

#### **Numerical Methods**

**Sampling Domains** 

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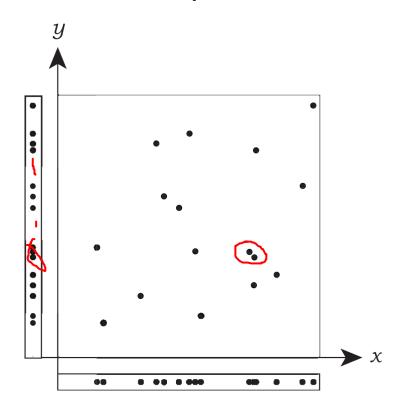
### 2D Sampling

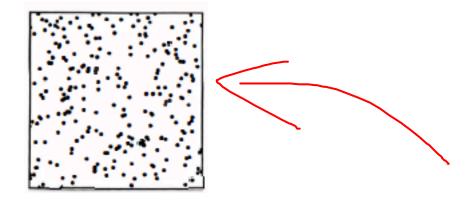
- Assume we are sampling a function on a unit square
- Good sampling
  - Not structured...involves some amount of randomness
  - Uniform(ish) distribution...avoid gaps and clumps
  - Projections into 1D along x and y are also uniform(ish)
  - There is a non-trivial minimum distance between all sample points
- Such a sample pattern is called Well-Distributed



#### Random

Too irregular
Oversamples some areas...
Undersamples others





By random we mean a floating point values generated by a pseudo-random number generator of reasonable quality like Python's:

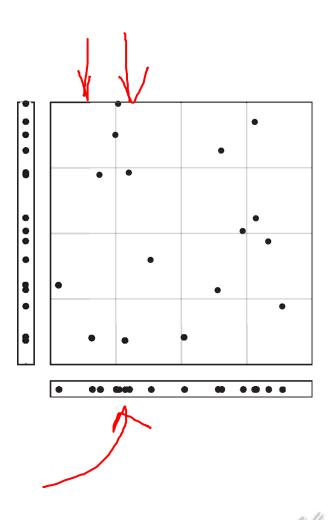
random. random()

Return the next random floating point number in the range [0.0, 1.0).



#### **Jittered**

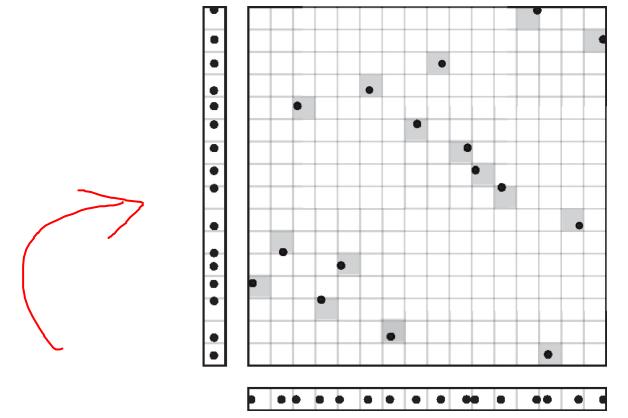
- Create a n x n grid covering the domain
  - Or a n<sup>d</sup> uniform grid in d dimensional space
- Randomly generate a sample in each cell
- Example of stratified sampling
  - Each cell is a strata
- Significantly better than random
- x-y projections can still be poorly distributed





### n-rooks

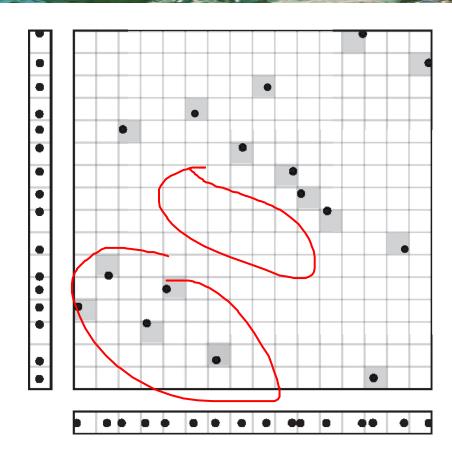
- Also called Latin hypercube sampling
- Use and n x n grid
- One sample exactly in each row and column
  - Again, randomly position a sample within the cell containing it
  - If samples were rooks in chess, no captures can occur





#### n-rooks

- Produced by random shuffle of diagonal samples
  - Must maintain the rook condition
- Use n samples instead of n<sup>2</sup> as in jittered
- 1D distributions are good...better than jittered
- 2D barely better than random...worse than jittered

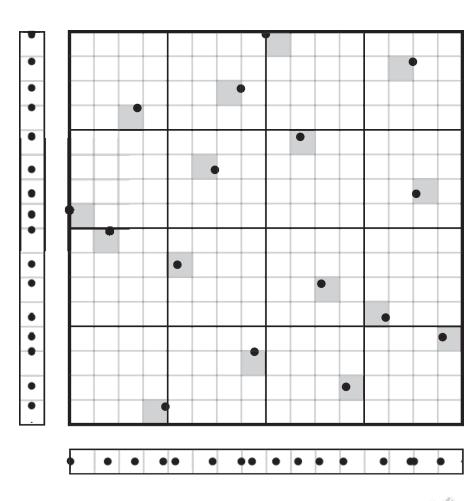


Developed by UIUC alum Pete Shirley in 1991



## Multi-Jittered Sampling

- We use two grids
- For n samples with n a perfect square
  - Coarse grid is √n X √n
  - Fine grid is n X n
  - 1 sample per coarse grid cell
  - For each select unique row & column of fine grid
  - Randomly position sample in fine grid cell
- Good 1D projections from the rook condition
- Good 2D distribution from stratification
- Very good sampling technique



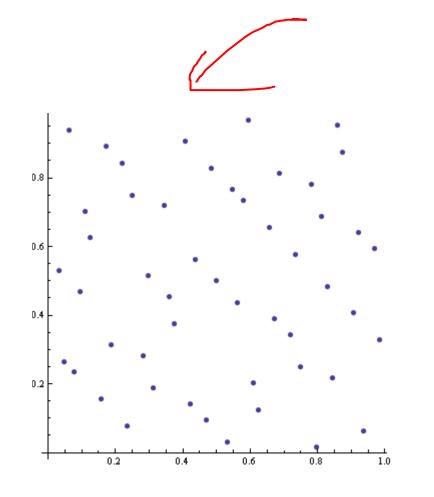


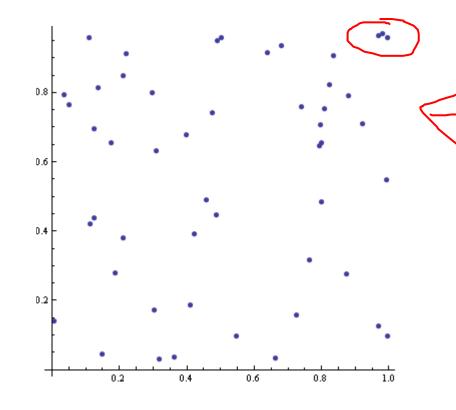
# **Hammersley Sampling**

#### Deterministic quasi-random sequence

Low-discrepancy sequences are also called quasirandom sequences, due to their common use as a replacement of uniformly distributed random numbers. The "quasi" modifier is used to denote more clearly that the values of a low-discrepancy sequence are neither random nor pseudorandom.









### Hammersley Sampling

$$\Phi_2(i) = \sum_{j=0}^n a_j(i) 2^{-j-1} = a_0(i) \frac{1}{2} + a_1(i) \frac{1}{4} + a_2(i) \frac{1}{8} \dots$$

#### Radical inverse function of integer i to base 2

- reflect binary digits of i across decimal point
- evaluate this new number now in [0,1)

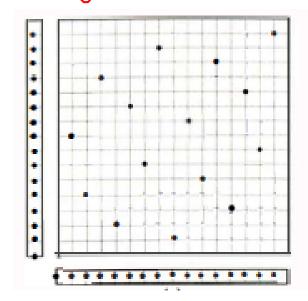
Y			$\overline{}$		•	
i			Φ <sub>2</sub> (i) (base 2)			
1	=	12	.12	-	1/2	0.5
2	=	102	.012	=	1/4	0.25
3	=	112	.112		1/2 + 1/4	0.75
4	=	1002	.0012	-	1/8	0.125
5	-	1012	.1012	=	1/2 + 1/8	0.635
6	-	1102	.0112	_	1/4 + 1/8	0.325
7	=	1112	.1112		1/2 + 1/4 + 1/8	0.875
8	-	1000 <sub>2</sub>	.00012	=	1/16	0.0625

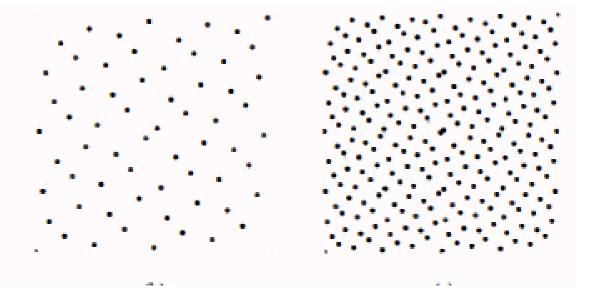


## Hammersley Sequence in 2D

Set of n 2D samples in unit square:

$$p_i = (x_i, y_i) = [i/n, \Phi_2(i)]$$

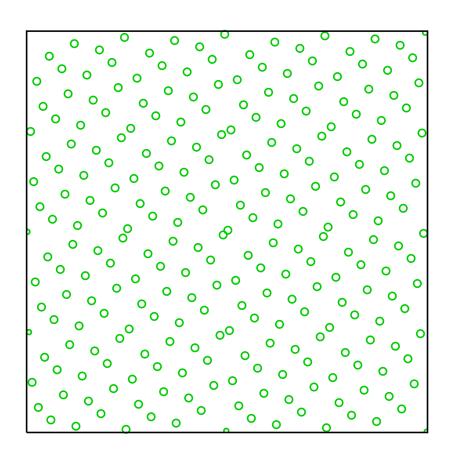






#### Hammersley Issues

- Pattern is well-distributed but...
- 1D projections are regular
  - Too much structure in the pattern
- For a given n only one sequence exists





### Halton Sequence

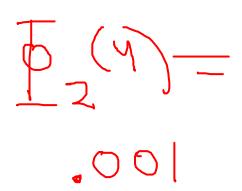
$$p_i = (\Phi_2(i), \Phi_3(i), \Phi_5(i), ...)$$

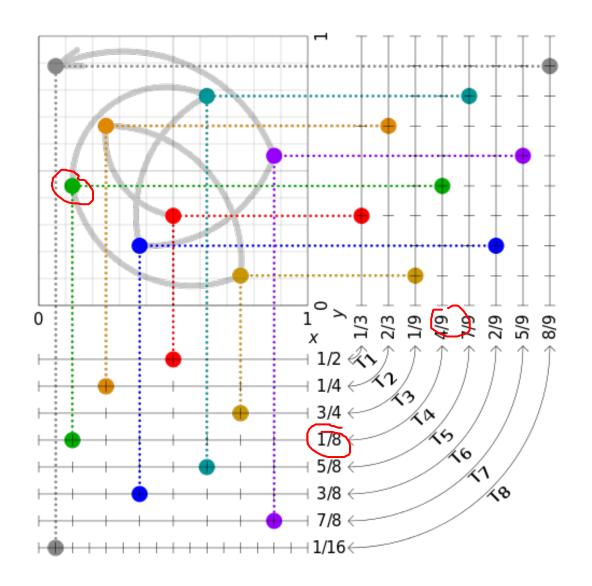
- Better low discrepancy sequence
- Generate n-dimensional points
  - though Hammersley can be generalized as well....
- Number of samples need not be known in advance



# Halton Sequence Example

The 2,3 Halton Sequence







0.11



## Definition: Discrepancy

- We often want a low discrepancy sequence
  - e.g. Monte Carlo methods
- Imagine points in some space S = [0,1]<sup>n</sup>
- Suppose we sample using K points...
- We can evaluate the quality by
  - take portion V of S
  - volume V/volume S should equal (number points in V)/(number points in S)
    - ...but it generally won't
  - the difference is the *discrepancy*
- Different formal ways to measure discrepancy...



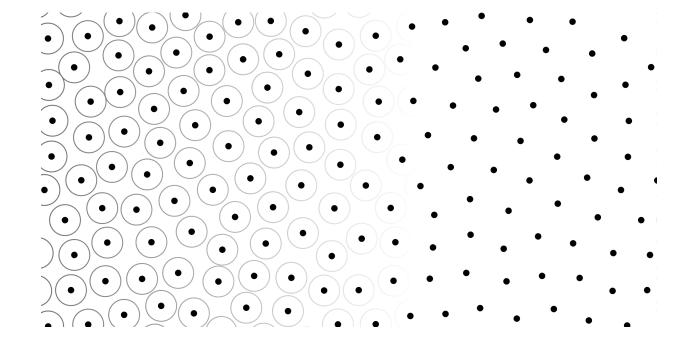


Not actually a low-discrepancy sequence

Does assure a minimum distance between points

Fast and generalizes well to higher dimensions

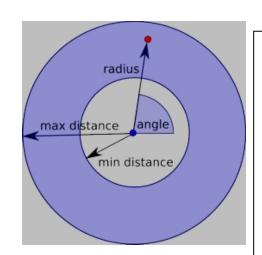
Popular in game industry





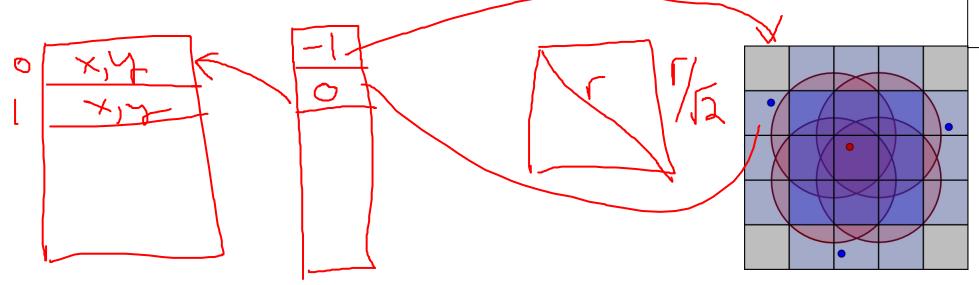
#### Algorithm

**Step 0.** Initialize an n-dimensional background grid for storing samples and accelerating spatial searches. We pick the cell size to be bounded by  $r/\sqrt{n}$ , so that each grid cell will contain at most one sample, and thus the grid can be implemented as a simple n-dimensional array of integers: the default -1 indicates no sample, a non-negative integer gives the index of the sample located in a cell.



r is minimum distance between two sample points

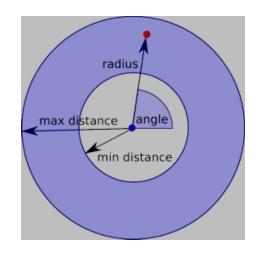
r will be the length of a cell diagonal

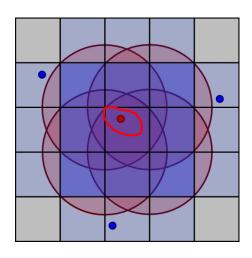




#### Algorithm

**Step 1.** Select the initial sample,  $x_0$ , randomly chosen uniformly from the domain. Insert it into the background grid, and initialize the "active list" (an array of sample indices) with this index (zero).

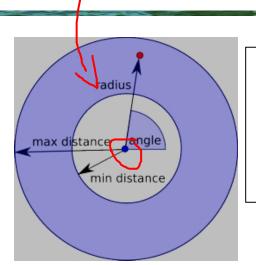






#### Algorithm

**Step 2.** While the active list is not empty, choose a random index from it (say i). Generate up to k points chosen uniformly from the spherical annulus between radius r and 2r around  $x_i$ . For each point in turn, check if it is within distance r of existing samples (using the background grid to only test nearby samples). If a point is adequately far from existing samples, emit it as the next sample and add it to the active list. If after k attempts no such point is found, instead remove i from the active list.



r will be the length of a cell diagonal

k=30 is a popular choice

