dimensions. This is a large prime number:
  - We have permutations with each dimension.

- Sobol: based on algebra of polynomials in  $\mathbb{F}_2$ :
  - It can be computed using Gray codes.

# **Poisson-Disk Sampling**

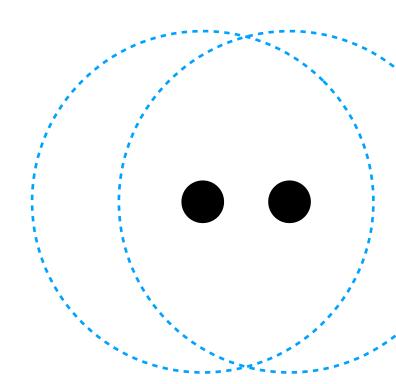
- Poisson-disk sampling is a sequential random process for generating samples in a domain.



#### • Each generated sample/point has to be "disk-free" for a minimum distance r:



- Poisson-disk sampling is a sequential random process for generating samples in a domain.



#### • Each generated sample/point has to be "disk-free" for a minimum distance r:

No Disk-Free

- without regularity.
- i.e., the spectrum of a sequence has certain properties:
  - uniformity.
  - isotropic.

#### • This method does not guarantee low-discrepancy, but it creates point-sets

• The goal of this sequence is to generate samples with blue noise properties;

- radius according a PDF, different distributions, etc.
- The most famous algorithms:
  - distance  $d \ge r$ .
  - close to others; i.e.,  $d \leq r$ .
  - Spatial data structures helps in reducing computational complexity:
    - Bridson 2007 algorithm.

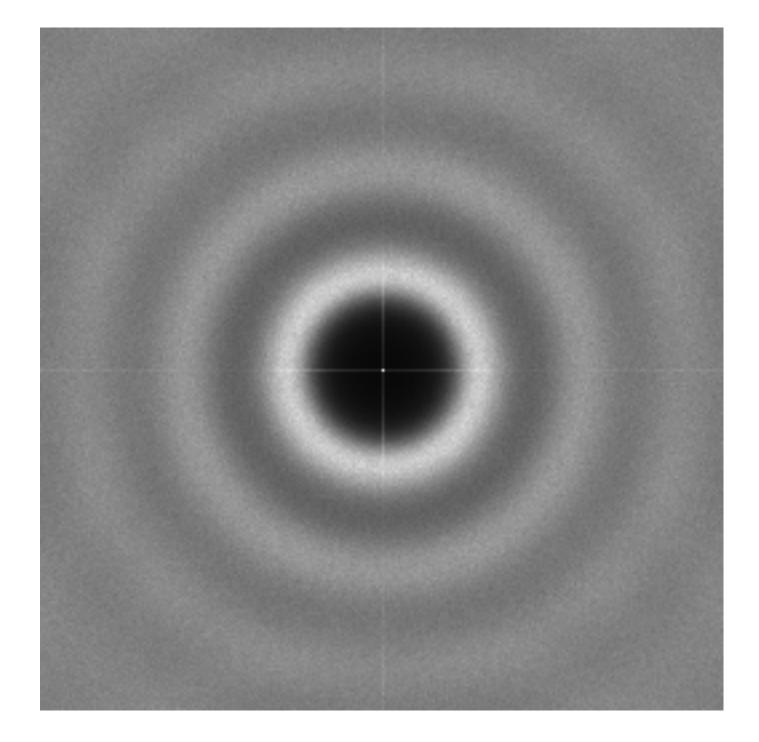
• To achieve Poisson-Disk Sampling, there are a huge literature: 2D, nD, spatially varying

• Dart Throwing: we draw a sample,  $\mathbf{x}_i$ , we accept it if its neighbors are at a minimum

• Samples removal: we draw a huge number of samples, we remove that samples that

#### **Possin-Disk Sampling** Example

Samples



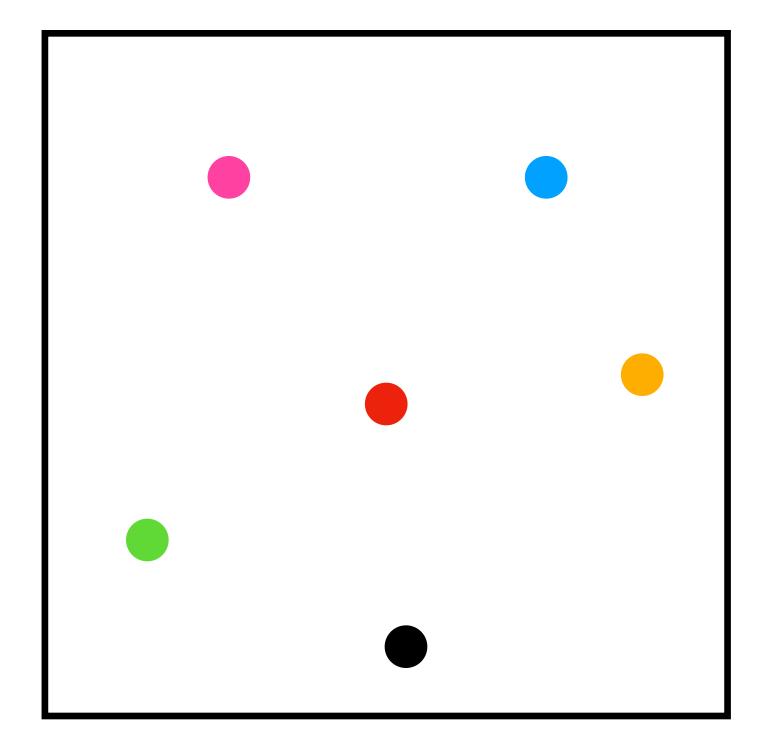
#### Periodogram

# Randomized QMC

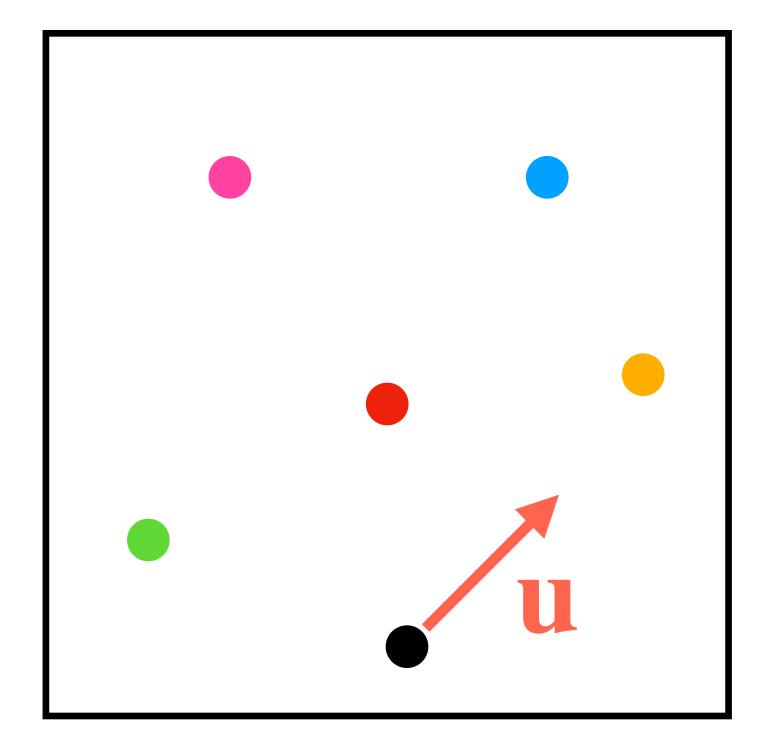
## **Randomized QMC** Main Idea: Cranley-Patterson Rotation

- One problem of QMC is that if we run it on parallel, all threads will start to generate exactly the same samples!
- Another issue is that we cannot have the error estimation that we have in classic Monte-Carlo.
- A solution is to apply a random shift to the sequence:
  - $\mathbf{x}'_i = \mathbf{x} + \mathbf{u} \mod 1 \qquad \mathbf{u} \in \mathbf{U}(0,1).$
- This solution is called Cranley-Patterson rotation.

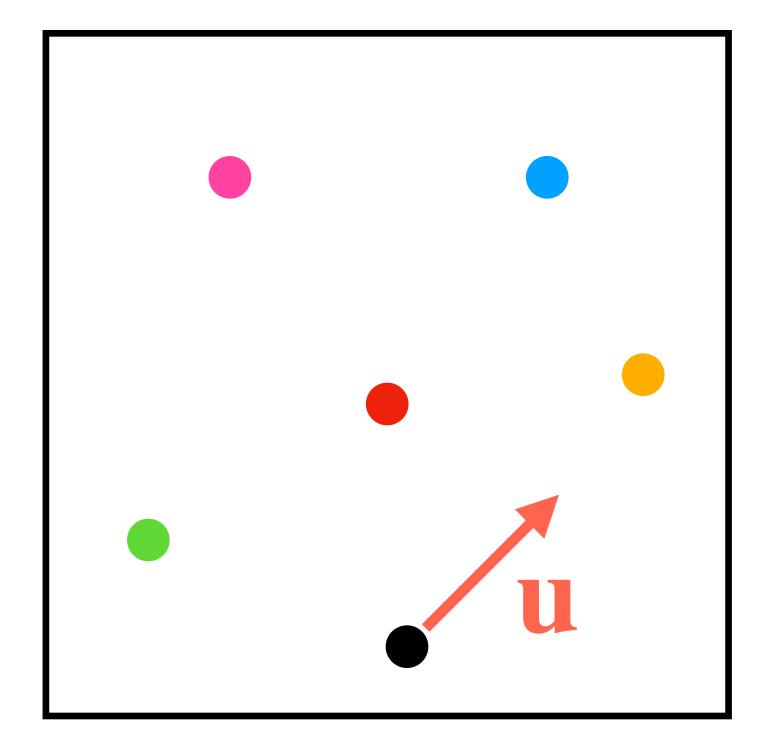
# Main Idea: Cranley-Patterson Example

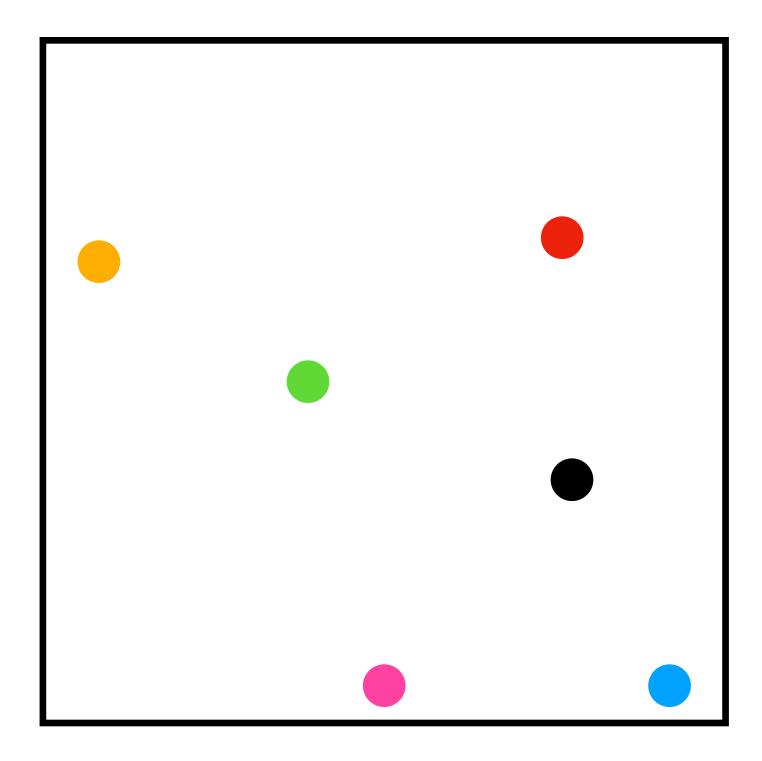


# Main Idea: Cranley-Patterson Example



# Main Idea: Cranley-Patterson Example





# **Randomized QMC** Main Idea: Scrambling

- Cranley-Patterson rotation works and is low discrepancy. However, it does not preserve stratification properties of a sequence.
- A solution is scrambling the digits of numbers in a sequence. For example in 1D:

$$x = \sum_{i=0}^{\infty} x_i b^{-i-1} \to x' = \sum_{i=0}^{\infty} x'_i b^{-i-1},$$

• Where we apply random permutations:

and  $\pi$  are permutations of  $\{0, \dots, b-1\}$ .

 $x'_{0} = \pi(x_{0})$   $x'_{1} = \pi_{x_{0}}(x_{1})$   $x'_{2} = \pi_{x_{0},x_{1}}(x_{2})$ • • •

# Bibliography

- Art Owen. "Chapter 15: The Quasi-Monte Carlo parts" from the book "Monte Carlo theory, methods and examples". 2019.
- Paolo Brandimarte. "Handbook in Monte-Carlo Simulation". Wiley. 2014.
- Matt Pharr, Greg Humphreys. "Chapter 7: Sampling and Reconstruction" from the book "Physically Based Rendering - Second Edition". Morgan Kaufmann. 2010.

Thank you for your attention!