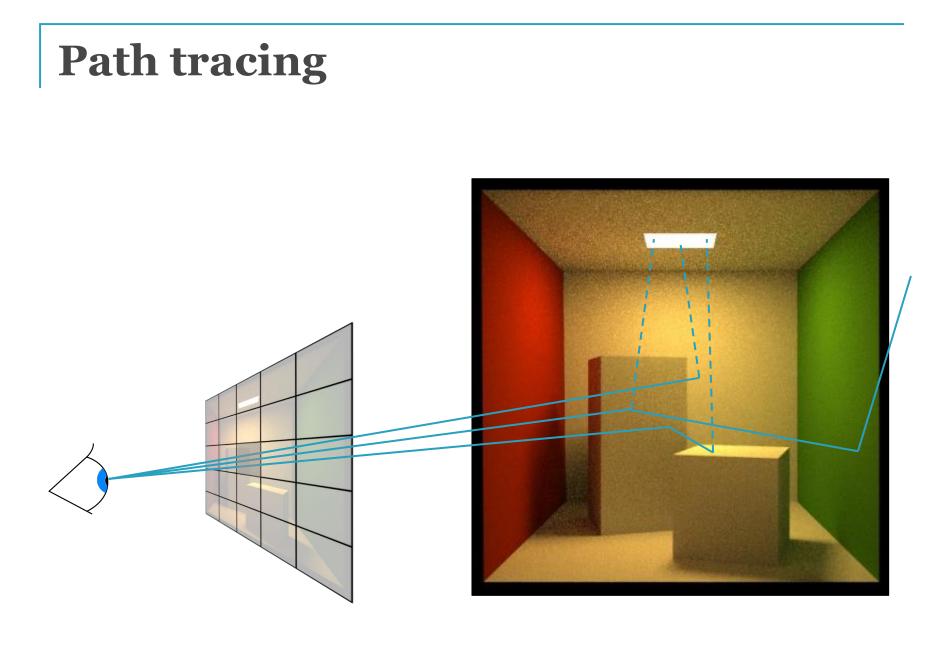
Advanced 3D graphics for movies and games (NPGR010)

Low-discrepancy sequences and quasi-Monte Carlo methods

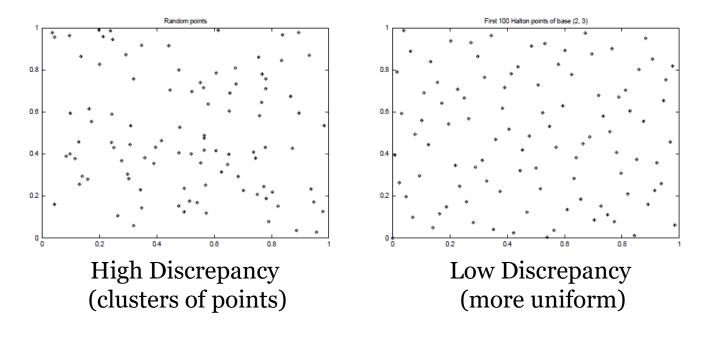
Jiří Vorba, MFF UK/Weta Digital jirka@cgg.mff.cuni.cz

Slides by prof. Jaroslav Křivánek, extended by Jiří Vorba



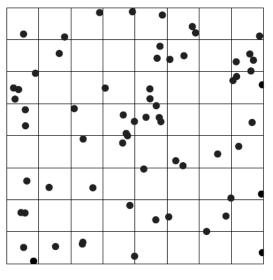
Quasi-Monte Carlo

 Goal: Use point sequences that cover the integration domain as uniformly as possible, while keeping a 'randomized' look of the point set

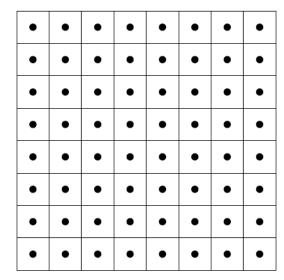


Recall: Stratified samples

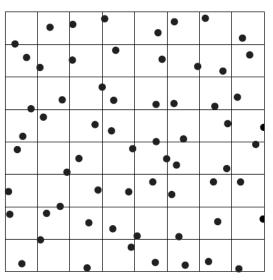
- Samples can still form clumps at borders
- Suffers from course of dimensionality



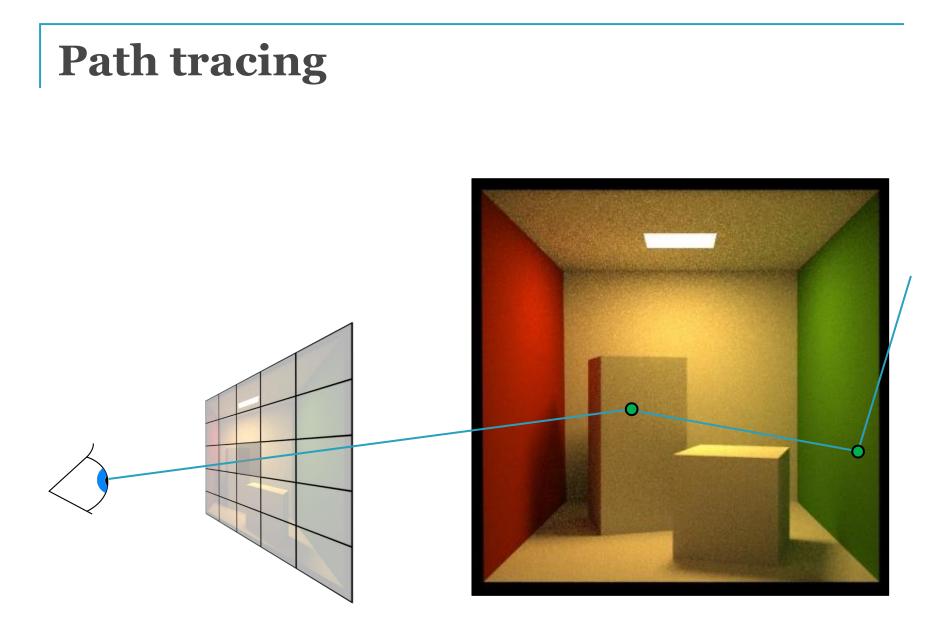
Random



One sample per stratum



Jittered stratified sampling



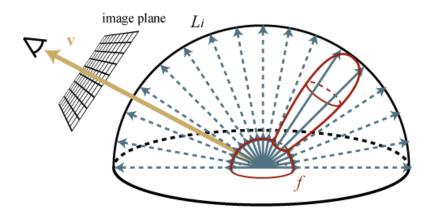
```
estimateLin(x, omegaInAtX) // radiance incident at "x" from the direction "omegaInAtX"
                          // ("omegaInAtX" is pointing *away* from "x")
{
 Spectrum throughput = (1, 1, 1)
 Spectrum accum = (0, 0, 0)
 while(1)
                          // we don't cut off the path length now
   hit = findNearestIntersection(x, omegaInAtX)
   if noIntersection(hit) // ray leaves the scene - it "hits" the background
     return accum + throughput * bkgLight.getLe(x, - omegaInAtX)
   omegaOut := -omegaInAtX // omegaOut at hit.pos
   if isOnLightSource(hit) // ray happened to directly hit a light source
     accum += throughput * getLe(hit.pos, omegaOut)
                                                           // "pick up" emission
    // now estimate the reflected radiance
   [omegaIn, pdfIn] := generateRandomDir(hit)
                                                           // omegaIn at hit.pos
   throughput *= 1/pdfIn * brdf(hit.pos, omegaIn, omegaOut) * dot(hit.n, omegaIn)
```

```
survivalProb = min(1, throughput.maxComponent())
if rand() < survivalProb // Russian Roulette - survive (reflect)
throughput /= survivalProb
x := hit.pos // "recursion"
omegaInAtX := omegaIn // "recursion"
else
break; // terminate path
}
return accum; Advanced 3D Graphics (NPGR010) - J. Vorba 2020</pre>
```

}

BSDF sampling

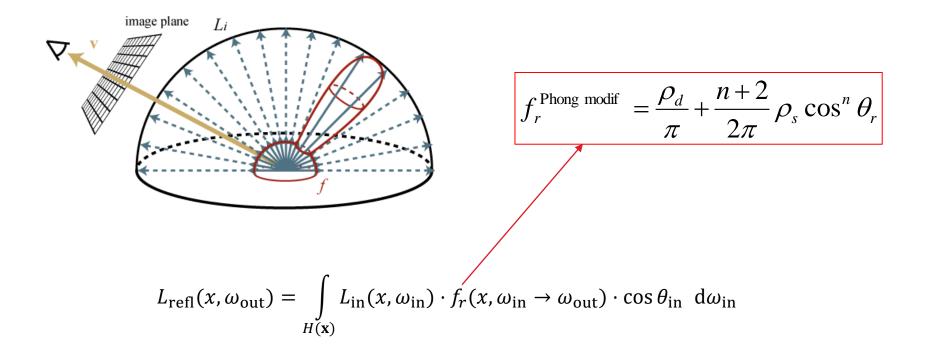
• We need 2 random samples to cover hemisphere



$$L_{\text{refl}}(x,\omega_{\text{out}}) = \int_{H(\mathbf{x})} L_{\text{in}}(x,\omega_{\text{in}}) \cdot f_r(x,\omega_{\text{in}} \to \omega_{\text{out}}) \cdot \cos\theta_{\text{in}} \, \mathrm{d}\omega_{\text{in}}$$

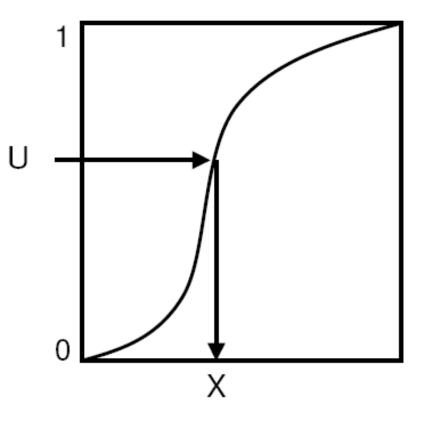
BSDF sampling

- We need 2 random samples to cover hemisphere
- + 1 to choose a component



Transformation method – cdf inversion

 U is uniformly distributed in [0,1]ⁿ



Transformation of point sets

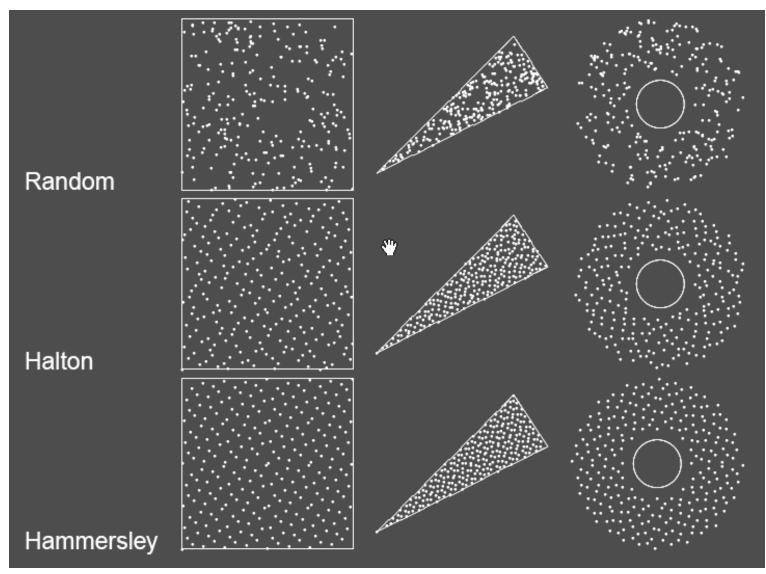
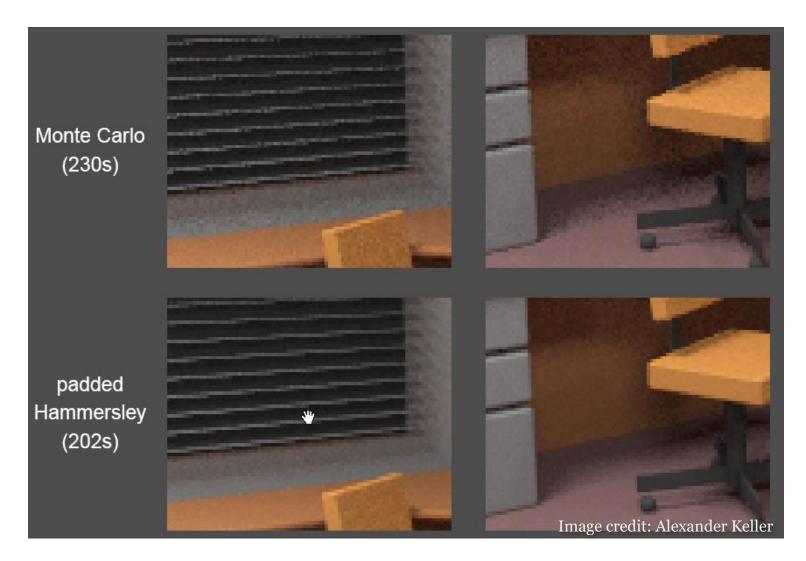


Image credit: Alexander Keller

MC vs. QMC



Random

- Mersenne twister
- Pseudorandom number generator
- Available in C++11
- 32-bit wide (period)

```
#include <random>
const uint32 t seed = 123;
std::mt19937 generator (seed);
std::uniform_real_distribution<float> dis(0.0, 1.0);
float ksi = dis(generator);
```

Large state – 2.5KB

Random - seeding





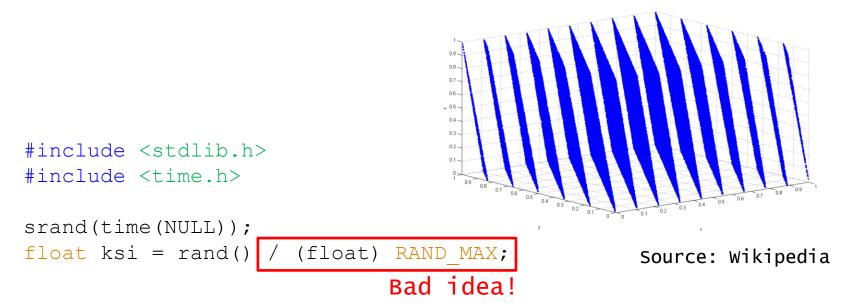
Truly random – seed by current time But we want deterministic renders!

Why?

Debugging! Imagine chasing a source of firefly

Stay away from srand() and rand()

- Bad statistical properties
- Short period (only 16-bit)



More details

https://channel9.msdn.com/Events/GoingNative/2013/rand-Considered-Harmful

Random

- Xorshift (by George Marsaglia)
- Pseudorandom generator
- Fast, tiny state
- **32-bit, (64-bit, 128-bit)**
- Xorshift+ (statistically as good as Mersenne twister)

```
#include <stdint.h>
struct xorshift32_state {
    uint32_t a;
};
/* The state word must be initialized to non-zero */
uint32_t xorshift32(struct xorshift32_state *state)
{
    /* Algorithm "xor" from p. 4 of Marsaglia, "Xorshift RNGs" */
    uint32_t x = state->a;
    x ^= x << 13;
    x ^= x << 5;
    return state->a = x;
}
```

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Source

Quasi Monte Carlo (QMC) methods

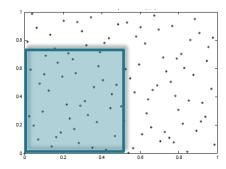
- Use of strictly deterministic sequences instead of random numbers
 - Also true for pseudo-random numbers
- All formulas as in MC, just the underlying proofs cannot reply on the probability theory (nothing is random)
- Based on low-discrepancy sequences

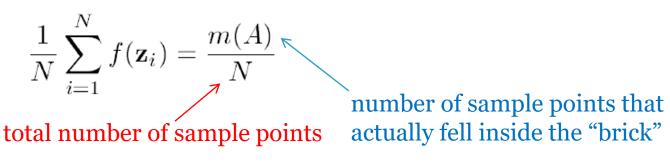
Defining discrepancy

s-dimensional "brick" function:

$$\mathcal{L}(\mathbf{z}) = \begin{cases} 1 \text{ if } 0 \leq \mathbf{z}|_1 \leq v_1, 0 \leq \mathbf{z}|_2 \leq v_2, \dots, 0 \leq \mathbf{z}|_s \leq v_s \\\\ 0 \text{ otherwise.} \end{cases}$$

- True volume of the "brick" function: $V(A) = \prod_{j=1}^{s} v_j$
- MC estimate of the volume of the "brick":





Discrepancy

 Discrepancy (of a point sequence) is the maximum possible error of the MC quadrature of the "brick" function over all possible brick shapes:

$$\mathcal{D}^*(\mathbf{z}_1, \mathbf{z}_2, \dots \mathbf{z}_N) = \sup_A \left| \frac{m(A)}{N} - V(A) \right|.$$

- serves as a measure of the uniformity of a point set
- must converge to zero as N -> infty
- the lower the better (cf. Koksma-Hlawka Inequality)

Koksma-Hlawka inequality

Koksma-Hlawka inequality "variation" of
$$f$$

$$\left| \int_{\mathbf{z} \in [0,1]^s} f(\mathbf{z}) \, d\mathbf{z} - \frac{1}{N} \sum_{i=1}^N f(\mathbf{z}_i) \right| \le \mathcal{V}_{\mathrm{HK}} \cdot \mathcal{D}^*(\mathbf{z}_1, \mathbf{z}_2, \dots, \mathbf{z}_N)$$

the KH inequality only applies to *f* with finite variation
QMC can still be applied even if the variation of f is infinite

i	binary form of i	radical inverse	H_i
1	1	0.1	0.5
2	10	0.01	0.25
3	11	0.11	0.75
4	100	0.001	0.125
5	101	0.101	0.625
6	110	0.011	0.375
7	111	0.111	0.875

- point placed in the middle of the interval
- then the interval is divided in half
- has low-discrepancy

i	binary form of i	radical inverse	H_i
1	1	0.1	0.5
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1

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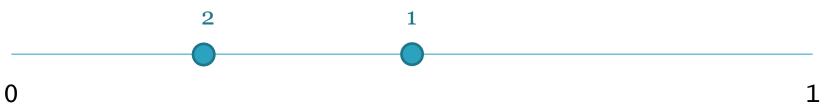
1

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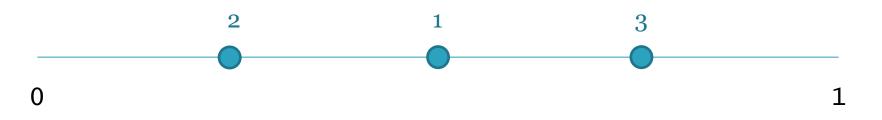
1

0

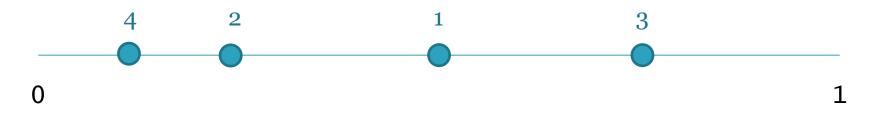
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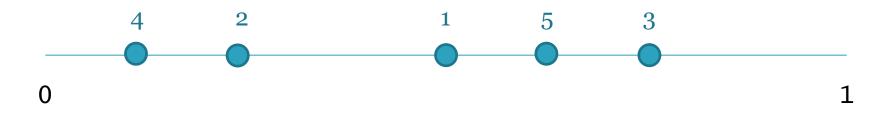
i	binary form of i	radical inverse	H_i
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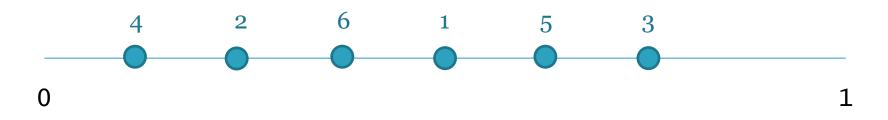
i	binary form of i	radical inverse	H_i
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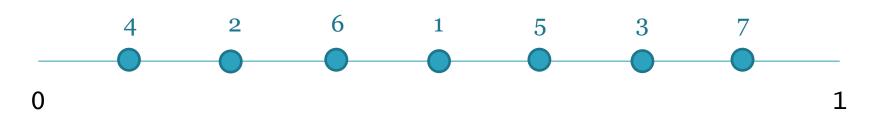
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7	111	0.111	0.875



Van der Corput Sequence

b ... **Base**

radical inverse

$$i = \sum_{j=0}^{\infty} a_j(i)b^j \quad \mapsto \quad \Phi_b(i) := \sum_{j=0}^{\infty} a_j(i)b^{-j-1}$$

Van der Corput Sequence

b ... **Base**

radical inverse

$$i = \sum_{j=0}^{\infty} a_j(i)b^j \quad \mapsto \quad \Phi_b(i) := \sum_{j=0}^{\infty} a_j(i)b^{-j-1}$$

• Example: i = 123 b = 10 $\Phi_{10}(123) = \frac{3}{10} + \frac{2}{10^2} + \frac{1}{10^3} = 0.321$

```
double radicalInverse(const int base, int i)
{
    double digit, radical;
    digit = radical = 1.0 / (double)base;
    double inverse = 0.0;
    while(i)
    {
        inverse += digit * (double)(i % base);
        digit *= radical;
        i /= base;
    }
    return inverse;
}
```

Discrepancy of sequence P

$$D_N^*(P) = O\left(\frac{\log N}{N}\right)$$

Sequences in higher dimension

Halton sequence $x_i := (\Phi_{b_1}(i), \dots, \Phi_{b_s}(i))$ where b_i is the *i*-th prime number

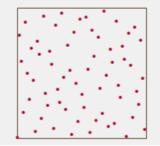
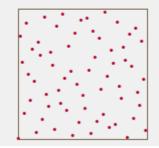


Image credit: Alexander Keller

Sequences in higher dimension

Halton sequence $x_i := (\Phi_{b_1}(i), \dots, \Phi_{b_s}(i))$ where b_i is the *i*-th prime number



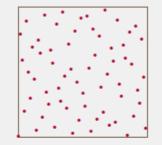
Discrepancy

$$D_N^*(x_i) = O\left(\frac{(\log N)^s}{N}\right)$$

Image credit: Alexander Keller

Sequences in higher dimension

Halton sequence $x_i := (\Phi_{b_1}(i), \dots, \Phi_{b_s}(i))$ where b_i is the *i*-th prime number



Hammersley point set $x_i := \left(\frac{i}{n}, \Phi_{b_1}(i), \dots, \Phi_{b_{s-1}}(i)\right)$

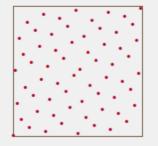


Image credit: Alexander Keller

Sequences vs sets

Set

- Number of samples need to be known a-priory
- Hammersley, stratified samples
- Usually needs to be recomputed if we change N

Sequence

- We don't need to know number of samples beforehand
- Points added without recomputing of previous points
- Halton

Progressive sequences/sets

- Any prefix preserves low-discrepancy
- Suitable for progressive rendering



Figure 2: Penumbra region with 100 samples per pixel. Left: nonprogressive sample set. Right: progressive sample sequence.

Quasi-Monte Carlo (QMC) Methods

- Disadvantages of QMC:
 - Regular patterns can appear in the images (instead of the more acceptable noise in purely random MC)
 - Random scrambling can be used to suppress it



(a) Image rendered with 512 random samples per pixel



(c) Aliasing artefacts from using the same 4 samples for each pixel



(b) Noise artefacts from using 4 samples per pixel with a random sampler

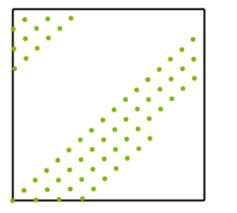


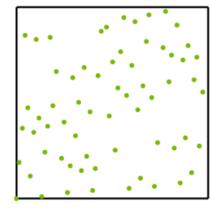
(d) Aliasing artefacts from using 4 samples per pixel with a structured sampler

Scrambling

- Low-dimensional projections show visible patterns
- Mitigated by scrambling

 \rightarrow

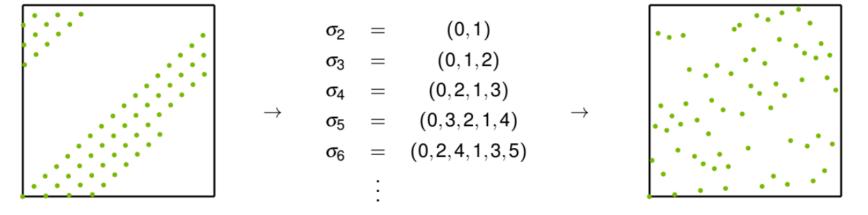




 $(\Phi_{17}(i), \Phi_{19}(i)), i = 0 \dots 63$

Scrambling - permutations

- Low-dimensional projections show visible patterns
- Mitigated by scrambling

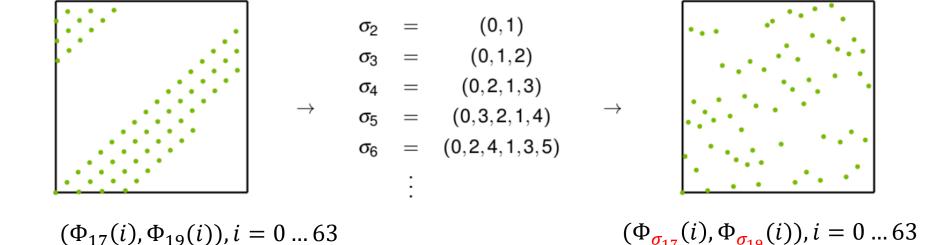


 $(\Phi_{17}(i), \Phi_{19}(i)), i = 0 \dots 63$

Scrambling - permutations

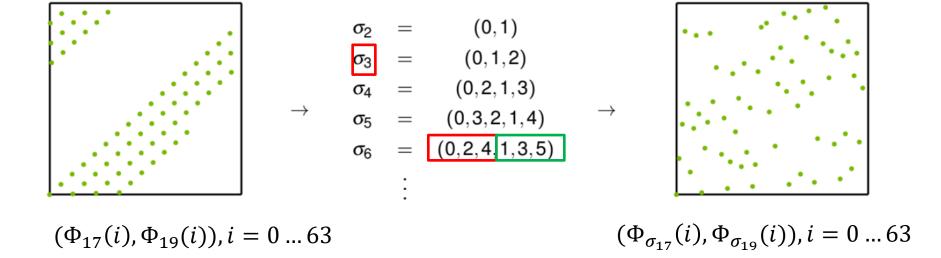
Apply the same permutation at every digit

$$\Phi_{\sigma_b}\left(i\right) = \sum_{j=0}^{\infty} \sigma_b\left(a_j(i)\right) b^{-j-1}$$



Scrambling – permutations by Faure

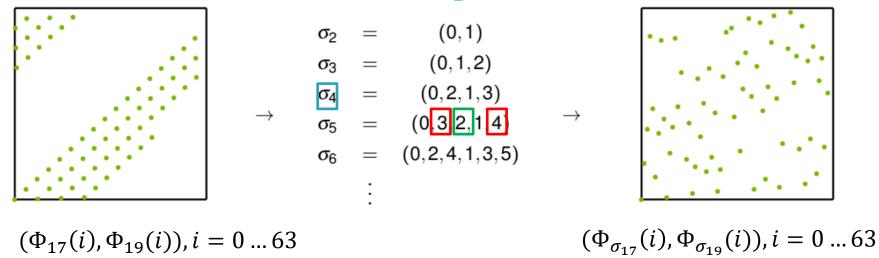
- In general, permutations for each base can be arbitrary
- Deterministic perms. by Faure
- When b is even: Take $2\sigma_{\frac{b}{2}}$ and append $2\sigma_{\frac{b}{2}} + 1$



Scrambling – permutations by Faure

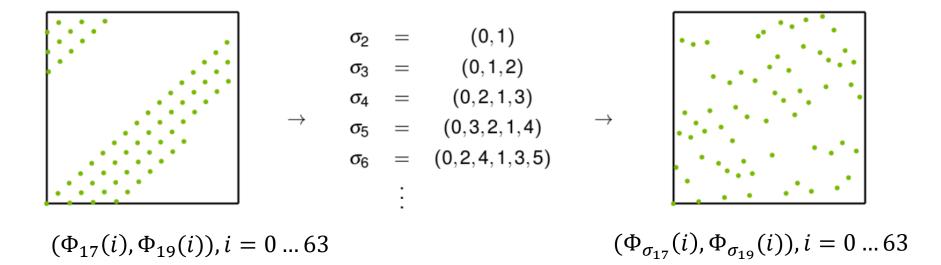
- In general, permutations for each base can be arbitrary
- Deterministic perms. by Faure

When b is odd: Take σ_{b-1} , increment each value $\geq \frac{b-1}{2}$, insert $\frac{b-1}{2}$ in the middle



Scrambling – permutations by Faure

Faure permutations can be implemented efficiently without branching



Use in path tracing

• **Objective**: Generated paths should cover the entire high-dimensional path space uniformly

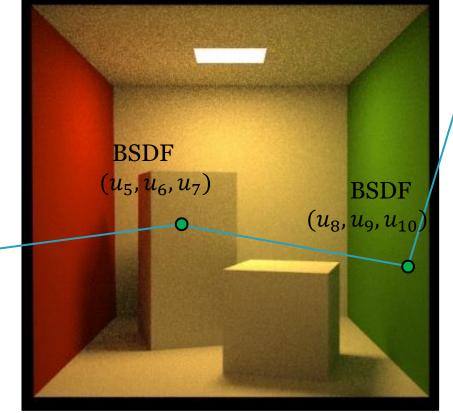
• Approach:

- Paths are interpreted as "points" in a high-dimensional path space
- Each path is defined by a long vector of "random numbers"
 - Subsequent random events along a single path use subsequent components of the same vector
- Only when tracing the next path, we switch to a brand new "random vector" (e.g. next vector from a Halton sequence)

Path tracing

Path "i" sampled based on the "i-th" sample from the sequence

 $U_i = (u_0, \dots, u_{10})$ **BSDF** (u_5, u_6, u_7) Time, lens, pixel $(u_0, u_1, u_2, u_3, u_4)$



Progressive rendering with Halton

• Option 1:

- Halton sequence per pixel, use different scrambling
- Loose low-discrepancy properties in the image plane
- □ In fact results into stratified sampling in the image plane

• Option 2:

- Use nearest power of two to your resolution
- It is possible to compute index of n-th sample in the Halton sequence given the pixels coordinates
- Skip samples falling outside your actual pixels
- Details: PBRT, 3rd edition Chapter 7.4.2.

Further study material

- (t,s)-sequences
 - (0,2)- sequence: Sobol sequence generator matrices
- Rank-1 lattice sequences
- Multi-jittered samples
- References
 - <u>https://sites.google.com/view/myfavoritesamples</u>
 - <u>Christiane Lemieux: Monte Carlo and Quasi-Monte Carlo</u> <u>Sampling [2008]</u>