

# Anti-aliasing and sampling

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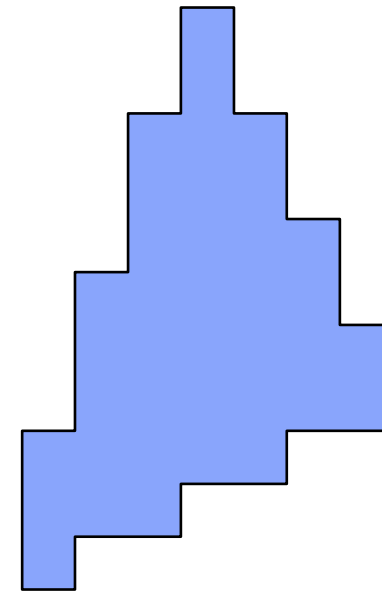
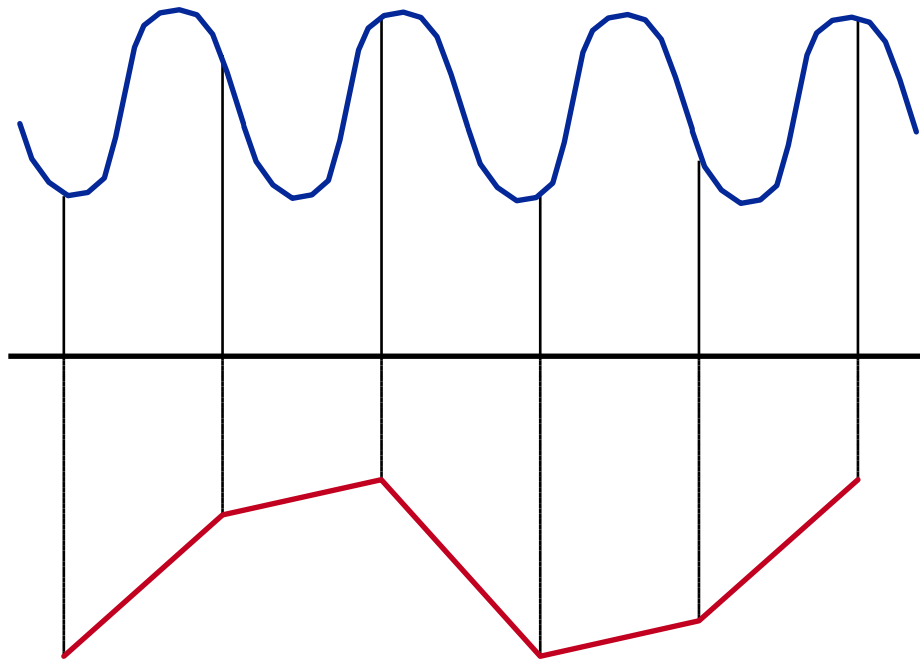
[pepca@cgg.mff.cuni.cz](mailto:pepca@cgg.mff.cuni.cz)

<http://cgg.mff.cuni.cz/~pepca/>



# Spatial and temporal alias

Artefacts caused by a regular-grid  
or regular sampling

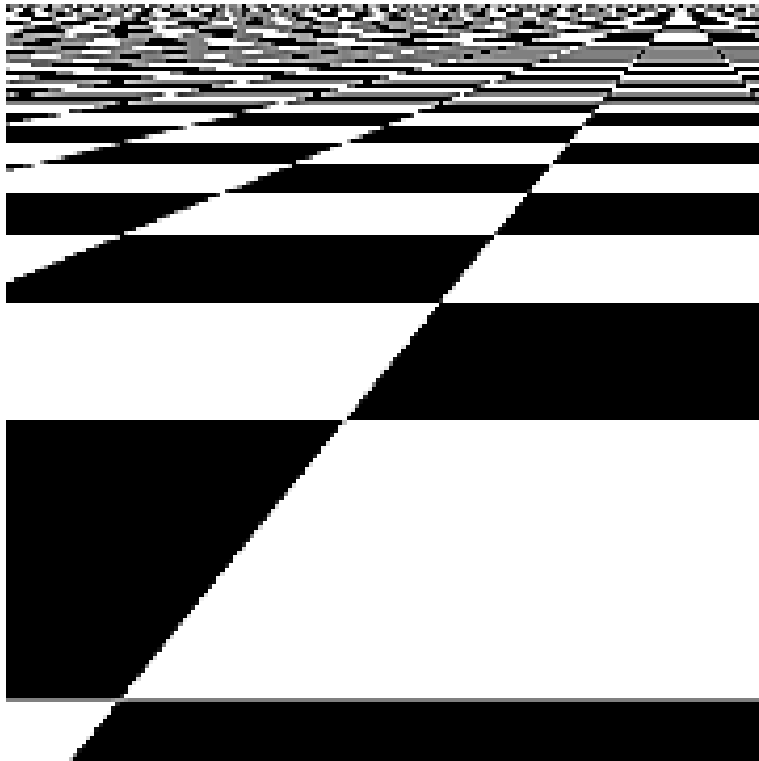


# Spatial alias

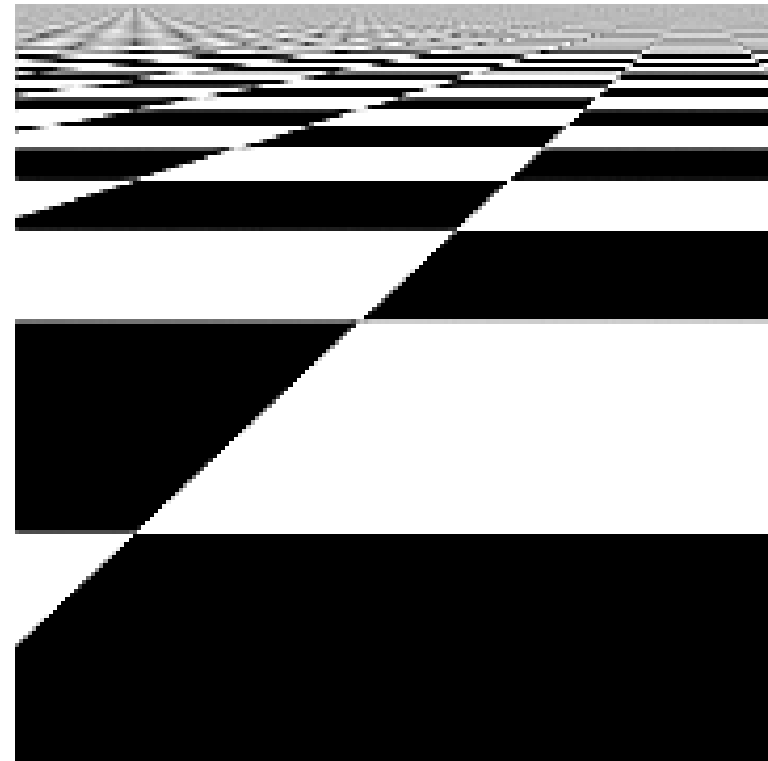


- **jagged** oblique lines
  - regular dense system of lines or stripes on a texture can lead to “Moiré effect”
- **interference** of fast periodic image changes with a pixel raster
  - example: picket fence in perspective projection
  - too fine or too distant regular texture (checkerboard viewed from distance)

# Spatial alias – checkerboard



1 sample per pixel



256 spp (jittering)



# Temporal alias

- ◆ shows in **slow motion animation**
- **blinking pixels** on contours of moving objects
  - the whole small objects can blink
- **interference** of a periodic movement with a frame frequency
  - spinning wheel seems to be still or even rotating the other direction

# Real world

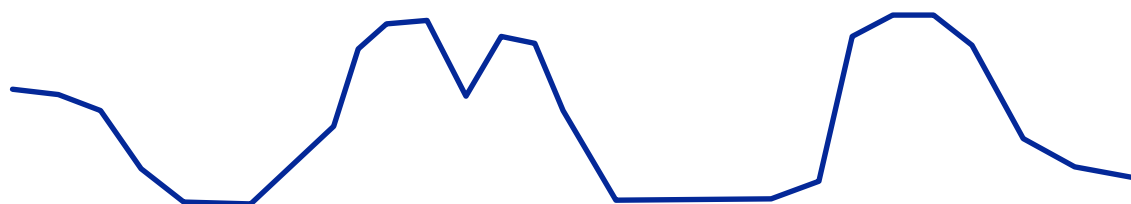


Human visual system has no alias.

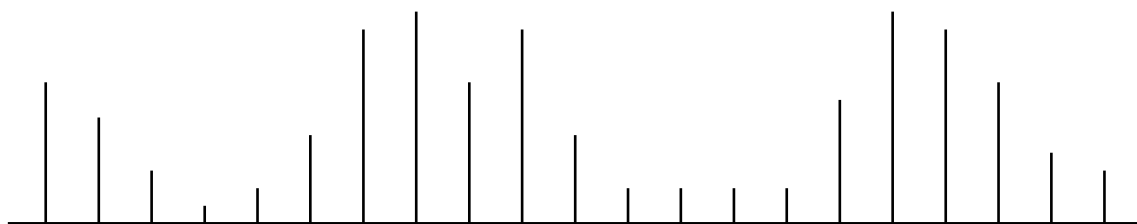
Alias manifests only mildly in photography.

- objects smaller than **resolution of a sensor** are without details / blurry
  - fence from large distance is perceived as an average-color area (mix of background + foreground colors)
- too fast movement generates **fuzzy** (blurred) **perception**

# Reconstruction in raster context

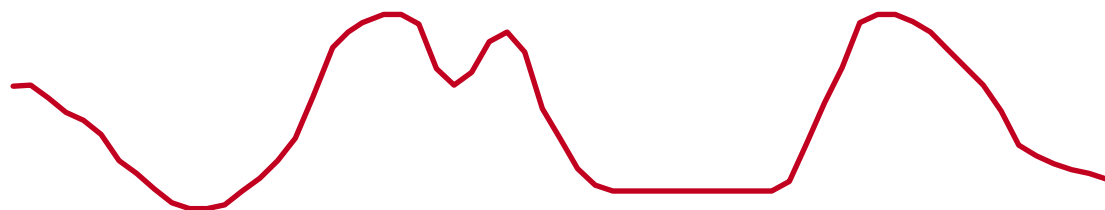
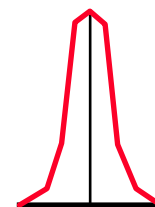


**original  
(image function)**



**sampling  
(computation)**

**reconstruction filter:**



**reconstruction  
(rendering)**

# Sampling and reconstruction



- **image sampling** or **computing** of image function
  - higher frequencies should be reduced/removed from an image before sampling
  - low pass filter (convolution – window averaging)
  - image synthesis can reduce higher frequencies directly (anti-aliasing by pixel supersampling)
- **reconstruction filter** is defined by an output device
  - e.g. neighbour CRT monitor pixels overlap
  - LCD pixels behave differently





# Anti-aliasing by supersampling

**Image function with continuous domain and unlimited spectrum:**

$$\underline{f(\mathbf{x}, \mathbf{y})}$$

**Anti-aliasing filter** (function with limited support):

$$\underline{h(\mathbf{x}, \mathbf{y})}$$

**Pixel color [i,j]:**

$$I(i, j) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f(\mathbf{x}, \mathbf{y}) \cdot h(\mathbf{x} - \mathbf{i}, \mathbf{y} - \mathbf{j}) \, dx \, dy$$



# Simple variant

Assuming **box smoothing filter** and **unit square pixel**:

$$I(i, j) = \int_j^{j+1} \int_i^{i+1} f(x, y) \, dx \, dy$$

(integral average value of the image function on the pixel area)



# Quadrature

- 1 **analytic** (close form solution)
  - in rare cases (simple image function)
- 2 **numeric solution** using sampling
  - finite set of samples  $[\mathbf{x}_i, \mathbf{y}_i]$
  - integral estimate by the sum

$$I(\mathbf{i}, \mathbf{j}) = \frac{\sum_{\mathbf{k}} \mathbf{f}(\mathbf{x}_{\mathbf{k}}, \mathbf{y}_{\mathbf{k}}) \cdot \mathbf{h}(\mathbf{x}_{\mathbf{k}}, \mathbf{y}_{\mathbf{k}})}{\sum_{\mathbf{k}} \mathbf{h}(\mathbf{x}_{\mathbf{k}}, \mathbf{y}_{\mathbf{k}})}$$

- stochastic sampling – **Monte-Carlo method**

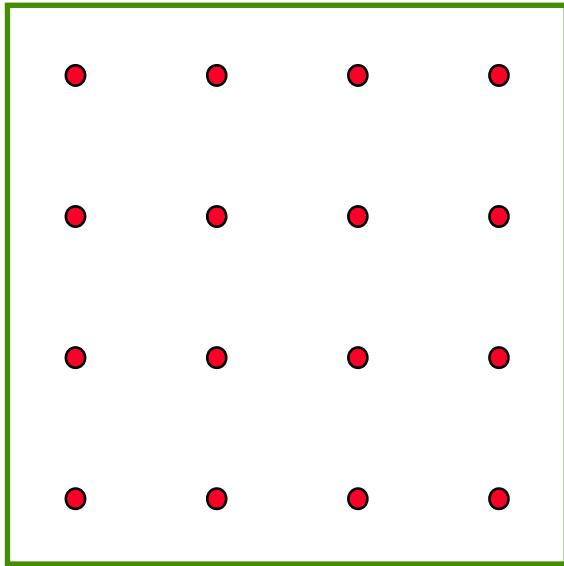


# Sampling methods

- ◆ **mapping:**  $\mathbf{k} \rightarrow [\mathbf{x}_k, \mathbf{y}_k]$ 
  - sample is selected from the given 2D region (domain): usually rectangular, square or circular
  - sampling in higher dimensions (dim=10)
- ➔ required **properties** of sampling algorithms
  - uniform probability over a domain
  - high regularity is not desirable (interference)
  - efficient computation



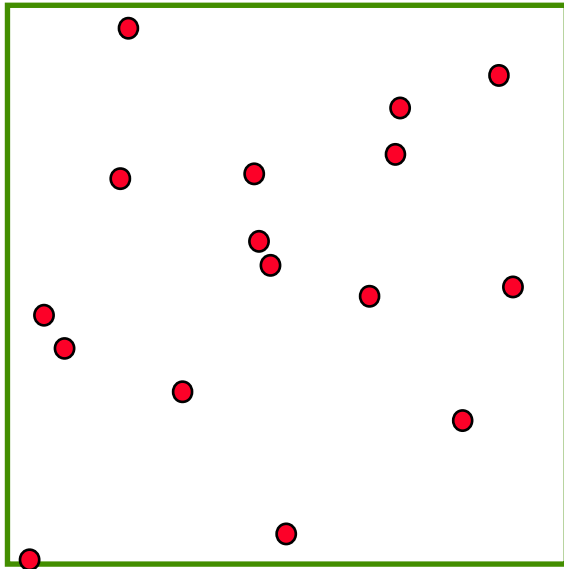
# Uniform sampling



Does not reduce **Moire / interference**  
(interference still appears in higher frequencies)



# Random sampling

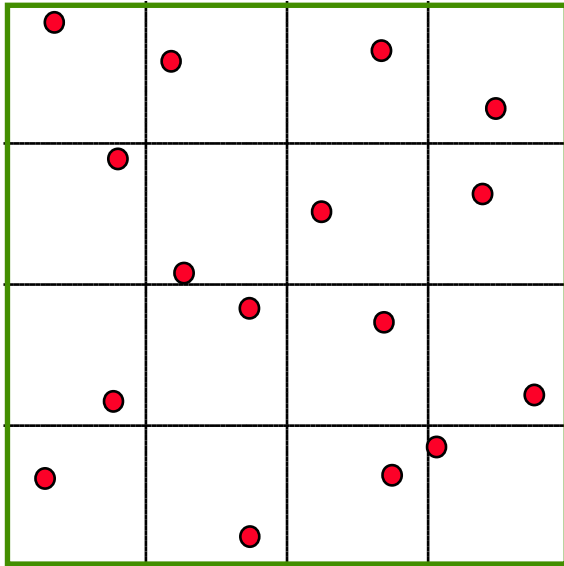


N independent random samples with uniform probability density (PDF)

Samples tend to form **clusters**

**Too much noise** in a result image

# “Jittering”

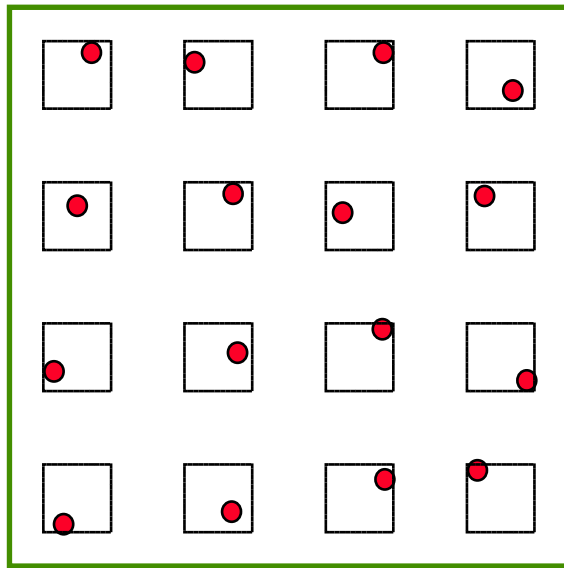


$K \times K$  independent random samples in  $K \times K$  equally sized sub-regions covering the original domain completely

Big clusters are not possible

**More regular coverage** of a domain

# Partial “jittering”, “semi-jittering”

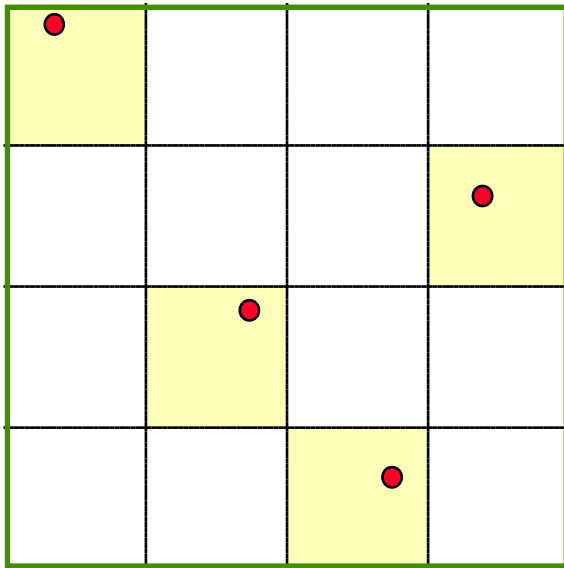


$K \times K$  independent random samples in  $K \times K$  equally sized sub-regions not covering the original domain

Clusters are impossible  
**Too** regular (interference)



# “N rooks” (“uncorrelated jitter”)

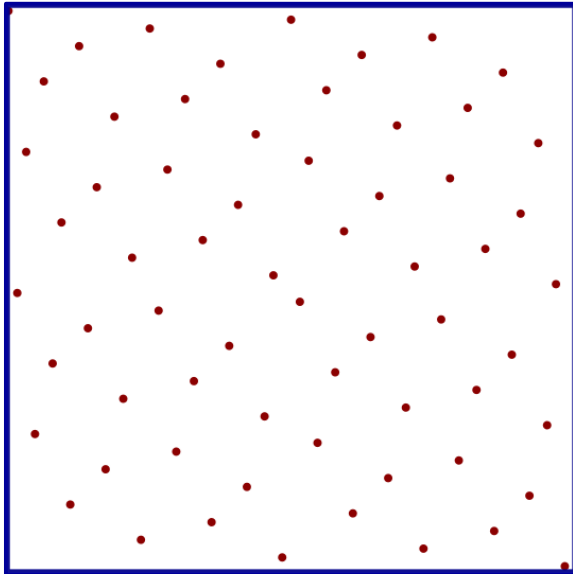


## “Low-cost jittering”:

there is exactly one sample in each row and in each column. Random permutation of a diagonal

Good properties of “jittering” are preserved  
Higher **efficiency** (especially in high dimensions)

# Hammersley



- + **excellent discrepancy**
- + **deterministic**
- + **very fast**
- difficult adaptive refinement
- bad spatial spectrum

Famous **Halton sequence** is based on similar principles..



# Deterministic sequences

- ◆ based on similar principles:
  - Halton, Hammersley, Larcher-Pillichshammer
- ◆ for a prime number  $\mathbf{b}$  let  $\mathbf{n}$  be positive integer expressed using  $\mathbf{b}$ -representation:

$$n = \sum_{k=0}^{L-1} d_k(n) b^k$$

- ◆ then there is a number from  $[0,1)$  range:

$$g_b(n) = \sum_{k=0}^{L-1} d_k(n) b^{-k-1}$$



# Halton, Hammersley

- ◆ famous Halton sequence (e.g.  $b_1=2, b_2=3$ ):

$$x(n) = \left[ g_{b_1}(n), g_{b_2}(n) \right]$$

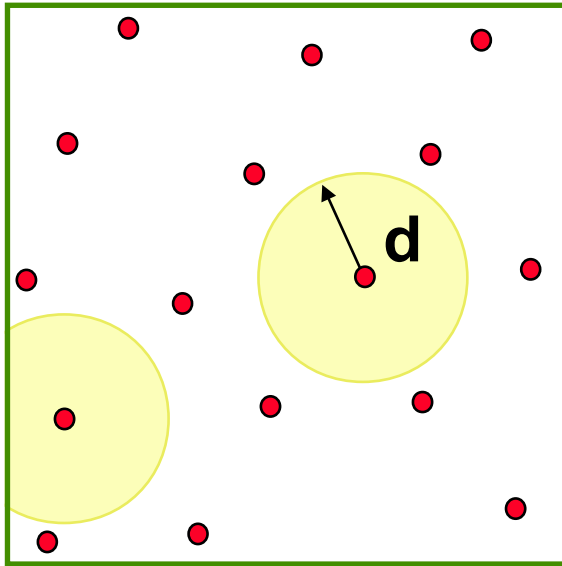
- ◆ Hammersley sequence (e.g.  $b=2$ ):

$$x(n) = \left[ \frac{n}{N}, g_b(n) \right]$$

- ◆ Larcher-Pillichshammer sequence uses **XOR** operation instead of addition (inside the  $g_b(n)$  )..



# Poisson disk sampling



**N random samples**

meeting condition:

$$| [x_k, y_k] - [x_l, y_l] | > d$$

for given value  $d$

Prevents creating **clusters**, imitates **distribution of light-perception cells** in retina of a mammal.  
Difficult efficient **implementation!**



# Implementation

- ◆ **rejection sampling**
  - candidate sample is rejected if too close to any previous accepted sample
  - less efficient for higher number of samples
- ➔ **choice of value  $d$  is problematic**
  - maximum number of placeable samples depends on  $d$
- ➔ **difficult (adaptive) refinement**
  - additional samples to a existing set of samples



# Mitchell's algorithm

(“best candidate” algorithm)

- ◆ generates **gradually refined sample set** (from Poisson sampling)
  - no problems with **d**
  - intrinsic refinement
- ◆ **compute-intensive** algorithm
  - sample set can be **precomputed and reused**
  - to **reduce dependency** between neighbour pixels random rotation and translation can be used

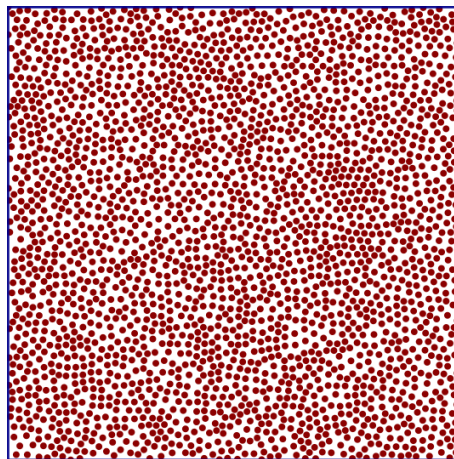
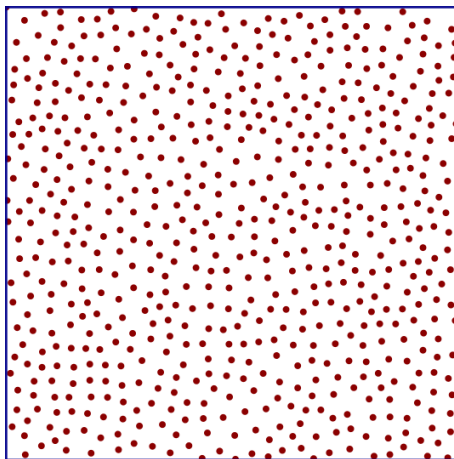
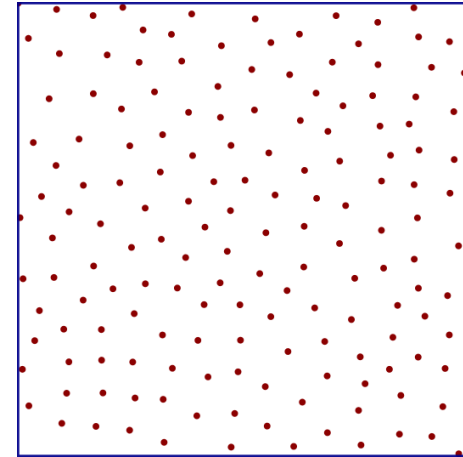
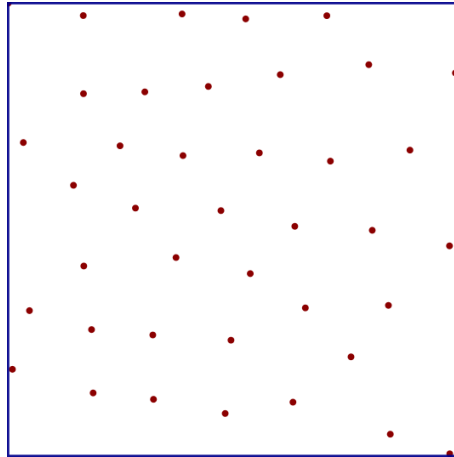
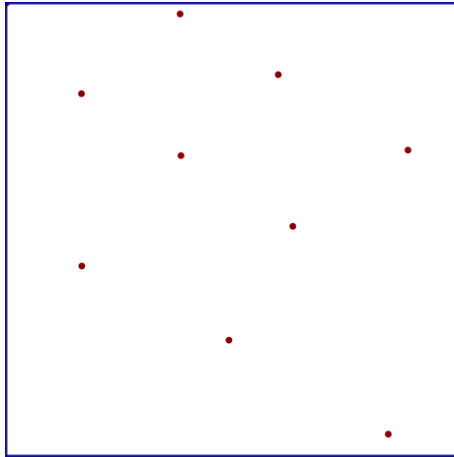


# Mitchell's algorithm

- ① the **1<sup>st</sup> sample** is chosen randomly
- ② choice of the **(k+1)<sup>th</sup> sample**:
  - generate **k · q** independent candidates (**q** determines sample-set quality)
  - the **most distant candidate** (from all previous **k** accepted samples) is selected and accepted
- ➔ for higher **q** we get better quality set
  - choose **q > 10** in demanding situations



# Incremental example



Sample numbers:

10, 40, 160,  
640, 2560

$K = 10$



# Adaptive refinement

- ◆ sampling based on **local importance** (importance sampling) or **interest**
  - some regions have higher weight (higher probability)
  - regions with **higher variance** should be sampled more densely
- ➔ “importance” or “interest” **need not be known in advance** (explicitly)
  - algorithm has to adapt to intermediate results (adaptability)

# Modification of static methods



## ① initial phase:

- compute small set of test samples (1-5)
- define **refinement criterion** based on previous samples

## ② refinement phase:

- sampling is refined in regions of higher need (criterion)
- efficiency: if we can reuse all generated samples

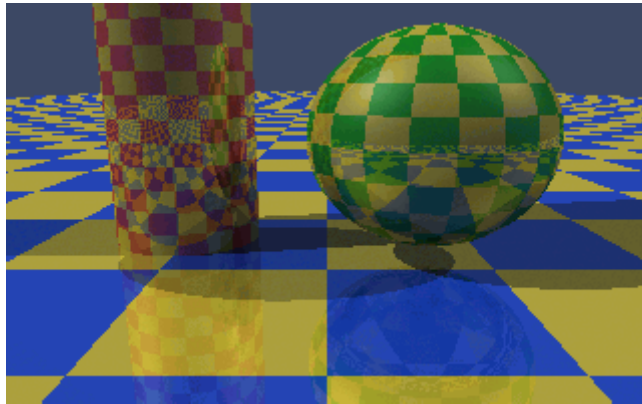
➔ almost every sampling algorithm can be **reformulated** in that way



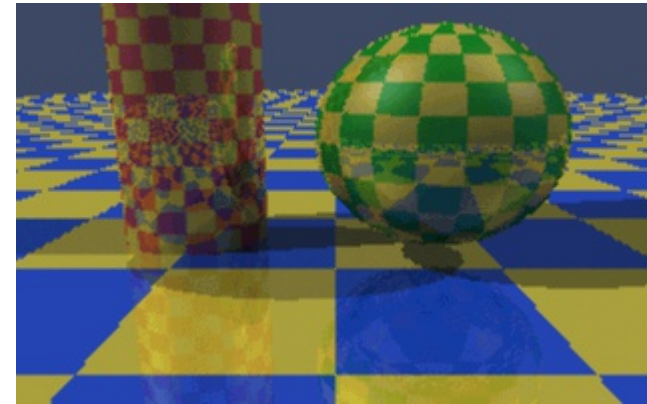
# Refinement criteria

- **function values** (difference, variance, gradient)
  - difference between neighbour samples, ..
- **Id's of hit solids** (Ray-tracing specific)
  - higher priority
  - textures with repeated patterns – use of signatures
- **trace tree** (recursive ray-tracing)
  - topologic comparison of complete or limited trace trees
  - **tree identifier** – recursive hash function using solid Id's, texture signatures, shadow/light, ..

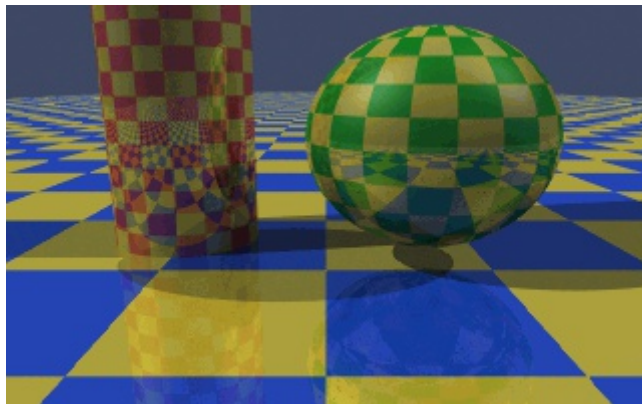
# Adaptive resampling example



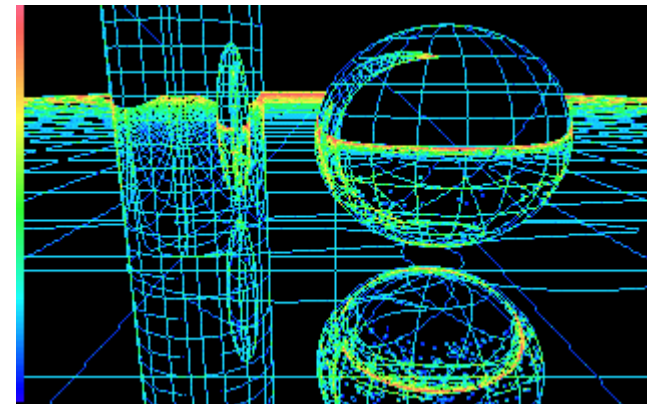
1 spp



1/2 spp

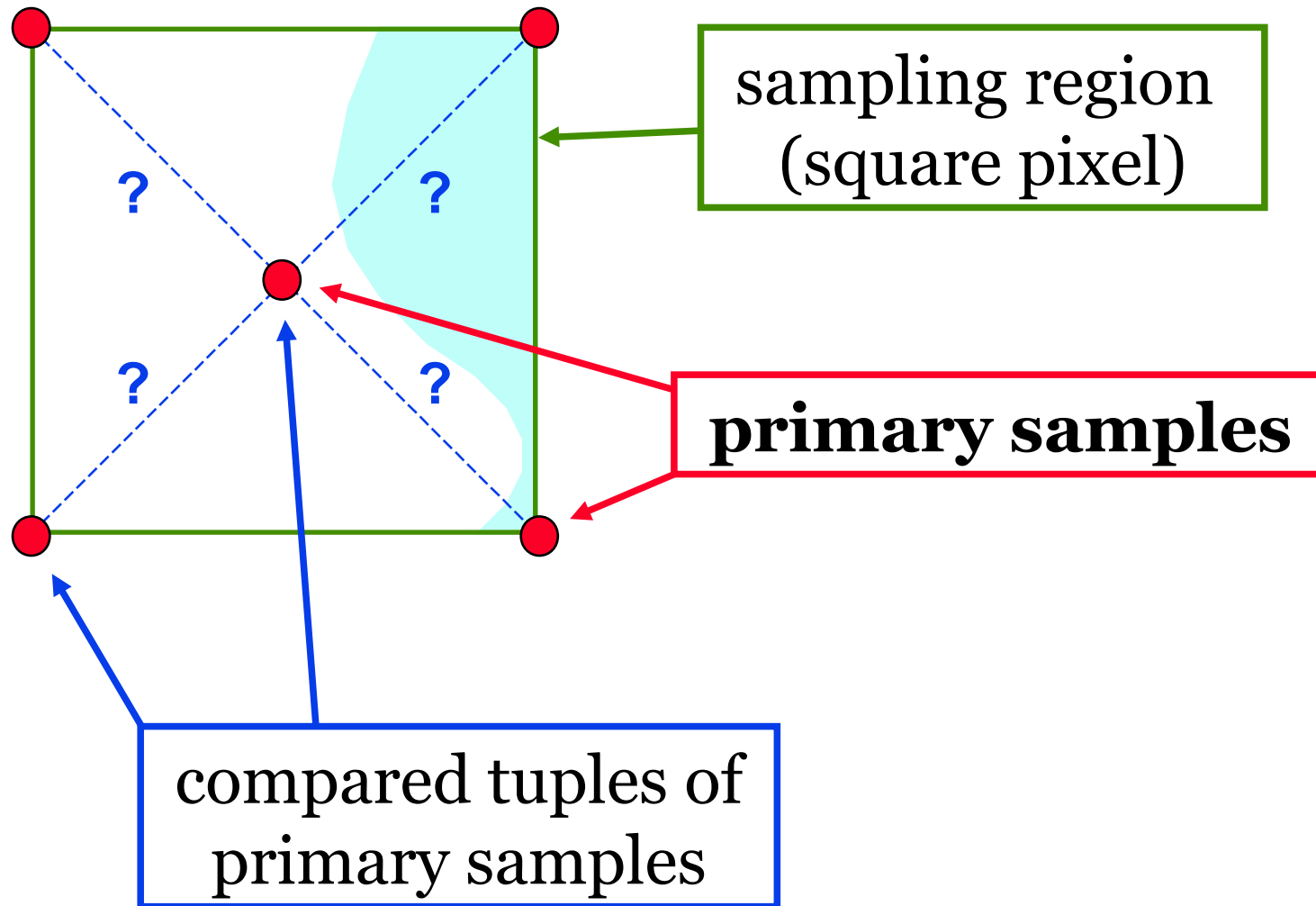


adaptive



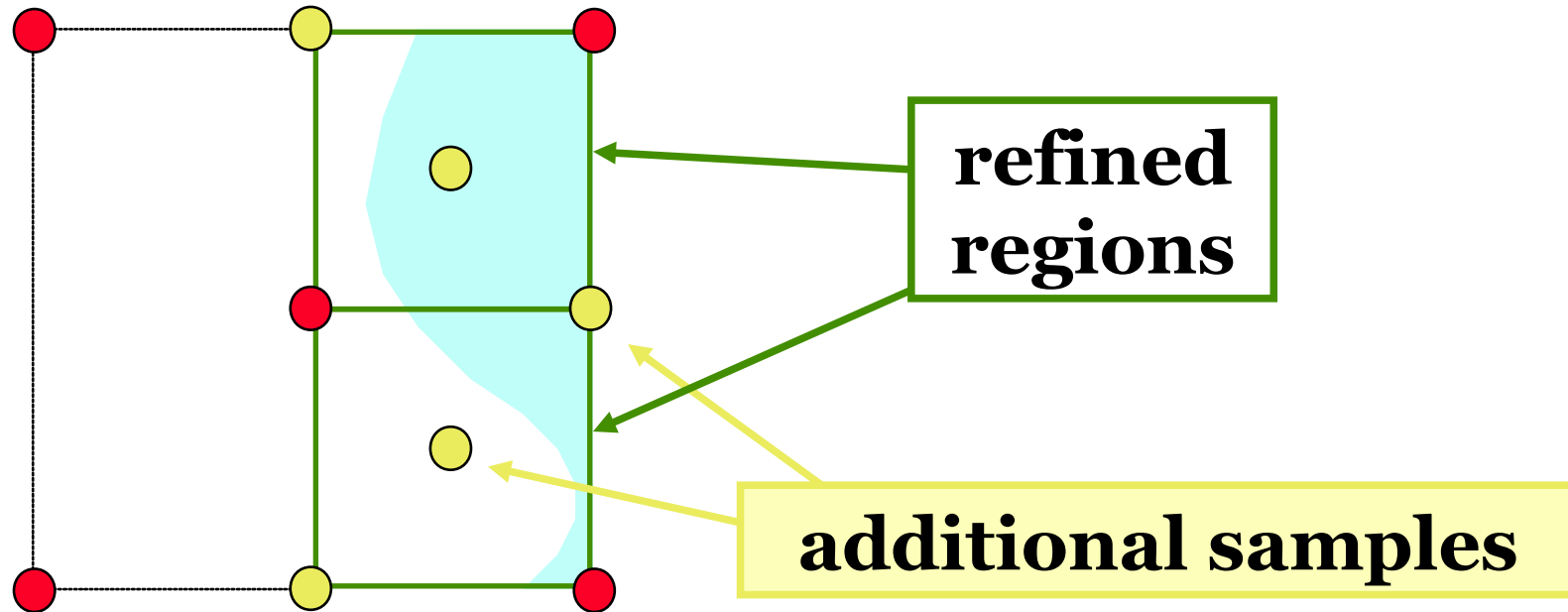
refinement map

# Recursive refinement (Whitted)





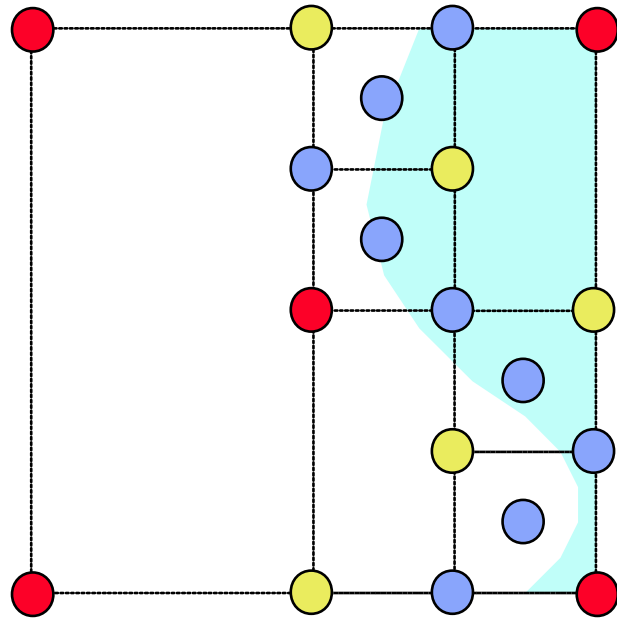
# Refinement phase



The same procedure is executed in refined regions **recursively** (up to the declared maximum level)



# Result sample set



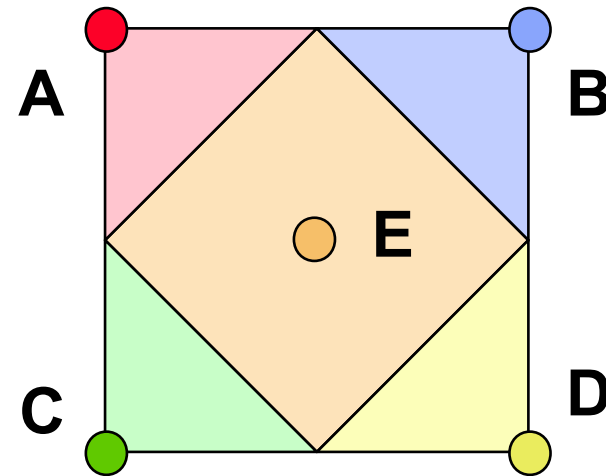
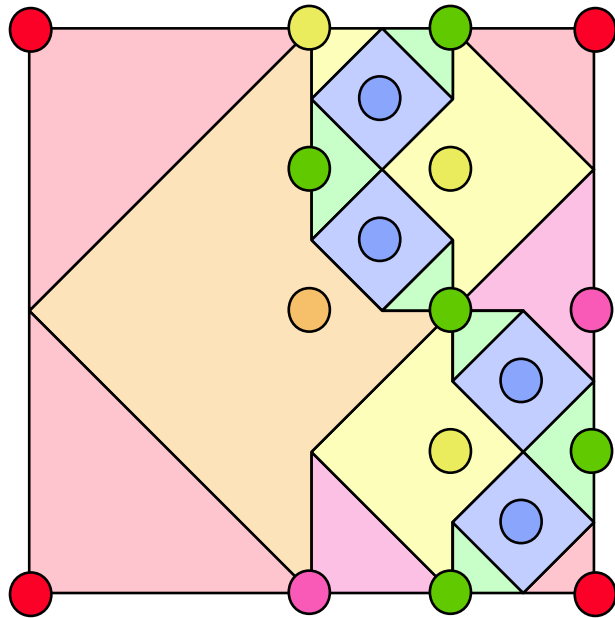
- **phase I**
- **phase II**
- **phase III**

Evaluated:  $5+5+9 = 19$  **samples**  
(from total number of **41**)





# Pixel value reconstruction



$$\frac{1}{2} \mathbf{E} + \frac{1}{8} [\mathbf{A} + \mathbf{B} + \mathbf{C} + \mathbf{D}]$$

If the refinement stops in a specific square,  
its area is split to two triangles (diagonal samples)

# References



- **A. Glassner: *An Introduction to Ray Tracing***, Academic Press, London 1989, 161-171
- **A. Glassner: *Principles of Digital Image Synthesis***, Morgan Kaufmann, 1995, 299-540
- **J. Pelikán: *Náhodné rozmístování bodů v rovině (Random point placement)***, CSGG 2014, slides & paper available online