

Statistical Methods in Particle Physics

4. Monte Carlo Methods

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Klaus Reygers, Martin Völkl (lectures)
Ulrich Schmidt, (tutorials)

Monte Carlo method

- Any method which solves a problem by generating suitable random numbers
- Useful for obtaining numerical solutions to problems which are too complicated to solve analytically
- The most common application of the Monte Carlo method is Monte Carlo integration

- Pioneers

- ▶ Enrico Fermi
- ▶ Stanislaw Ulam
- ▶ John von Neumann
- ▶ Nicholas Metropolis

<https://en.wikipedia.org>



Enrico Fermi



Stanislaw Ulam



J. von Neumann



N. Metropolis

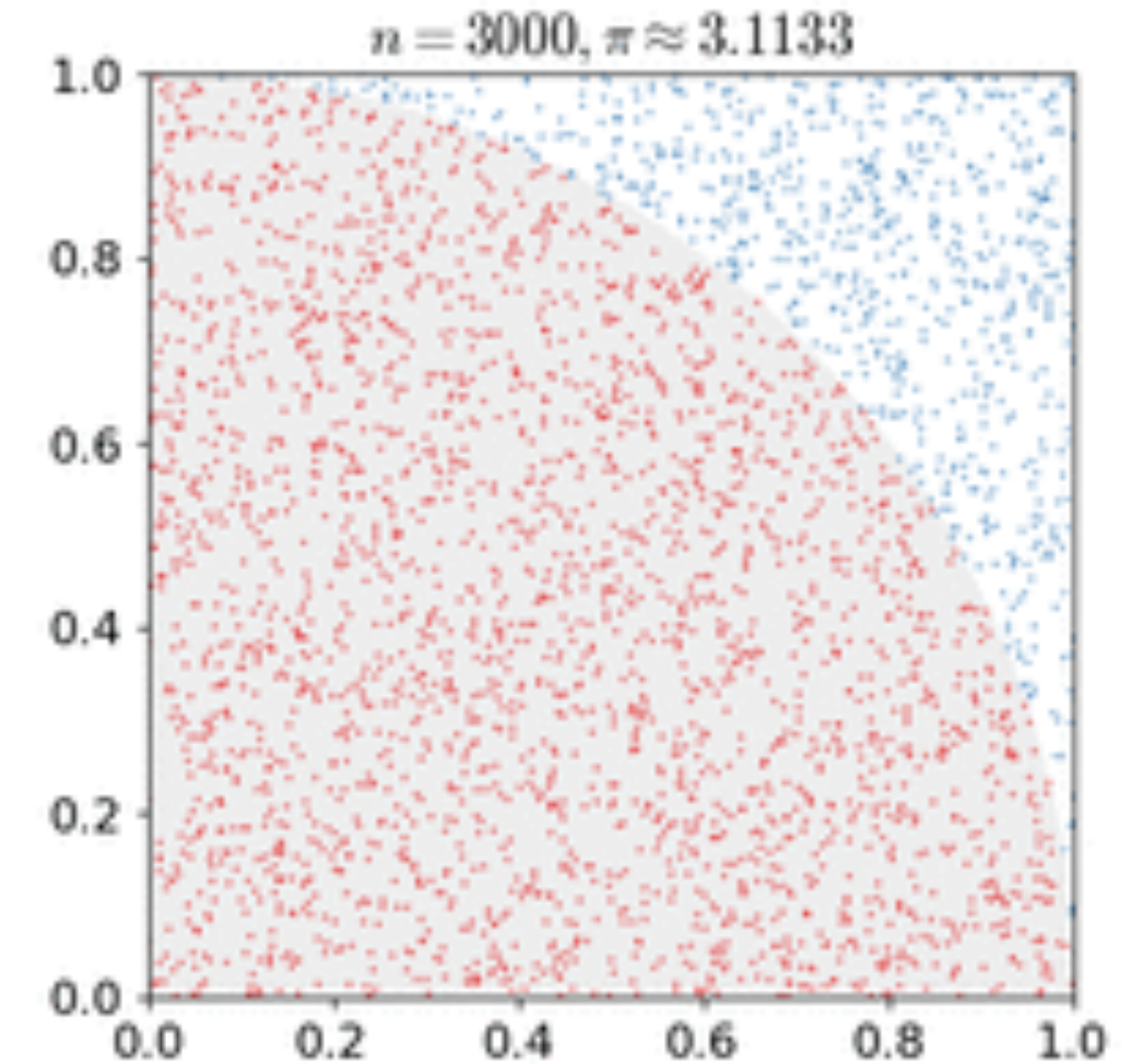
<http://mathworld.wolfram.com/MonteCarloMethod.html>

Monte Carlo method: Examples

[from Bohm, Zech: Introduction to Statistics and Data Analysis for Physicists]

- Area of a circle
- Volume of the intersection of a cone and a torus
 - ▶ Hard to solve analytically
 - ▶ Easy to solve by scattering points homogeneously inside a cuboid containing the intersect
- Efficiency of particle detection with a scintillator
 - ▶ Produced photons are reflected at the surfaces and sometime absorbed
 - ▶ Almost impossible to calculate analytically for different parameters like incident angle, particle energy, ...
 - ▶ Monte Carlo simulation is the only sensible approach

- Complicated function $f(x_1, x_2, \dots, x_n)$, what is the marginal $f_j(x_j) = \int \dots \int f dx_1 \dots dx_{j-1} dx_{j+1} \dots dx_n$?



These problems are easy if we can just sample from the relevant distributions.

Pseudo-random numbers

- Principle: Use insignificant digits of an operation to generate next number
 - ▶ choose large integers λ and m , $\lambda < m$
 - ▶ choose integer $n_0 < m$ (“seed”)
 - ▶ uniformly distributed random numbers r_i :

$$n_{i+1} = \lambda n_i \bmod m$$

$$r_i = n_i / m, \quad r_i \in [0, 1]$$

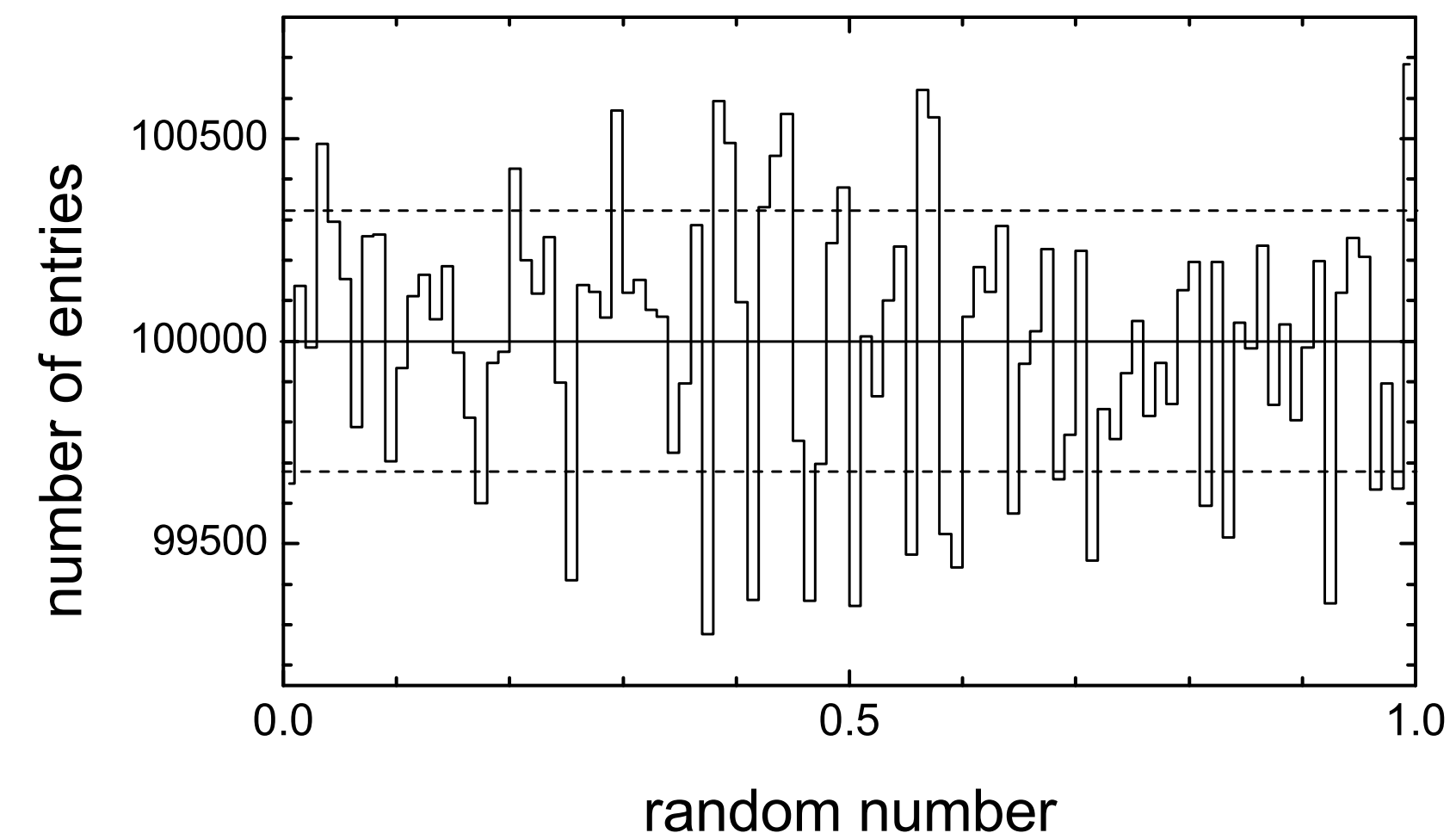
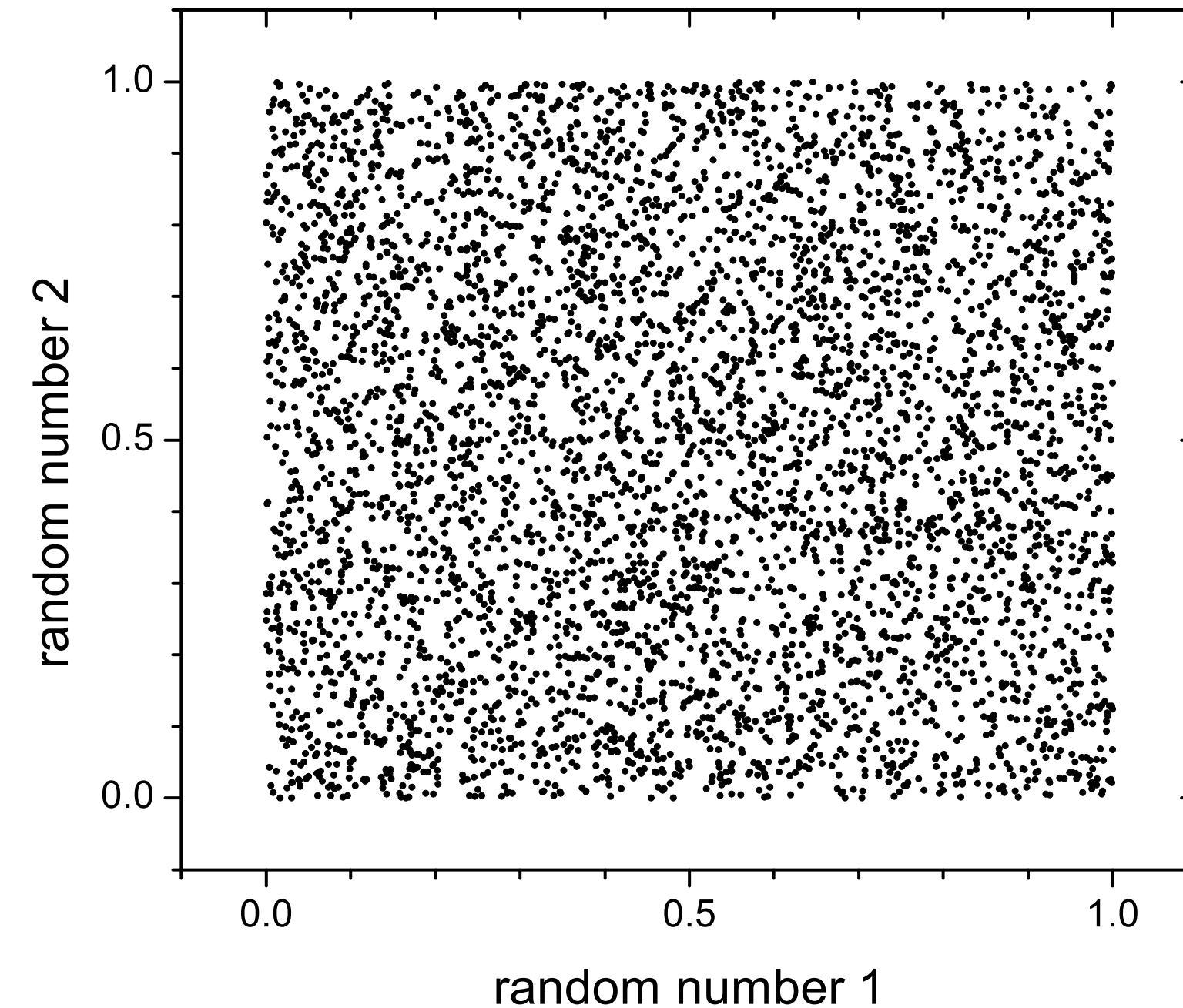
“Multiplicative linear congruential algorithm”
(period at maximum $m - 1$)

- *Mersenne twister*

- ▶ Invented 1997 by M. Matsumoto and T. Nishimura
- ▶ Sequence repeats after 2^{19937} calls, i.e., never ...

- Quality checks

- ▶ Frequency of occurrence
- ▶ Plot correlations between consecutive random numbers



Bohm, Zech:
http://www-library.desy.de/preparch/books/vstatmp_engl.pdf

Random Numbers from distributions: Inverse transform method

Consider a distribution f from which we want to draw random numbers. Let $u(r)$ be the uniform distribution in $[0, 1]$:

$$\int_{-\infty}^x f(x') dx' = \int_0^{r(x)} u(r') dr' = r(x)$$

With $F(x)$ = cumulative distr.:

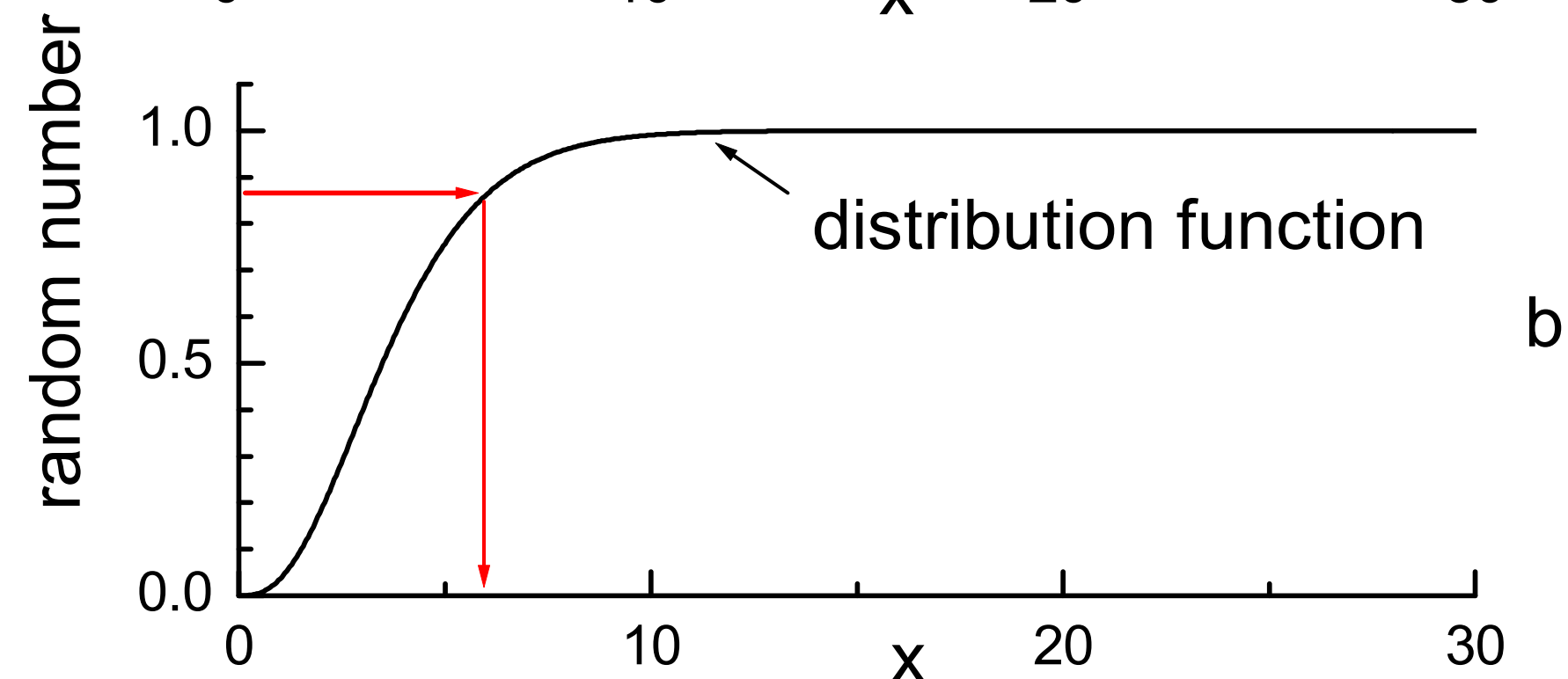
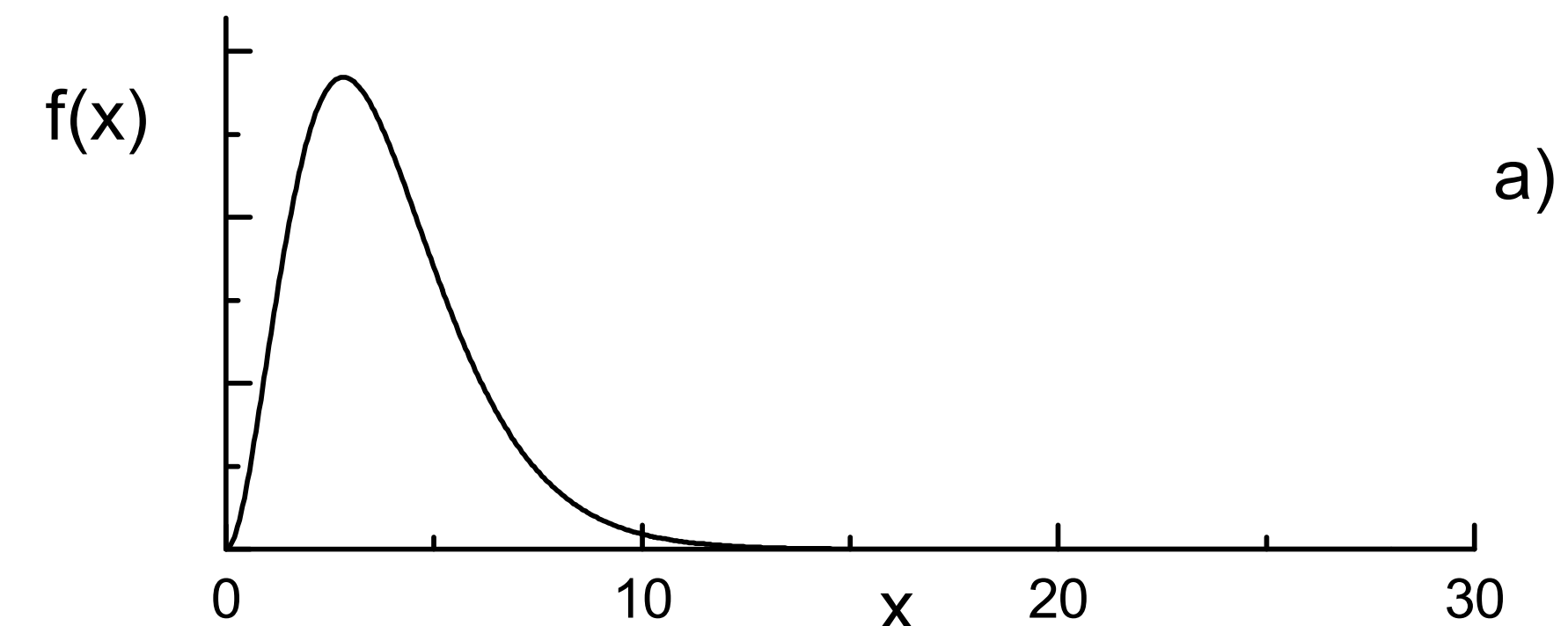
$$F(x) = r$$

We get the random number x from the inverse of the cumulative distribution:

$$x(r) = F^{-1}(r)$$

Bohm, Zech:

http://www-library.desy.de/preparch/books/vstatmp_engl.pdf



Cross check:

$$\frac{dp}{dx} = \underbrace{\frac{dp}{dr}}_{=1} \frac{dr}{dx} = \frac{dF(x)}{dx} = f(x) \quad \square$$

Example I

Linear function:

$$f(x) = 2x, \quad 0 \leq x \leq 1$$
$$F(x) = x^2 \quad \rightarrow \quad x = \sqrt{r}$$

Exponential:

$$f(x) = \gamma e^{-\gamma x}, \quad x \geq 0$$
$$F(x) = 1 - e^{-\gamma x} \quad \rightarrow \quad x = -\frac{\ln(1 - r)}{\gamma}$$

One can store $F(x)$ as a histogram if there is no analytical solution, cf. root's `GetRandom()` function:

```
root [0] TF1 f("f", "x^3/(exp(x)-1)", 0., 15.);
root [1] cout << f.GetRandom() << endl;
13.9571
```

Box-Muller algorithm for creating Gaussian distributed random numbers

1. Generate two uniformly distributed random numbers u_1 and u_2 in the range $[0,1]$

2. Set

$$\phi = 2\pi u_1, \quad r = \sqrt{-2 \ln u_2}$$

3. Then

$$z_1 = r \cos \phi \quad \text{and} \quad z_2 = r \sin \phi$$

are two independent rv's following a standard normal distribution

Why?

$$u_2(r) = e^{-\frac{r^2}{2}}$$

$$\frac{dp}{dr} = \frac{dp}{du_2} \cdot \left| \frac{du_2}{dr} \right| = e^{-\frac{r^2}{2}} r$$

$$dp = \underbrace{\frac{1}{2\pi} e^{-\frac{r^2}{2}} r dr d\phi}_{\text{2d standard normal distribution in polar coordinates}} = \underbrace{\frac{1}{2\pi} e^{-\frac{z_1^2+z_2^2}{2}} dz_1 dz_2}_{\text{2d standard normal distribution in cartesian coordinates}}$$

Inverse transform method using histograms in Python

```
def get_random(f, xmin, xmax, n_samples):  
    """Generate n_samples random numbers within range [xmin, xmax]  
    from arbitrary continuous function f  
    using inverse transform sampling  
    """  
  
    # number of points for which we evaluate F(x)  
    nbins = 10000  
  
    # indefinite integral F(x), normalize to unity at xmax  
    x = np.linspace(xmin, xmax, nbins+1)  
    F = integrate.cumtrapz(f(x), x, initial=0)  
    F = F / F[-1]  
  
    # interpolate F^{-1} and evaluate it for  
    # uniformly distributed rv's in [0,1[  
    inv_F = interpolate.interp1d(F, x, kind="quadratic")  
    r = np.random.rand(n_samples)  
    return inv_F(r)
```

[\[random numbers from distribution.ipynb\]](#)

Example II: Uniform points on a sphere

$$\frac{dp}{d\Omega} = \frac{dp}{\sin \theta d\theta d\phi} = \text{const} \equiv k$$

$$\frac{dp}{d\theta d\phi} = k \sin \theta \equiv f(\phi)g(\theta)$$

Distributions for θ and ϕ :

$$f(\phi) \equiv \frac{dp}{d\phi} = \text{const} = \frac{1}{2\pi}, \quad 0 \leq \phi \leq 2\pi$$

$$g(\theta) \equiv \frac{dp}{d\theta} = \frac{1}{2} \sin \theta, \quad 0 \leq \theta \leq \pi$$

Calculating the inverse of the cumulative distribution we obtain:

$$\phi = 2\pi r_1$$

$$\theta = \arccos(1 - 2r_2) \quad [\text{as } G(\theta) = \frac{1}{2}(1 - \cos \theta)]$$

Upshot: ϕ and $\cos \theta$ need to be distributed uniformly

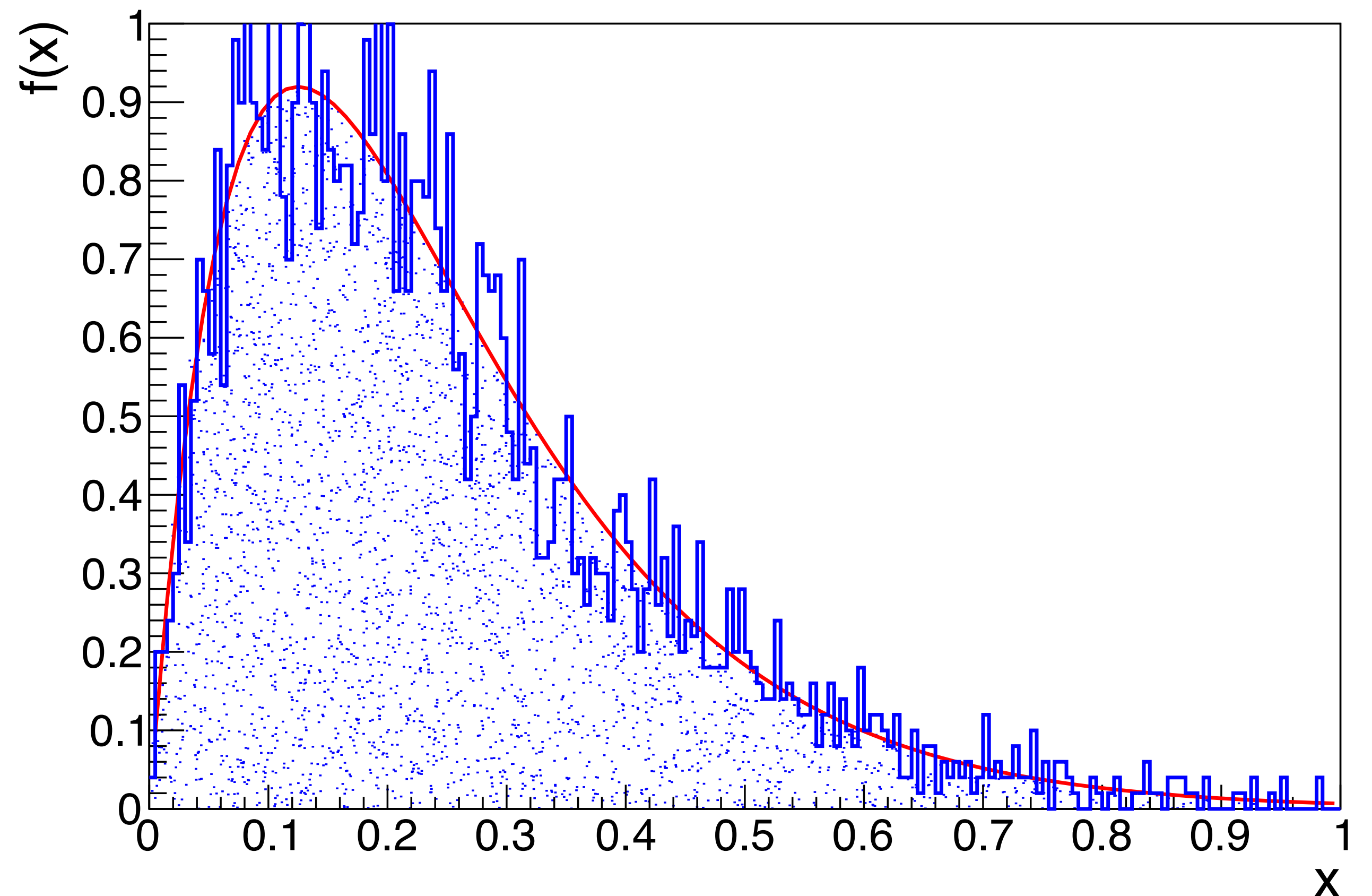
Random numbers from distributions: Acceptance-rejection method

■ Idea:

- ▶ Create flat distribution in x, y
- ▶ Use points below function value
- ▶ X values are distributed like the distribution

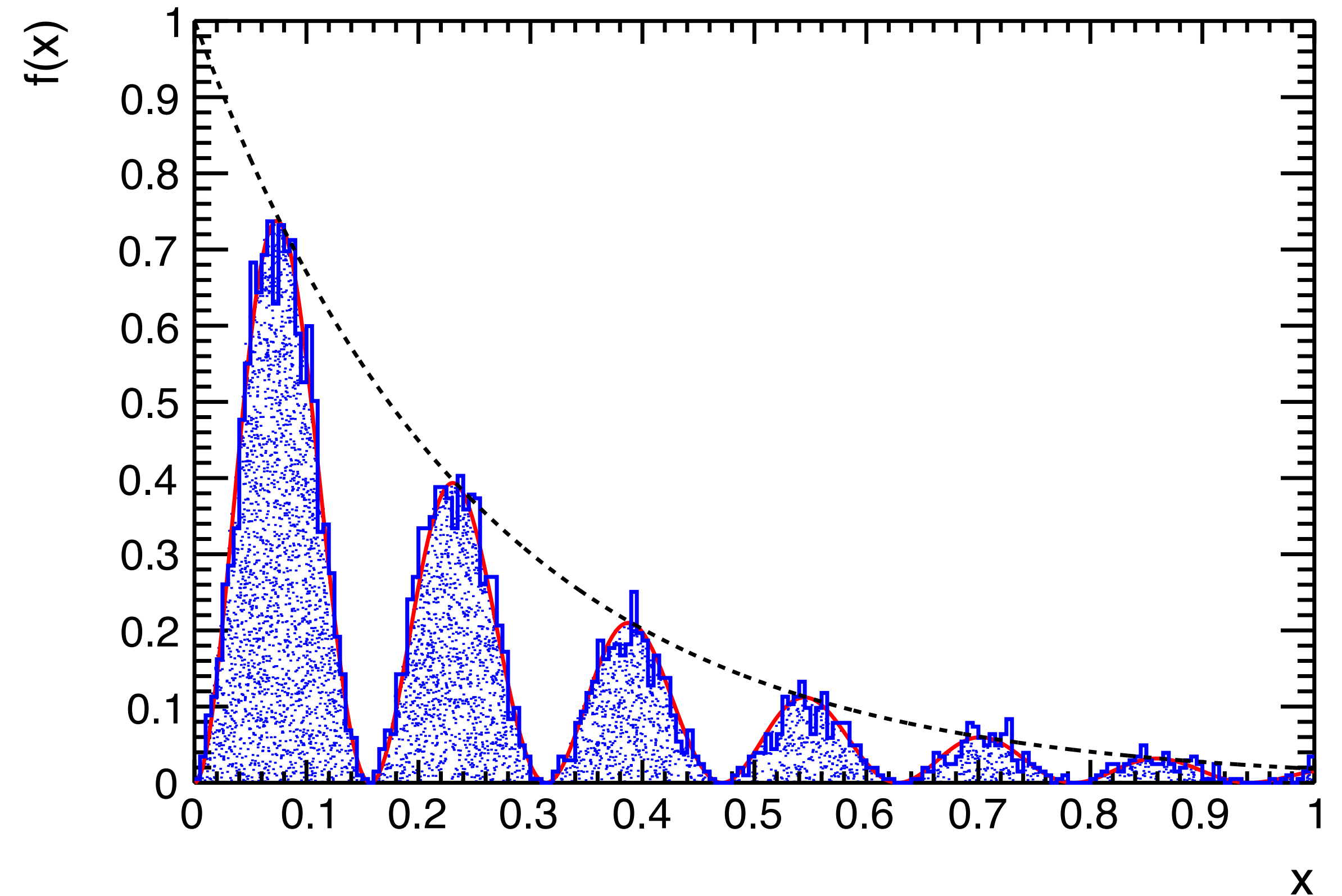
■ Algorithm

- ▶ Generate random number x uniformly between a and b
- ▶ Generate another number $y \in [0, 1]$
- ▶ For some number A larger than the maximum of the function: accept if $y < f(x)/A$
- ▶ Repeat many times



Random numbers from distributions: Acceptance-rejection method (II)

- The efficiency of this algorithm can be quite small (e.g. long, small tails)
- Find another distribution p_m that is easy to sample, and where $f_m = \alpha p_m$ is always larger than the target distribution for some α
- This is called a *majorant*
- Now sample from p_m and accept points with the condition $y < f(x)/f_m(x)$
- (Example: $\sin^2(c_1 x) \exp(-c_2 x)$ is hard to sample from, but the majorant $\exp(-c_2 x)$ is easy to sample)

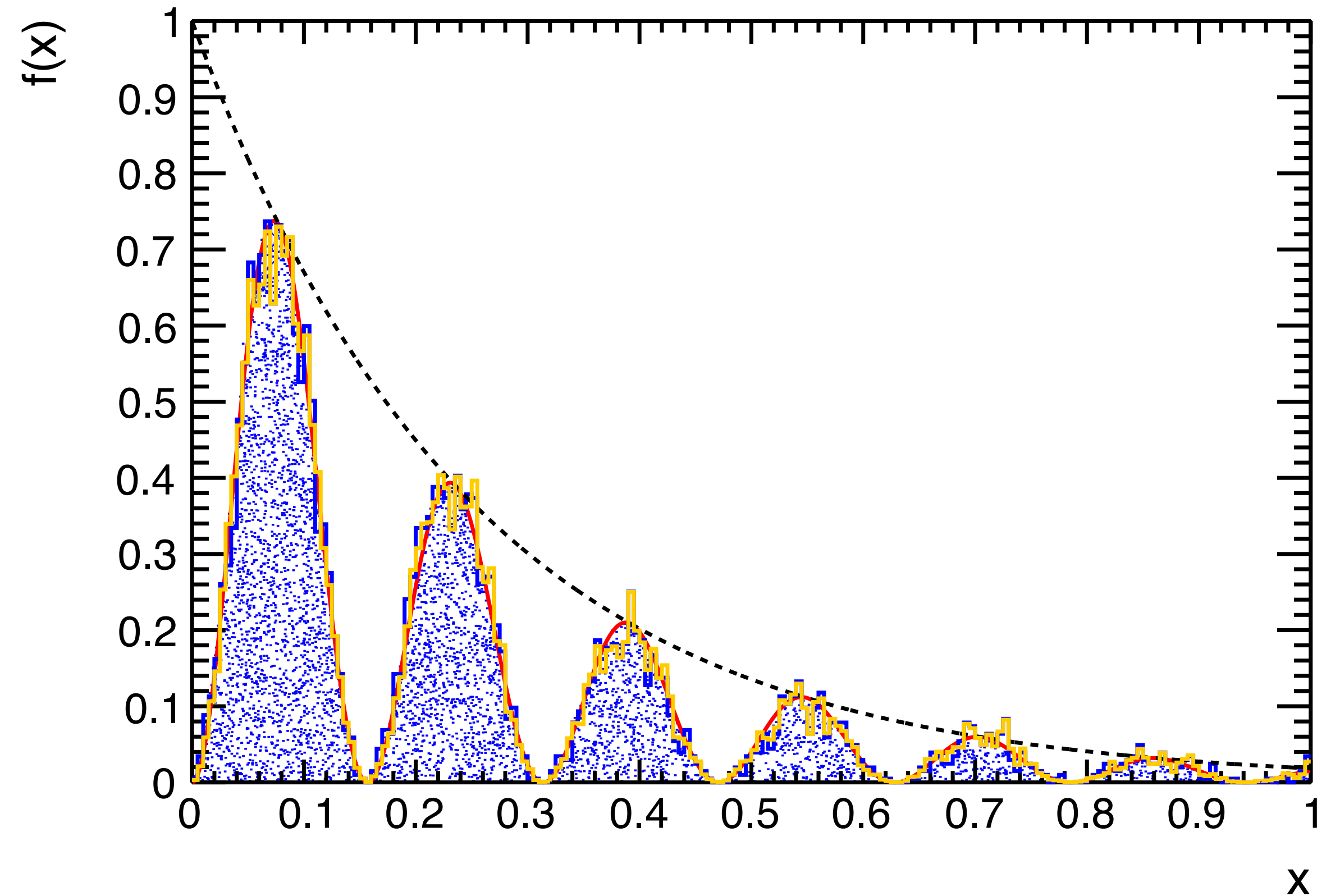


Importance Sampling

- Almost the same as accept-reject
- Instead of accepting point with probability $f(x)/f_m(x)$, always accept, but with weight

$$w = f(x)/f_m(x)$$

- Comparison function does not need to be a majorant



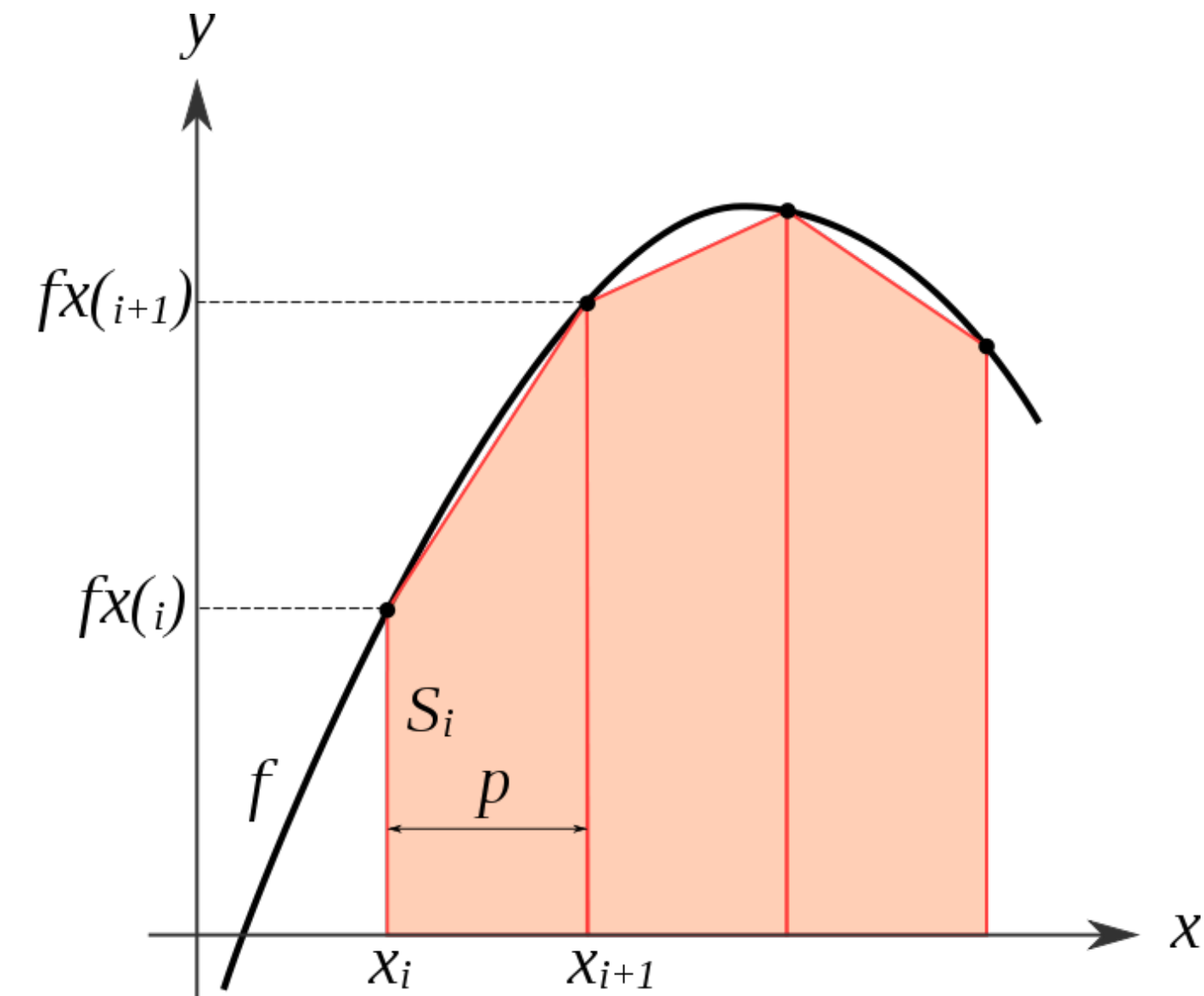
The curse of dimensionality

Trapezoidal rule in one dimension

- ▶ accuracy improves as $1/n^2$ with the number of points

Monte Carlo integration in d dimensions:

- ▶ Averages, fractions from independent points all scale with $1/\sqrt{n}$
- ▶ n is accepted number of points



https://en.wikipedia.org/wiki/Trapezoidal_rule

Trapezoidal rule in d dimension:

- ▶ accuracy improves as $1/n^{2/d}$ with the number of points
- ▶ for $d > 4$ the dependence on n is better for MC integration
- ▶ (However, fraction of accepted points tend to lower a bit with higher dimensions)

same as in 1d case

For multidimensional integrals
MC integration outperforms
other numerical integration
methods

Metropolis-Hastings algorithm (1)

Bayesian inference often involves marginalization of a high-dimensional posterior distribution:

$$P(\theta_0|\text{data}) = \int P(\theta_0, \theta_1, \dots, \theta_n|\text{data}) d\theta_1 \dots d\theta_n$$

Typically, the integral cannot be solved in closed form. Moreover, repeated one-dimensional integration becomes inefficient (“curse of dimensionality”).

Idea: sample distribution many times and consider only parameter of interest.

Method: Markov Chain Monte Carlo (MCMC)

MCMC has revolutionized Bayesian analysis.

A sequence of random numbers is a Markov chain if the probability of the next number only depends on the previous one:

$$f(x_{n+1}|x_n, x_{n-1}, \dots, x_0) = f(x_{n+1}|x_n)$$

Metropolis-Hastings algorithm (2)

Goal: sample from a distribution $f(\vec{x})$ known up to a normalization constant.

Take initial \vec{x}_0 with $f(\vec{x}_0) > 0$ and repeat the following steps many times:

1. Generate candidate \vec{y} according to proposal distribution $q(\vec{y} | \vec{x}_k)$
2. Generate uniformly distributed random number r in $[0, 1]$ and set

$$\vec{x}_{k+1} = \begin{cases} \vec{y}, & \text{if } r \leq \alpha(\vec{x}_k, \vec{y}) \\ \vec{x}_k, & \text{otherwise} \end{cases}$$

where

$$\alpha(\vec{x}, \vec{y}) = \min \left\{ 1, \frac{f(\vec{y}) q(\vec{x} | \vec{y})}{f(\vec{x}) q(\vec{y} | \vec{x})} \right\}$$

$\alpha(\vec{x}, \vec{y})$ is called the acceptance probability.

Metropolis-Hastings algorithm (3)

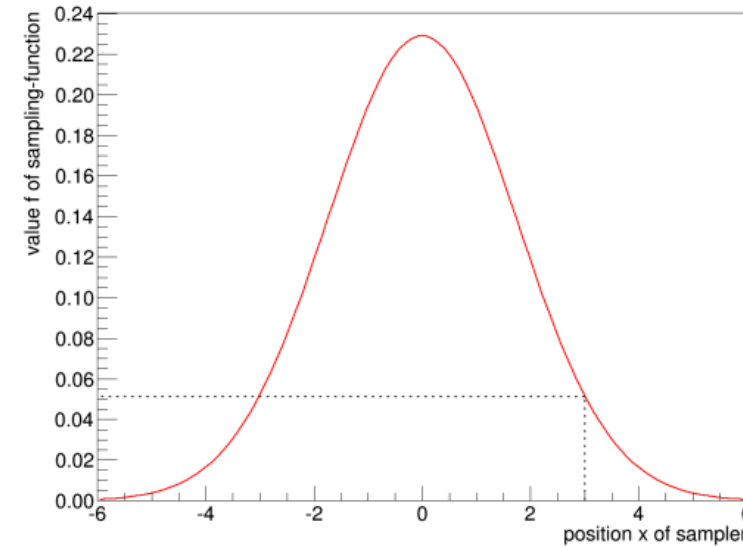
- The algorithm generates a *correlated* sequence of points (not suited for many applications, but okay for marginalization)
- If a finite initial sequence of points is discarded, the remaining points can be shown to follow $f(\vec{x})$
- Not easy to figure out when the sequence has started to converge to $f(\vec{x})$
- The proposal function $q(\vec{y} | \vec{x})$ can be almost anything. Often, a multi-dimensional Gaussian is used.
- Often the proposal function is symmetric, i.e., $q(\vec{y} | \vec{x}) = q(\vec{x} | \vec{y})$. Then the acceptance probability reduces to

$$\alpha(\vec{x}, \vec{y}) = \min \left\{ 1, \frac{f(\vec{y})}{f(\vec{x})} \right\}$$

and a step to a higher $f(\vec{y})$ is always taken.

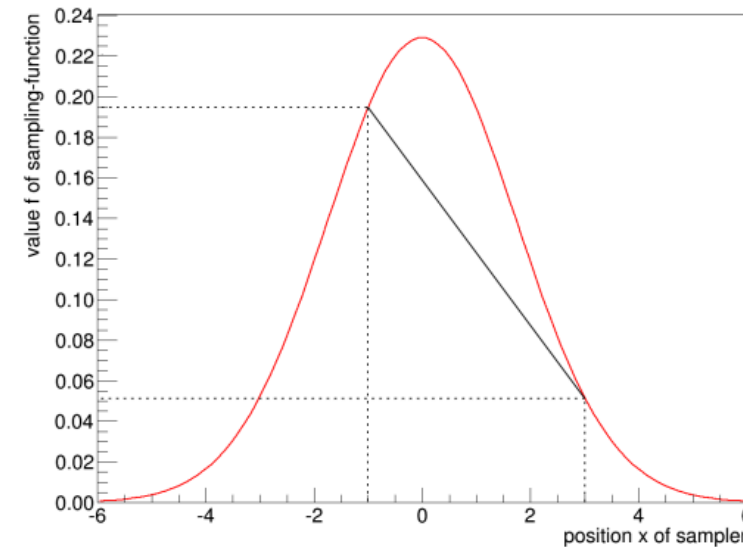
- Original Metropolis algorithm suggested symmetric proposal functions, Hastings modified original rules by using non-symmetric functions.

Example



$$x_{start} = 3$$

Figure 1.5: A starting point is chosen.



$$x_{old} = x_{start} = 3$$

$$p = -4$$

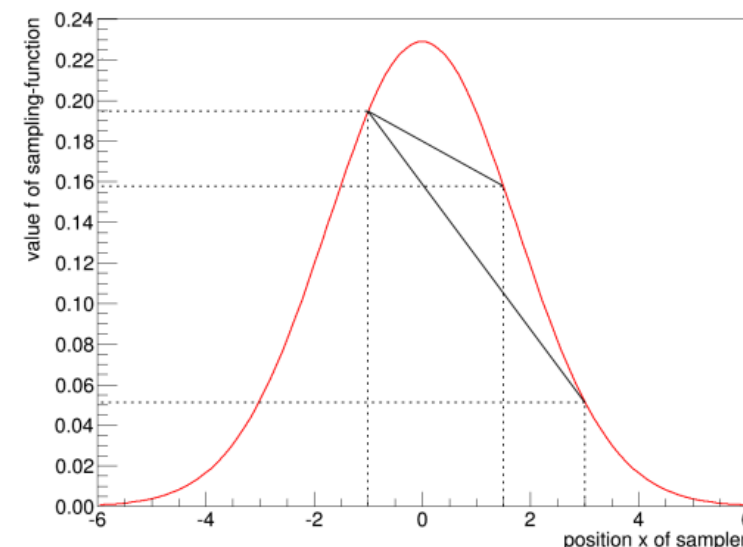
$$x_{new} = x_{old} + p = -1$$

$$\rho = \min\left(1, \frac{f(x_{new})}{f(x_{old})}\right) = \min(1, 3.75) = 1$$

$$u = 0.9$$

$$\rho > u \Rightarrow \text{accept}$$

Figure 1.6: For $f(x_{new}) > f(x_{old})$ the step is always accepted.



$$x_{old} = -1$$

$$p = 2.5$$

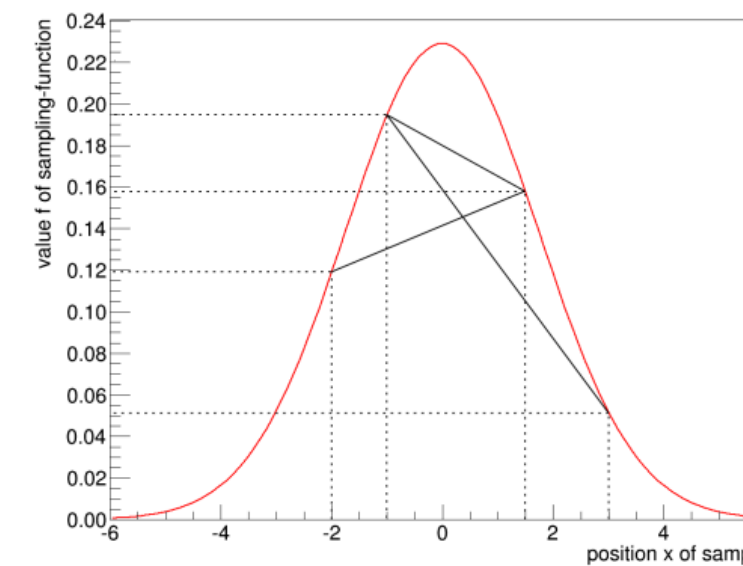
$$x_{new} = x_{old} + p = 1.5$$

$$\rho = \min\left(1, \frac{f(x_{new})}{f(x_{old})}\right) = \min(1, 0.81) = 0.81$$

$$u = 0.4$$

$$\rho > u \Rightarrow \text{accept}$$

Figure 1.7: For $f(x_{new}) < f(x_{old})$ it depends on u whether a step is accepted.



$$x_{old} = 1.5$$

$$p = -3.5$$

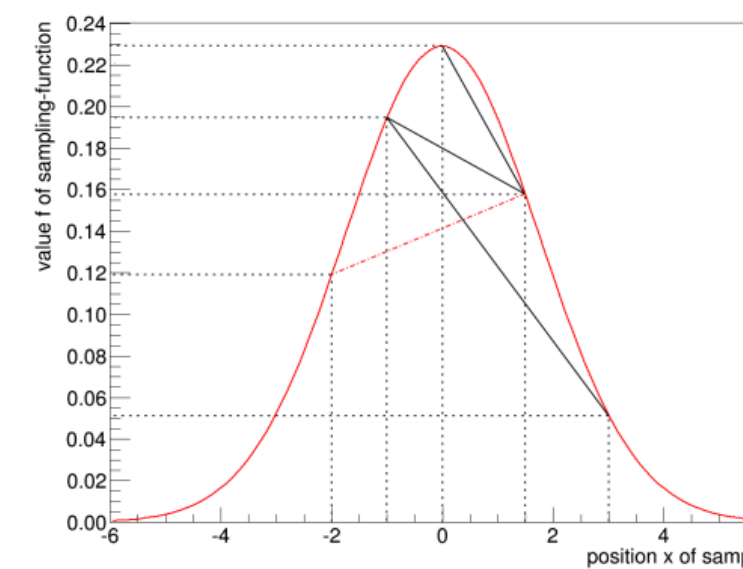
$$x_{new} = x_{old} + p = -2$$

$$\rho = \min\left(1, \frac{f(x_{new})}{f(x_{old})}\right) = \min(1, 0.75) = 0.75$$

$$u = 0.8$$

$$\rho < u \Rightarrow \text{reject}$$

Figure 1.8: For $\rho < u$ the step is rejected.



$$x_{old} = 1.5$$

$$p = -1.5$$

$$x_{new} = x_{old} + p = 0$$

$$\rho = \min\left(1, \frac{f(x_{new})}{f(x_{old})}\right) = \min(1, 1.45) = 1$$

$$u = 0.6$$

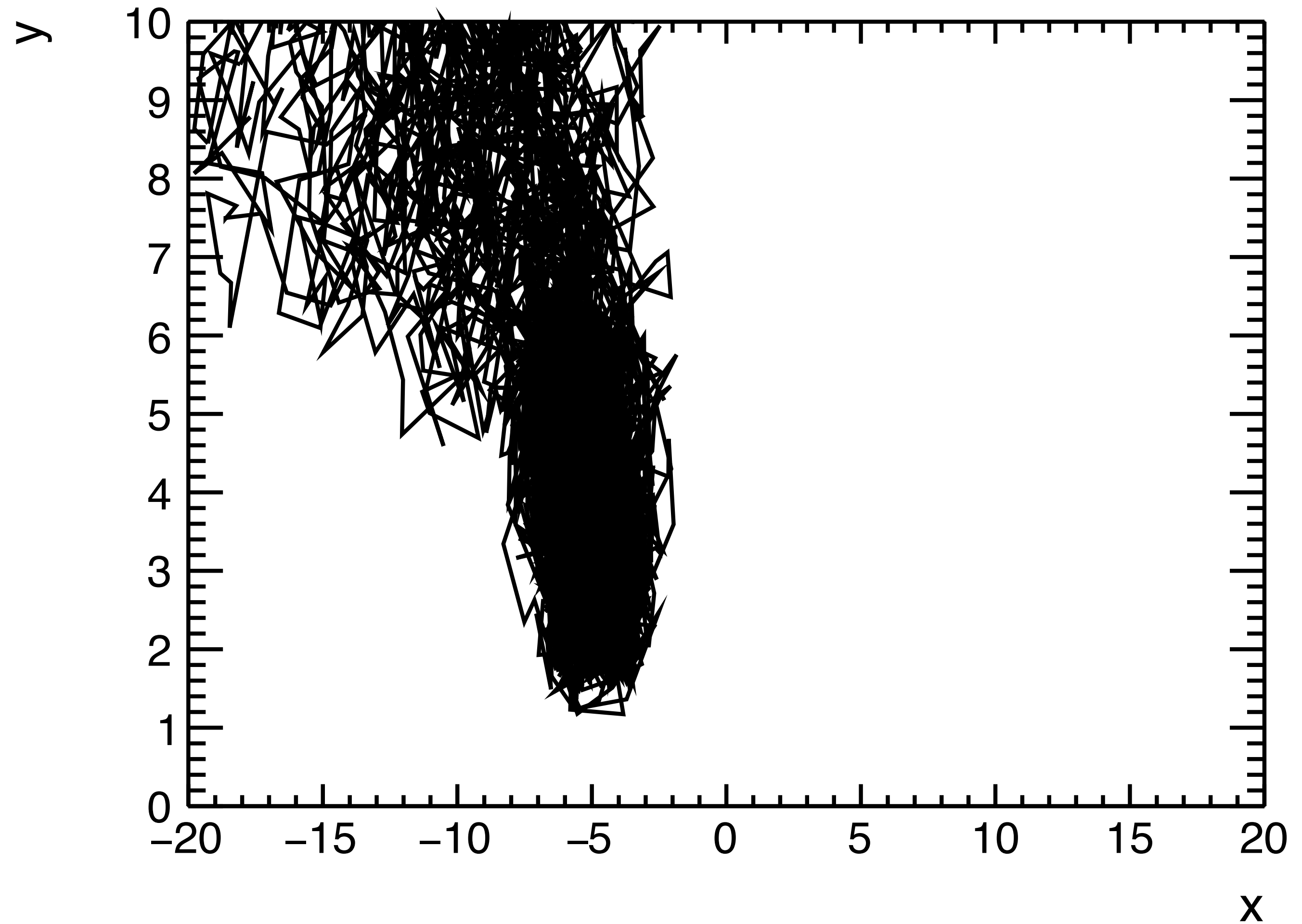
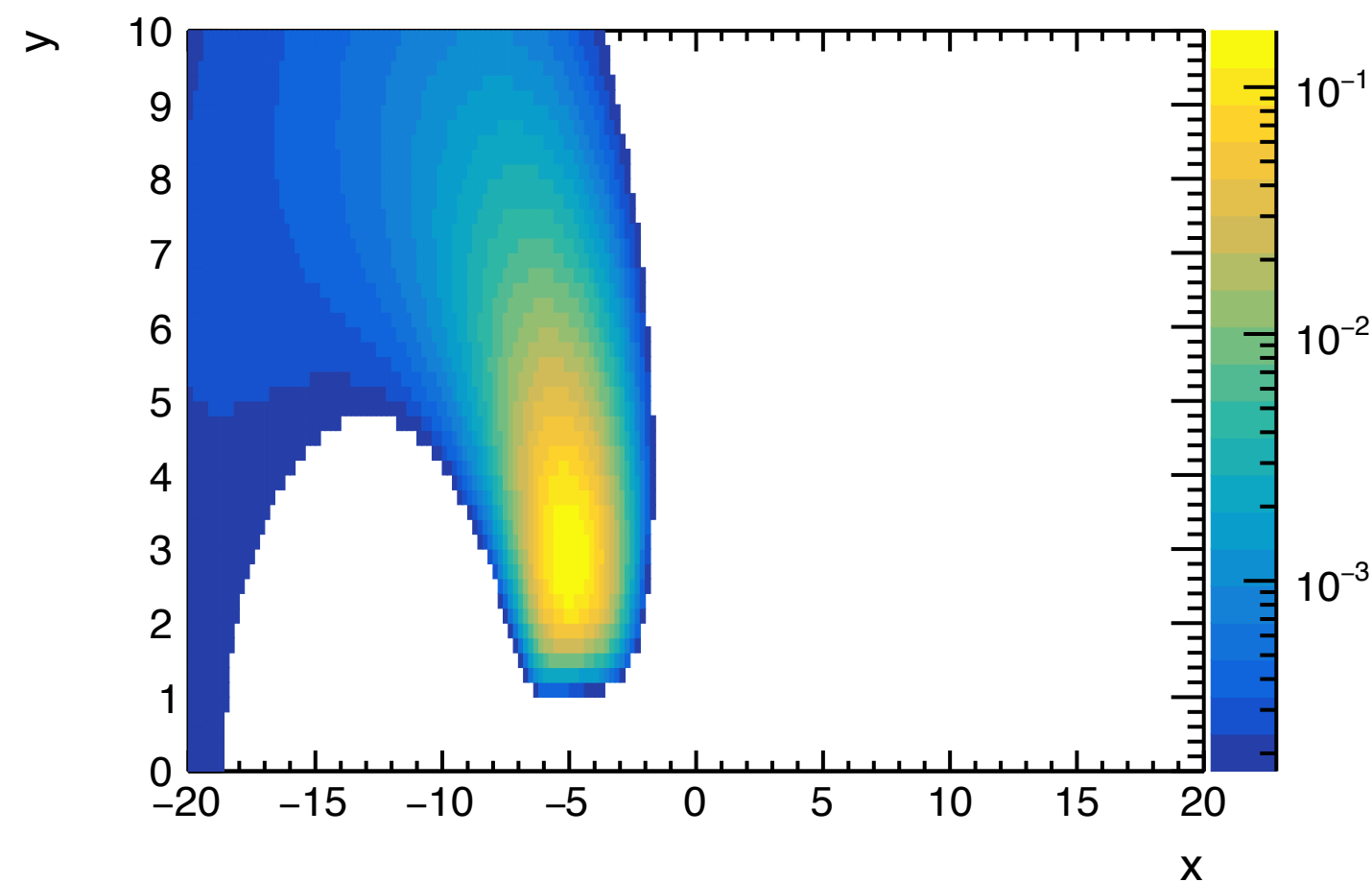
$$\rho > u \Rightarrow \text{accept}$$

Figure 1.9: For $\rho > u$ the step is accepted again.

- Each step only depends on the previous point

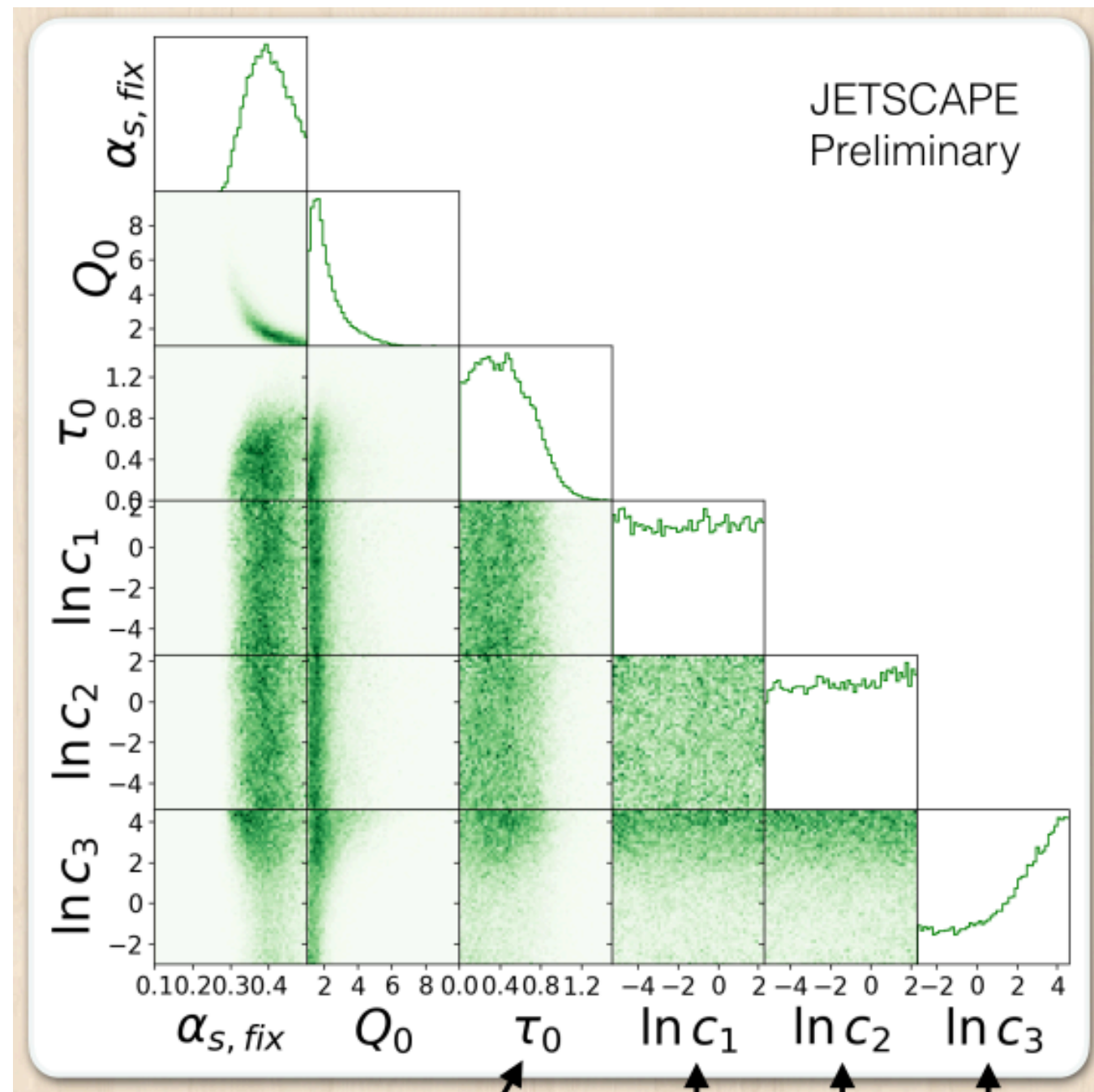
Metropolis-Algorithm Example 2

- Random-walk-like behaviour
- More time is spent in regions of high probability
- True distribution:

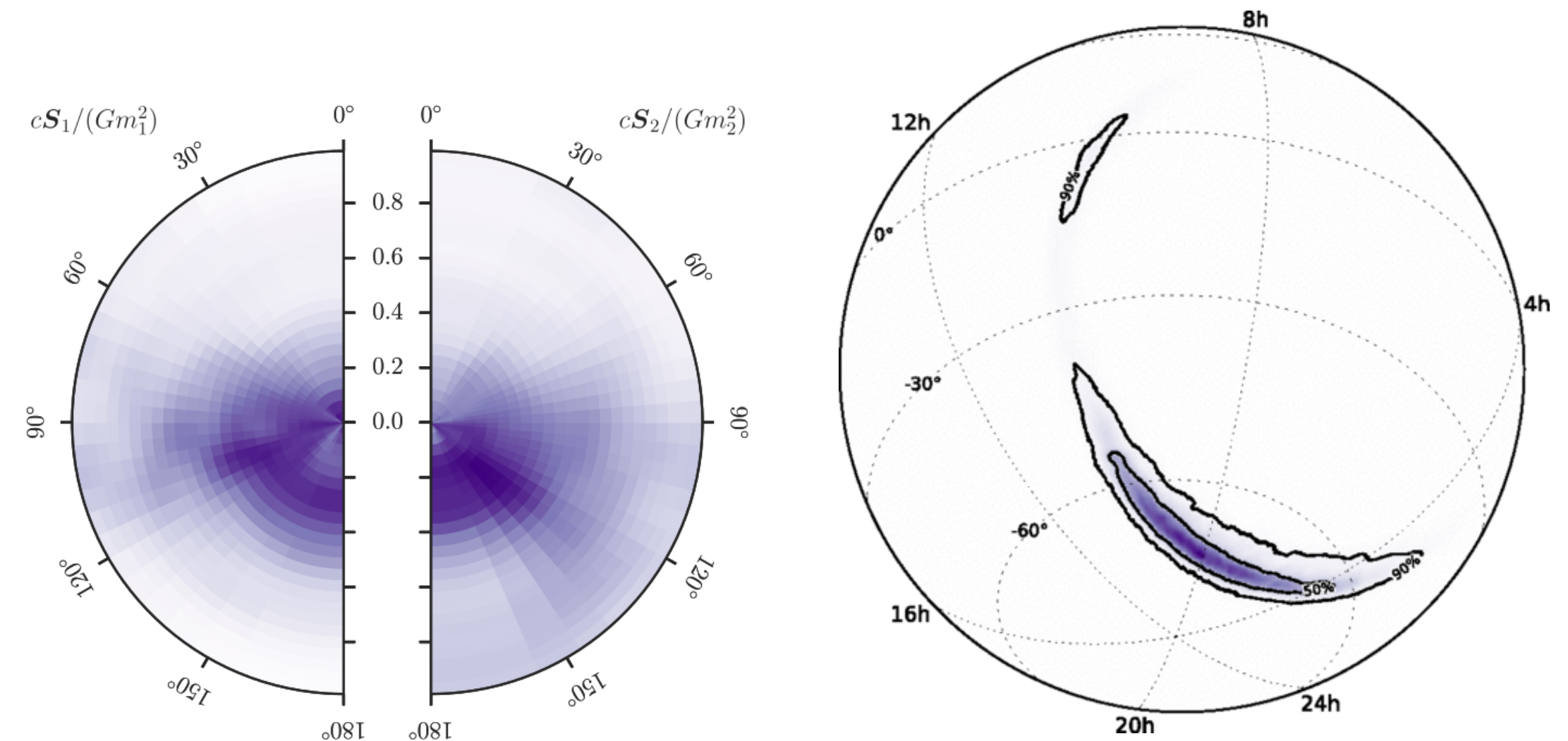
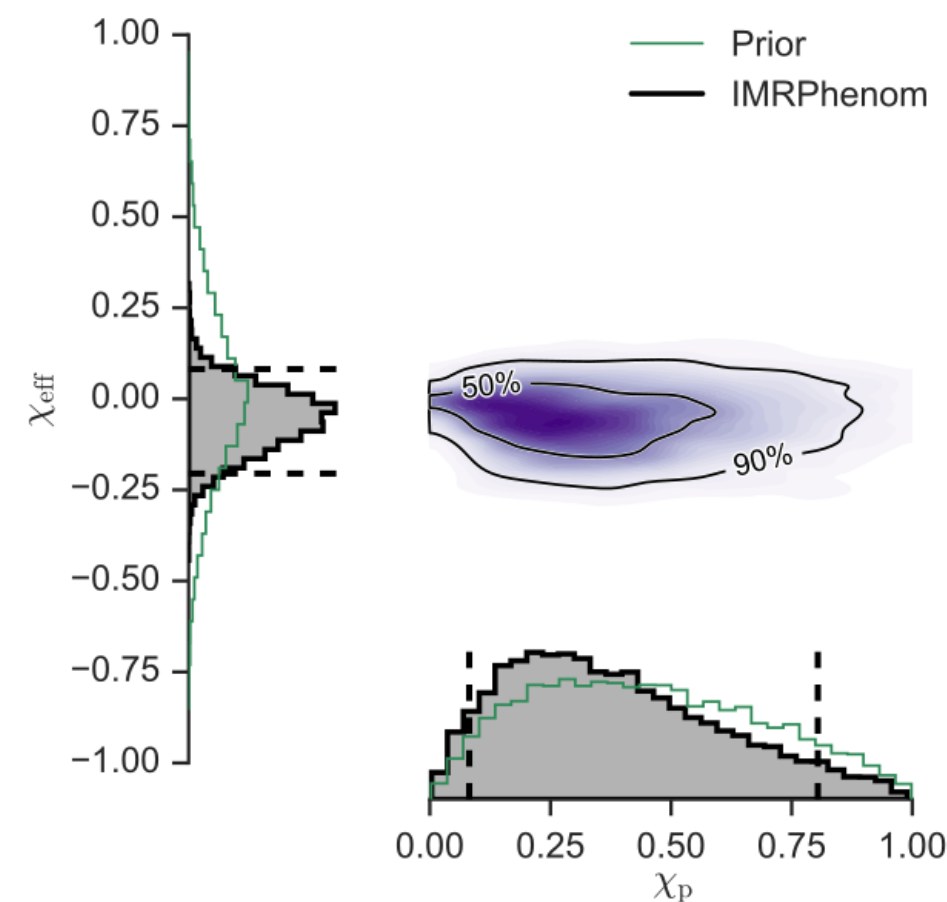


MCMC marginals

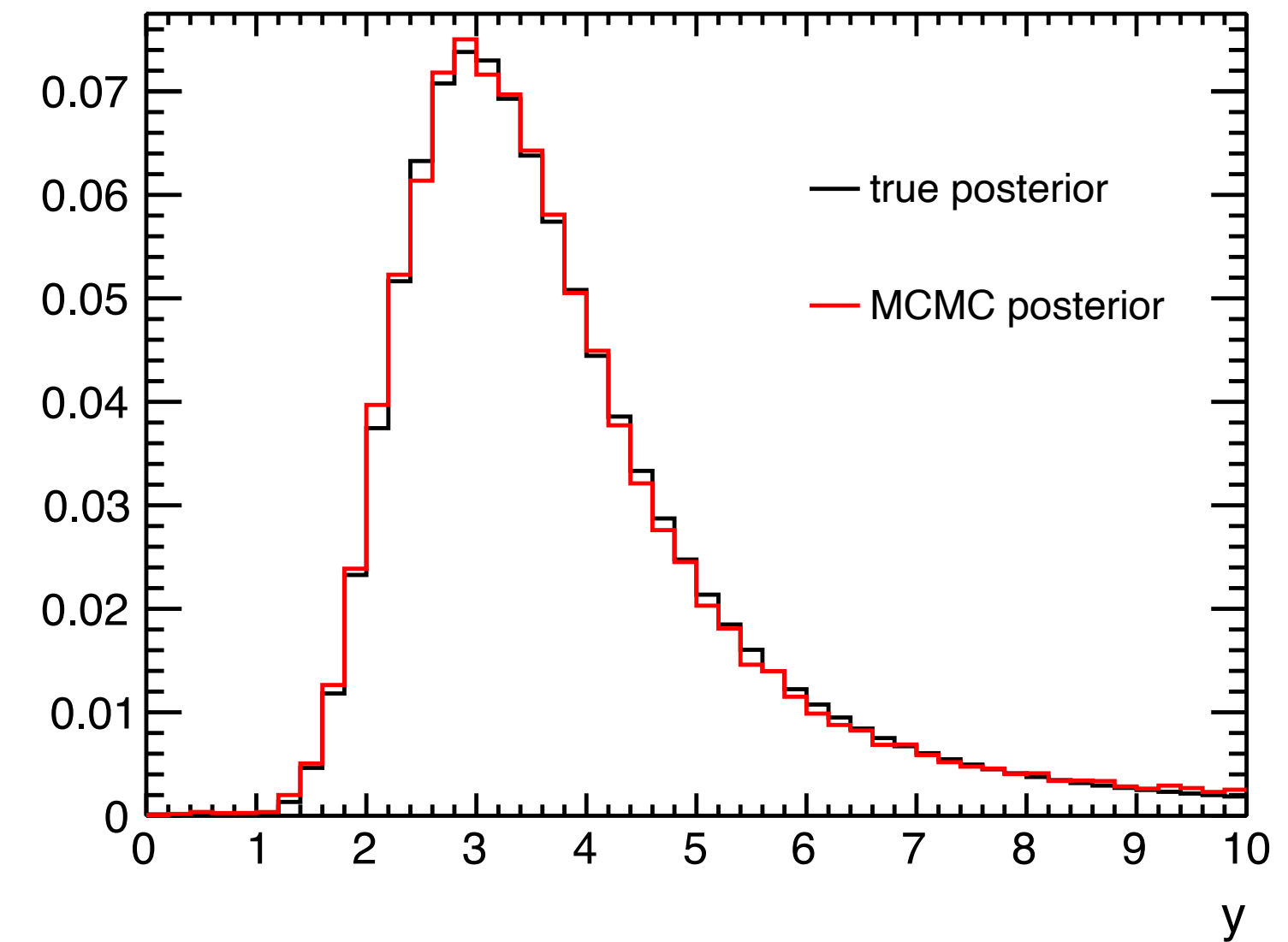
- With sampling, marginals become trivially easy: Just take the distribution of the coordinate
- Means and variances are estimated by the means and variances of the sampled points



JETSCAPE Collaboration

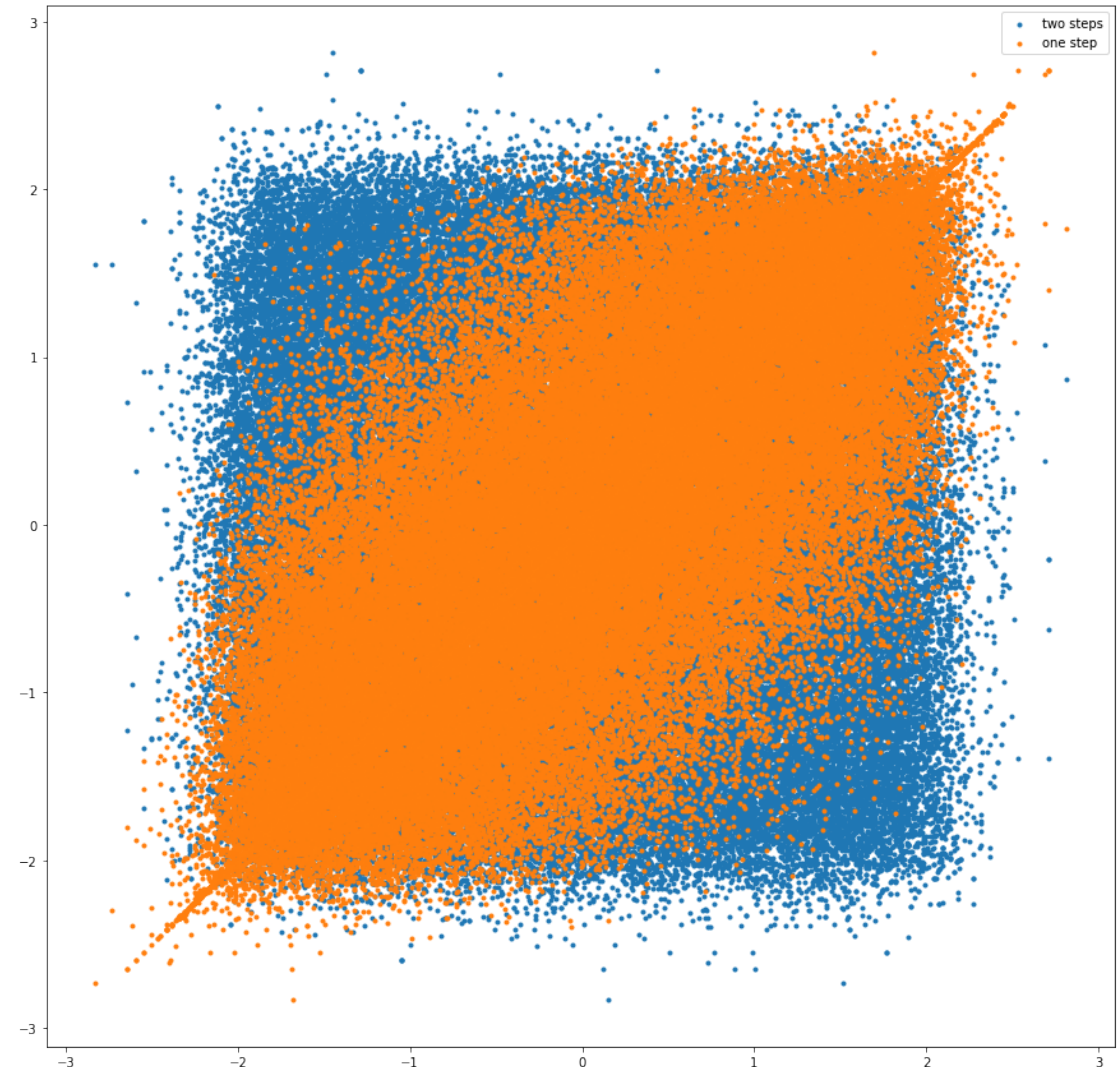


LIGO Collaboration



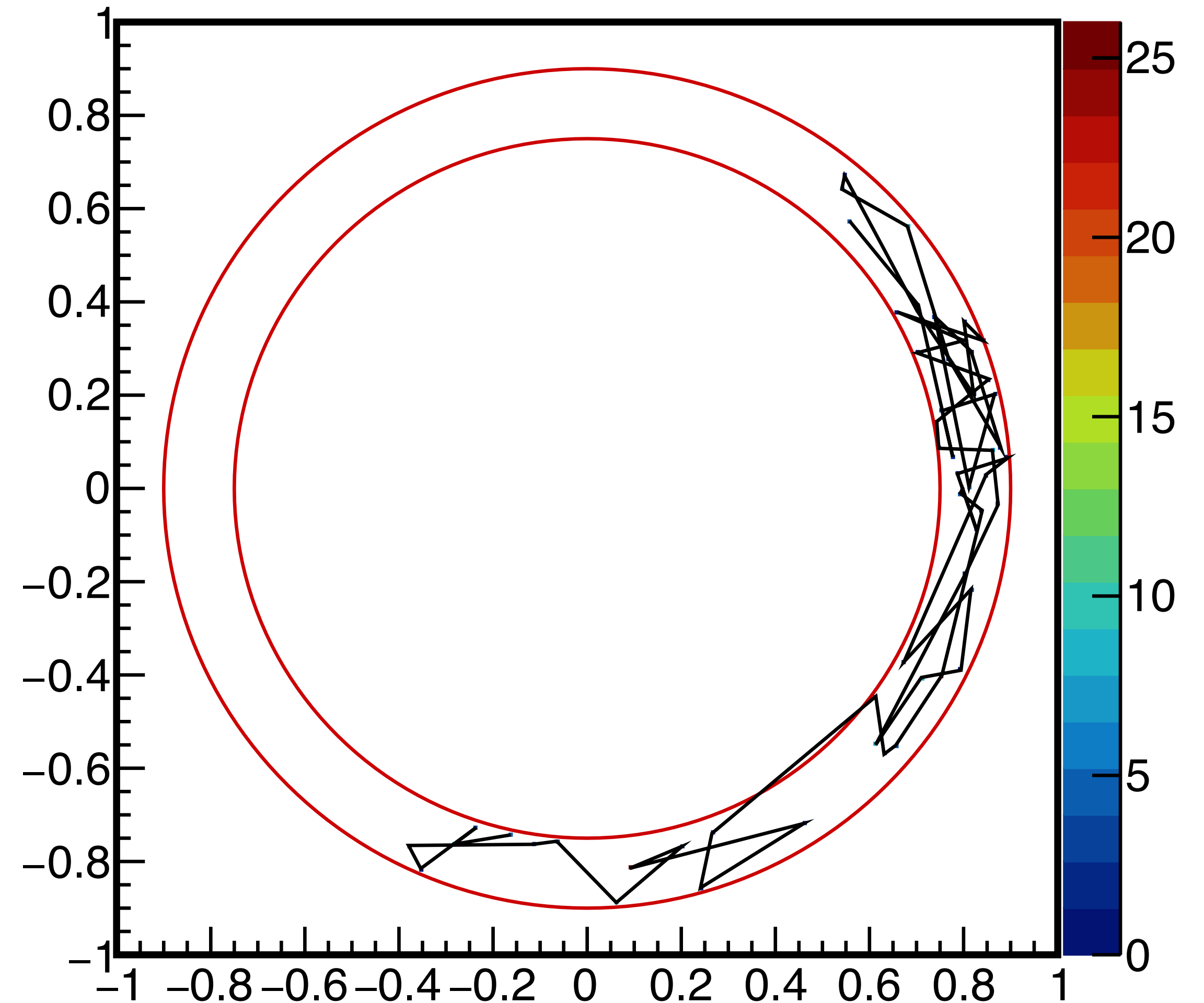
Properties of MCMCs

- Next position depends on current one - samples are not uncorrelated
 - ▶ But after a few steps they essentially are
- As long as all parts of the distribution are reachable by the steps, the result will always converge to the true distribution
 - ▶ But this can take a long time
- If the algorithm starts at a point of very low probability, then these initial points take a long time to become unimportant for e.g. averages
 - ▶ Ignore some of the initial steps - the “burn-in” - for faster convergence



A difficult example

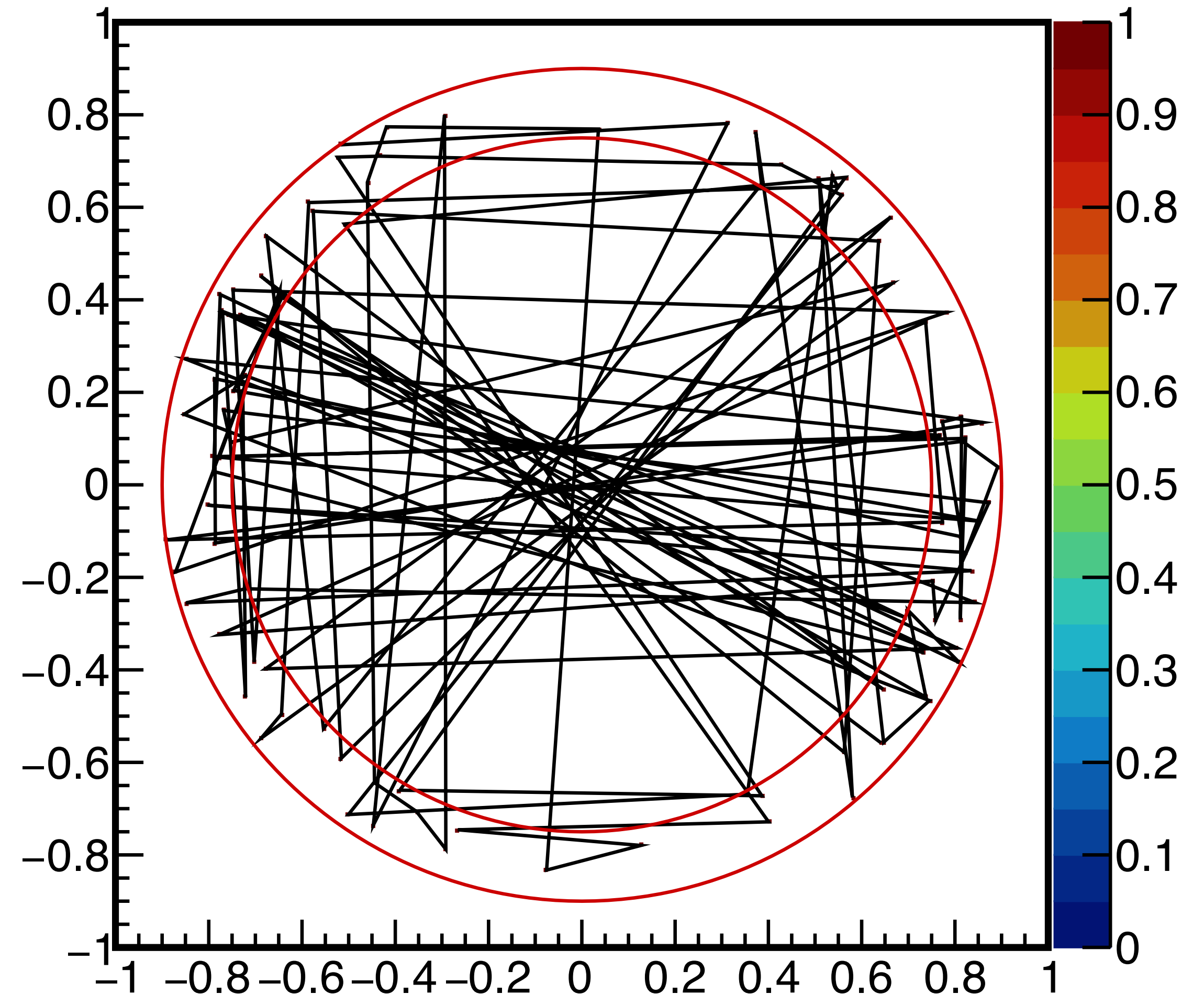
- Thin distributions can make it difficult for the walker to traverse the distribution
 - ▶ Small steps: many iterations to go around
 - ▶ Large steps: most steps lead out of the ring and are rejected
- Many types of MCMC algorithms are developed to deal with difficult distributions, high dimensions, improve convergence etc.



The distribution is constant on the ring and 0 elsewhere

The Gibbs-sampler

- Useful if we can sample easily from the conditional probability (keeping all coordinates except for one fixed)
 - ▶ e.g. $p(x_j | x_1, x_2, \dots, x_{j-1}, x_{j+1}, \dots, x_n)$
- Start sampling from $p(x_1 | x_2, \dots, x_n)$ - new value x'_1
- Now sample from $p(x_2 | x'_1, x_3, \dots, x_n)$
- Repeat for all variables, new position is \vec{x}'
 - ▶ No rejected steps
 - ▶ Can move through distributions quickly
- Special case of the Metropolis-Hastings algorithm



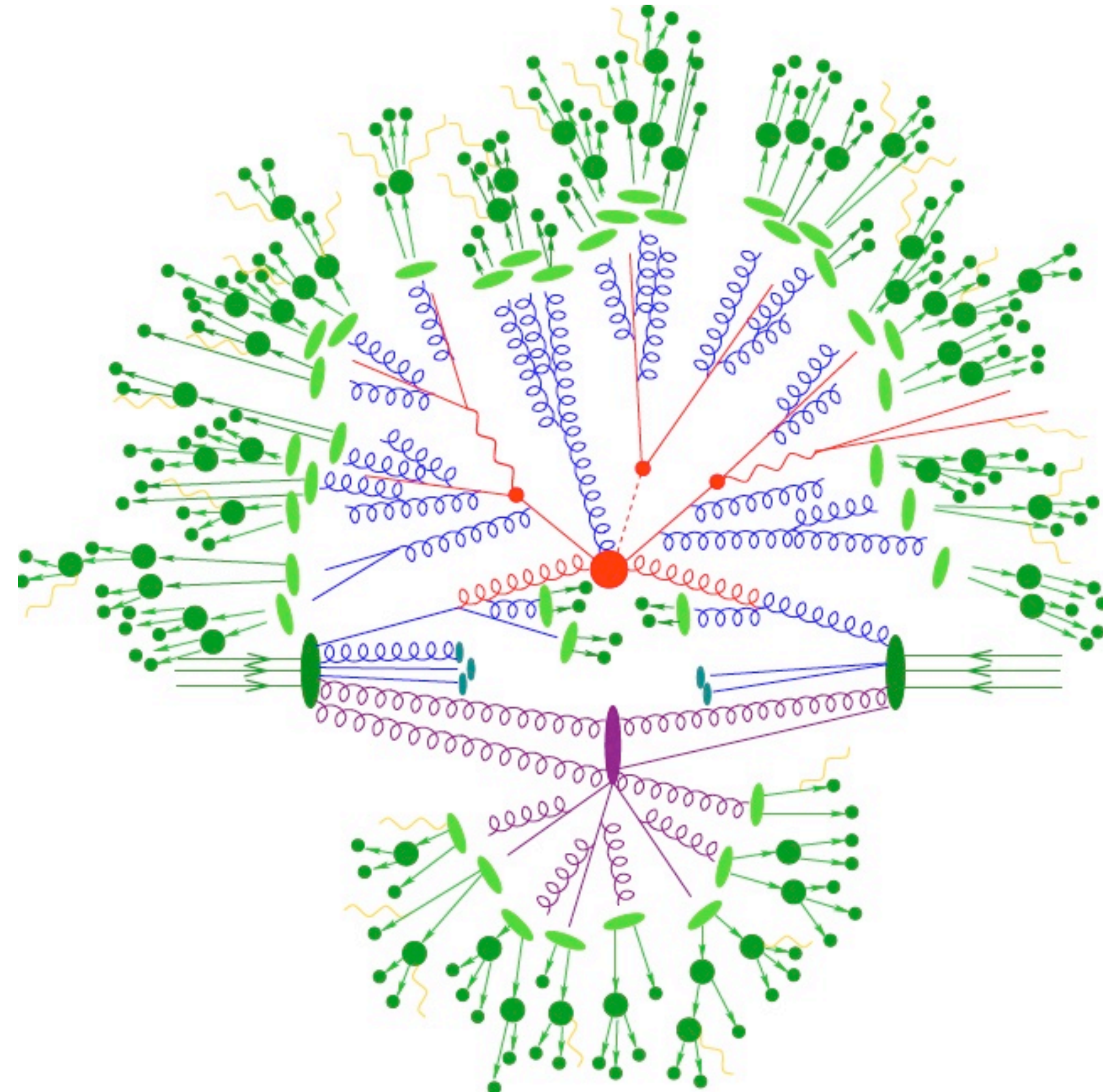
Event Generators

- Imagine a likelihood for getting the measured detector signals based on a parameter of interest and a bunch of others:
 - ▶ Consider all possible processes; all possible angles in multiple scattering, all fluctuations in detector signal
 - ▶ Then integrate out all other variables
 - ▶ Not viable
- Instead simplify:
 - ▶ Tracking algorithms, detector signal reconstruction, track end event selections/triggers
 - ▶ Results in efficiencies - would still need to integrate over all processes and detector signals to find them
- Sampling from physics events and detector responses makes this easier (“Monte Carlo” generators/simulations)

Monte Carlo simulation I: Event generators (Pythia, Sherpa, ...)

Examples: Pythia

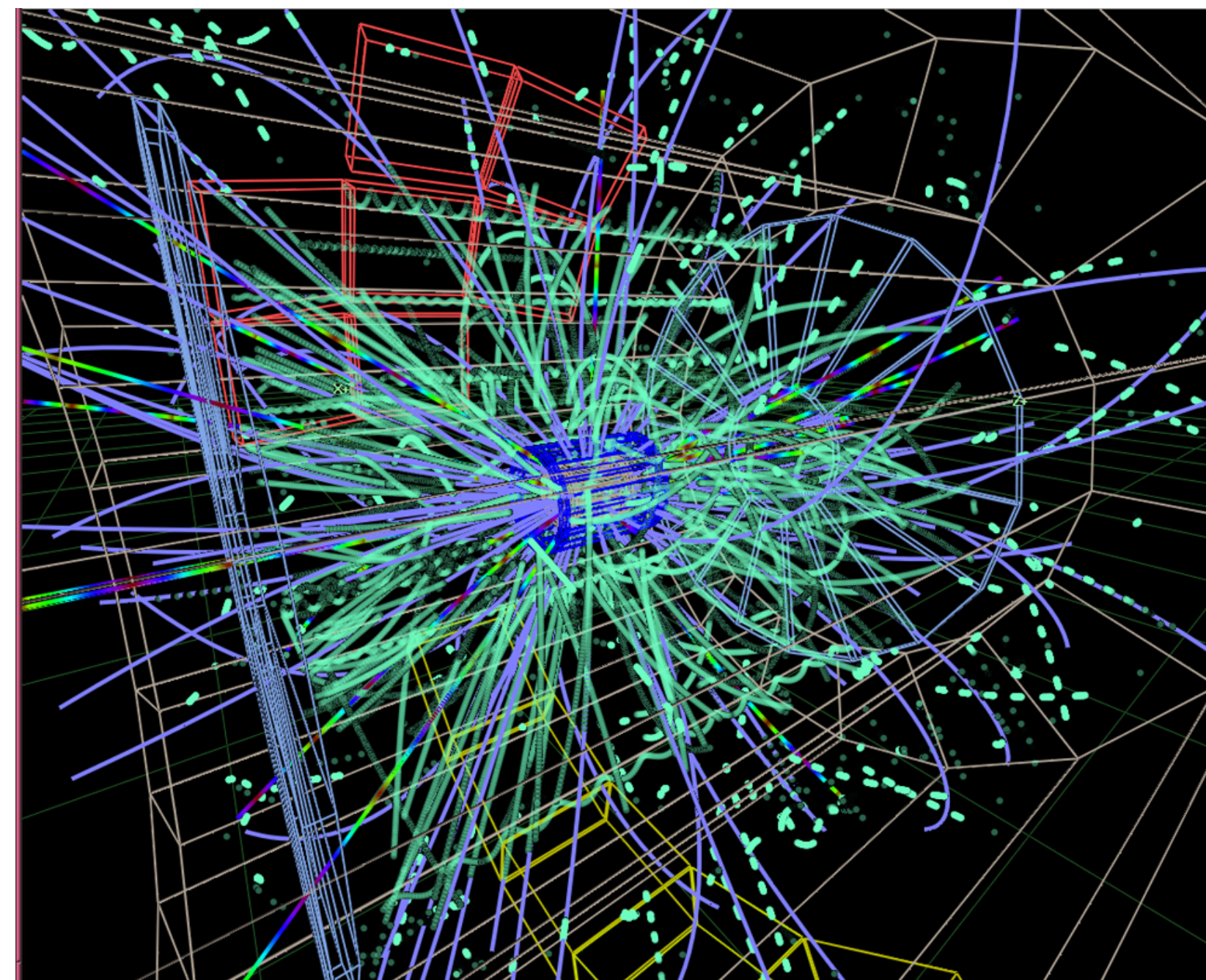
- ▶ Simulation of pp and e^+e^- collision on quark and gluon level
- ▶ Hard and soft interactions, parton showers, fragmentation and particle decay
- ▶ Many applications
 - Test underlying physics, e.g., perturbative QCD
 - Calculate QCD background processes, e.g., in Higgs searches
 - Calculation of detector efficiencies



Pythia

Output:

Four-vectors of produced particles



Event listing (summary)

| I | particle/jet | KS | KF | orig | p_x | p_y | p_z | E | m | |
|----|--------------|----|----|-------|-----|--------|--------|---------|--------|--------|
| 1 | (u) | A | 12 | 2 | 0 | 0.000 | 0.000 | 10.000 | 10.000 | 0.006 |
| 2 | (ubar) | V | 11 | -2 | 0 | 0.000 | 0.000 | -10.000 | 10.000 | 0.006 |
| 3 | (string) | | 11 | 92 | 1 | 0.000 | 0.000 | 0.000 | 20.000 | 20.000 |
| 4 | (rho+) | | 11 | 213 | 3 | 0.098 | -0.154 | 2.710 | 2.856 | 0.885 |
| 5 | (rho-) | | 11 | -213 | 3 | -0.227 | 0.145 | 6.538 | 6.590 | 0.781 |
| 6 | pi+ | | 1 | 211 | 3 | 0.125 | -0.266 | 0.097 | 0.339 | 0.140 |
| 7 | (Sigma0) | | 11 | 3212 | 3 | -0.254 | 0.034 | -1.397 | 1.855 | 1.193 |
| 8 | (K*+) | | 11 | 323 | 3 | -0.124 | 0.709 | -2.753 | 2.968 | 0.846 |
| 9 | p~- | | 1 | -2212 | 3 | 0.395 | -0.614 | -3.806 | 3.988 | 0.938 |
| 10 | pi- | | 1 | -211 | 3 | -0.013 | 0.146 | -1.389 | 1.403 | 0.140 |
| 11 | pi+ | | 1 | 211 | 4 | 0.109 | -0.456 | 2.164 | 2.218 | 0.140 |

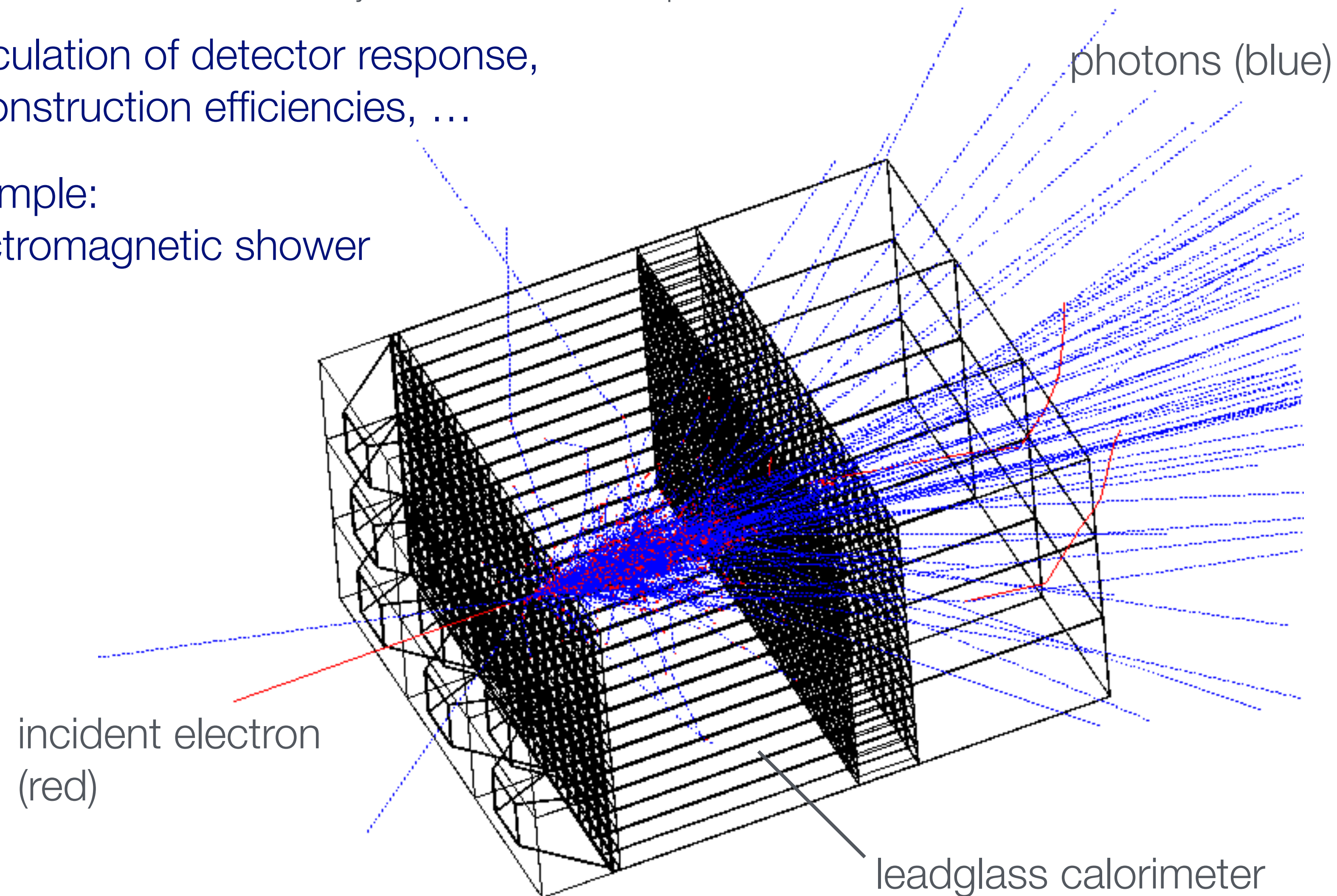
Monte Carlo simulation II: Detector simulation with GEANT

<http://geant4.cern.ch/>

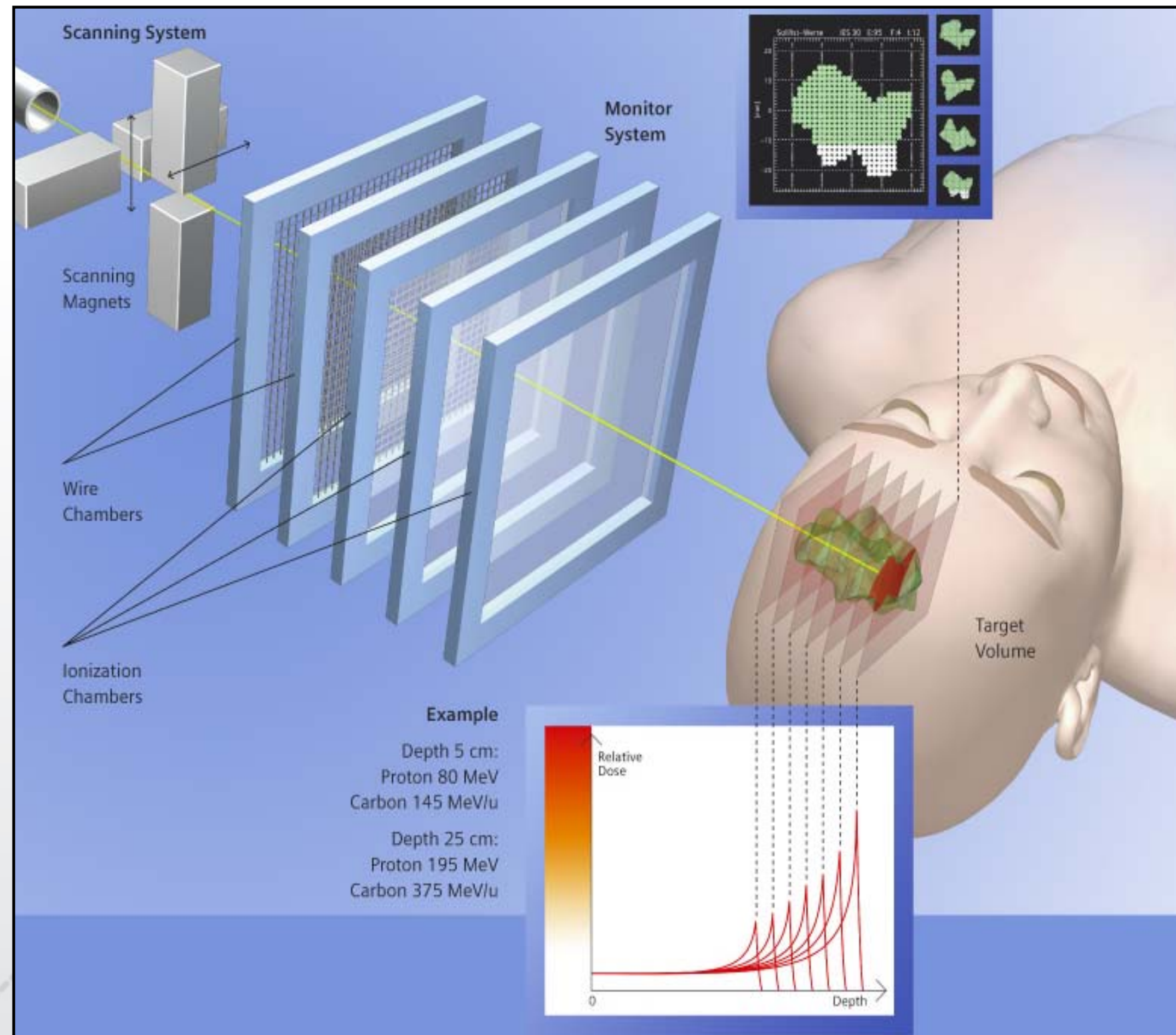
<http://www.uni-muenster.de/Physik.KP/santo/thesis/diplom/kees>

Calculation of detector response,
reconstruction efficiencies, ...

Example:
electromagnetic shower

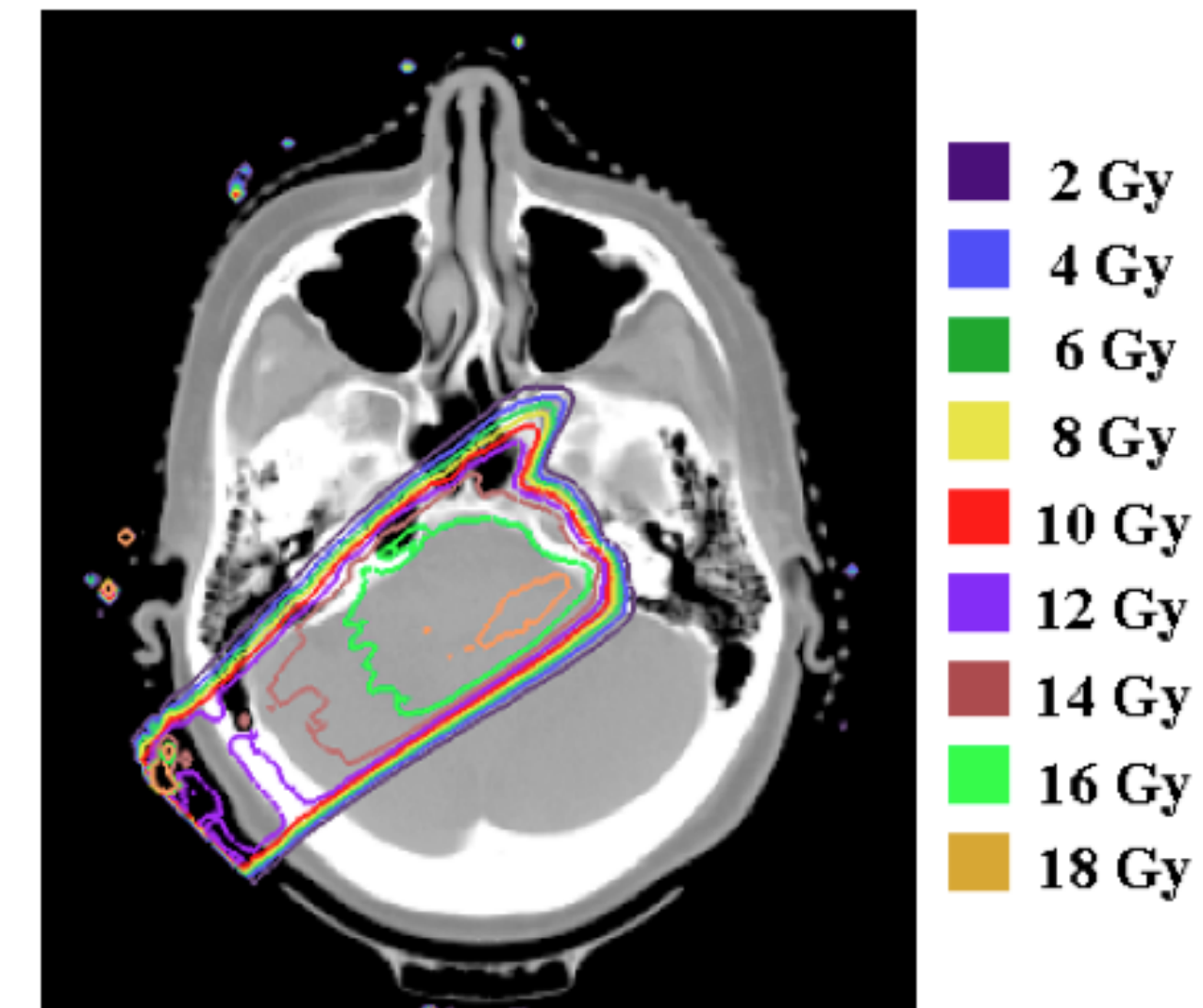


Monte Carlo simulation III: Treatment planning in radiation therapy



Intensity-Controlled Rasterscan Technique, Haberer et al., GSI, NIM A, 1993

Source: GSI



Codes

- ▶ GEANT 4
- ▶ FLUKA
- ▶ ...

