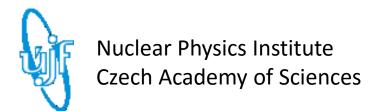
Reverse tracing in SIMRES & McStas Towards faster ray-tracing simulations

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Variance reduction in MC ray-tracing

Ray-tracing employs the *rejection sampling* method, which suffers from very low event probability, p ... becomes worse with increasing dimension

Relative variance for binomial distribution:

$$\frac{\sigma}{p} = \sqrt{\frac{1-p}{Np}}$$

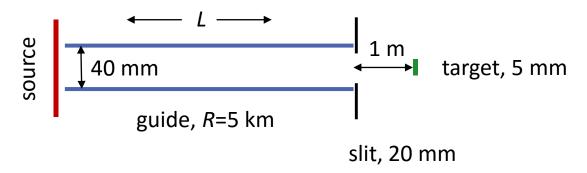
p can be increased by optimization of the sampling volume (typically n-dimensional box, $n \ge 6$)

What is the optimum sampling volume?

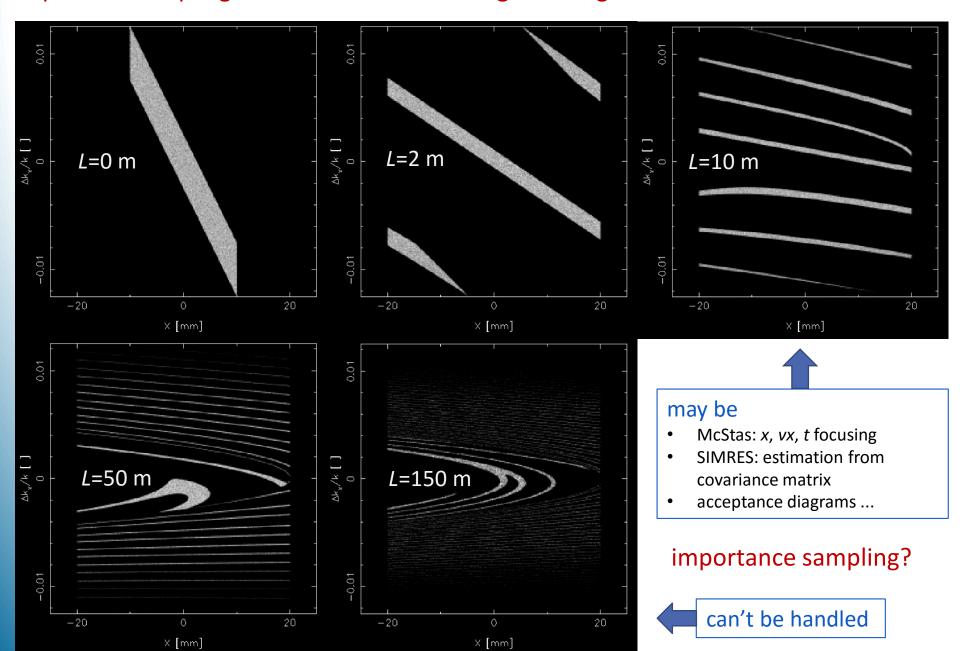
Consider simple 2-dimensional example:

Usual situation: sample << source

How to choose x, v_x at the source to hit the sample?



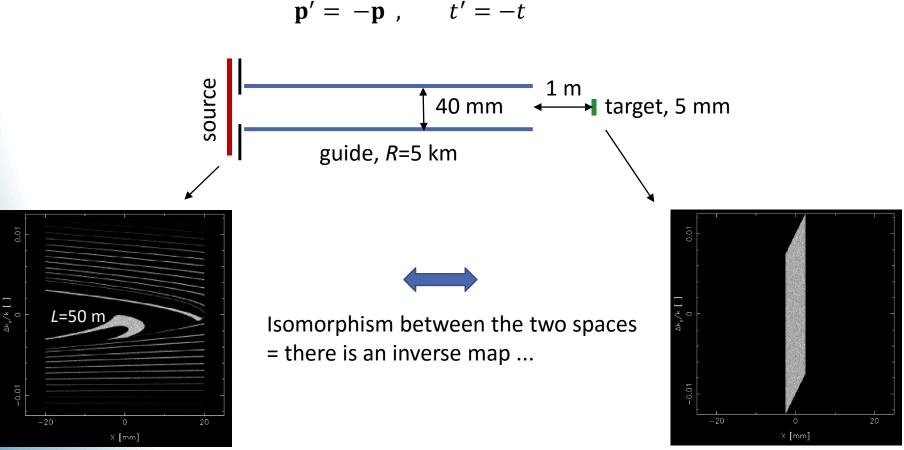
Optimum sampling volume as a function of guide length



Reversibility of ray-tracing

Reversibility is guaranteed by the invariance of underlying physical laws with respect to time inversion:

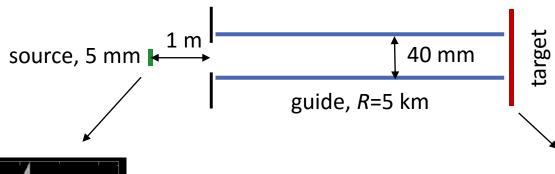
$$\mathbf{p}' = -\mathbf{p} , \qquad t' = -t$$

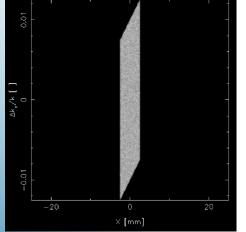


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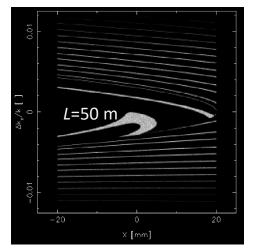
$$\mathbf{p}' = -\mathbf{p}$$
 , $t' = -t$







Isomorphism between the two spaces = there is an inverse map ...



Solutions

- allow tracing in any direction (SIMRES)
- or build an inverted instrument model (McStas)

Variance reduction by reverse ray-tracing

Prerequisites for true reverse tracing

- SW components must handle tracing in any direction
- Neutron generator must be decoupled from the moderator.
- Simulation kernel must handle the **p**, *t* reversal.

Not implemented in McStas



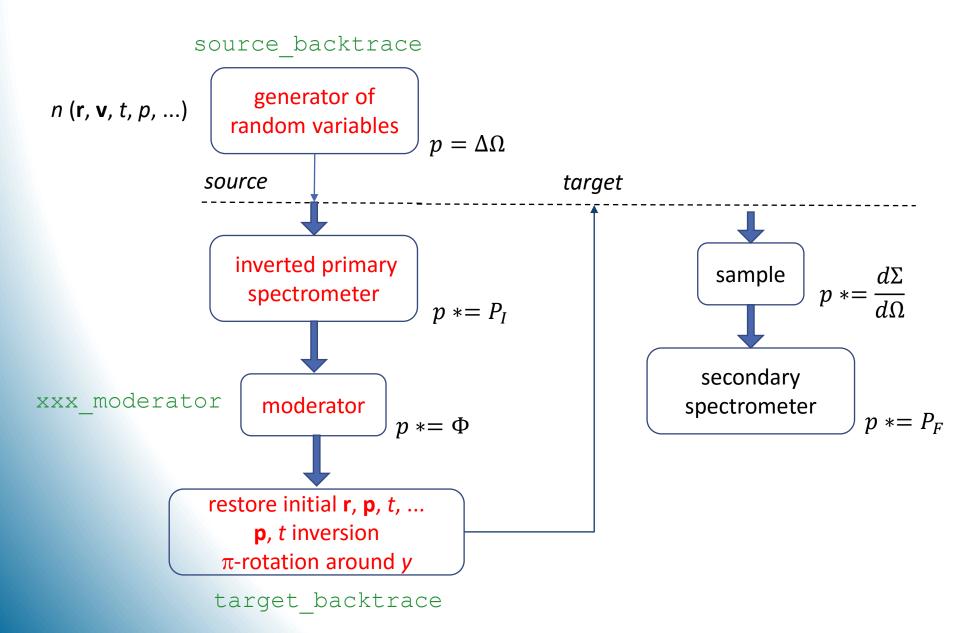
the workaround solution requires

- instrument inversion
- two special components for tracing reversal
- modification of moderators

Work done in early 2016:

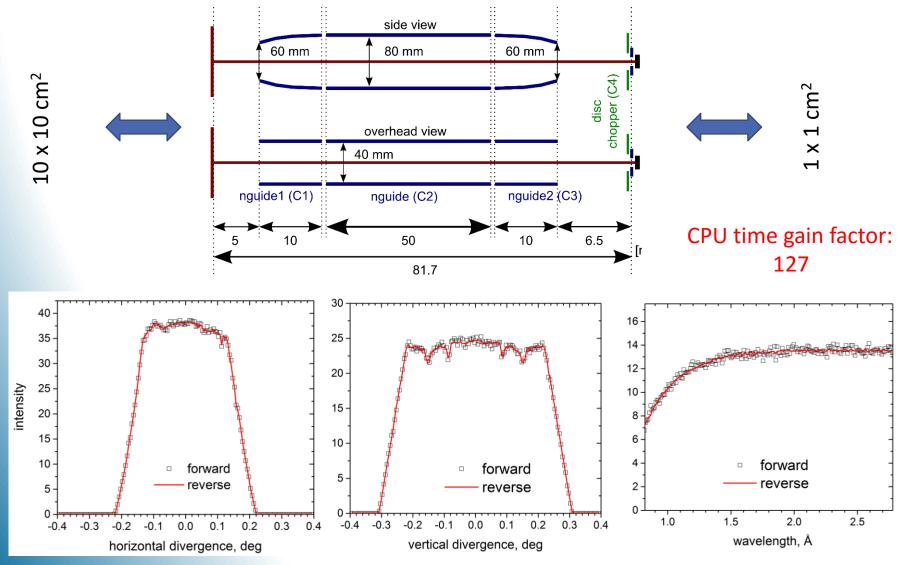
Simple example for McStas

- Includes required new components and instrument files for testing
- Can be used as a template for real work
- Presented at the last WP8 meeting



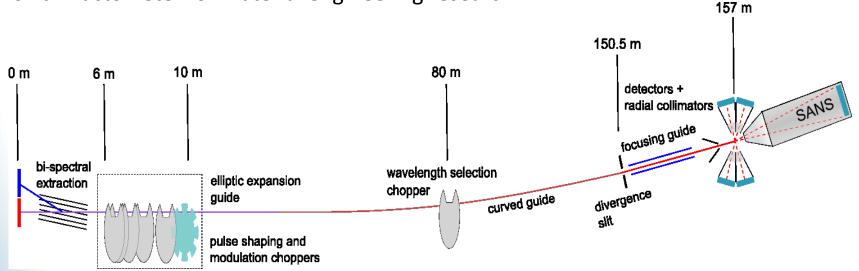
Simple instrument example

- includes commented source code
- short manual to be finished



Application to BEER@ESS

ToF diffractometer for material engineering research



McStas model:

- includes 35 components
- primary spectrometer written in two versions: direct and inverted (both are defined by the same set of input parameters)
- 2 adapted components (ESS moderator and diffraction monitor)
- 1 new component (semi-transparent multichannel guide)

Application to BEER@ESS

Added code:

ESS Moderator backtrace.comp

plus ess_source-lib-NPI (extension to the McStas library)

Provides ESS moderator brightness functions without actually generating neutrons (just adding weights). Only version 2014 (pancake) is done, implementation of the butterfly model is planned after the McStas update release.

PSDcyl_dhkl_monitor.comp

Adapted PSDcyl_monitor.comp, generates diffractograms from detected neutron positions in 3D (x, y, t).

Runtime parameters for high-level instrument control:

```
DEFINE INSTRUMENT BEER(lam0 = 1.8, dlam = 1, frac = 1, int operating_mode = 6, Guide_curvature = 0.06, int verbose = 0, S1_width = 0.040, S1_height = 0.0684, S2_width = 0.040, S2_height = 0.0502, S3_width = 0.005, S3_height = 0.010, S3_dist = 0.050, int GF3a_on = 1)
```

TRACE

```
%include "input_parameters.instr"
// choose tracing direction:
//%include "BEERforward.instr"
%include "BEERreverse.instr"
%include "secondary part.instr"
```

Other parameters are provided in a file

TRACE body is just an envelope which allows easy choice of tracing direction

Test with γ -Fe sample (Fe_Gamma.laz)

Conditions: *cylindrical sample*: 7 x 20 mm²

input slit: 5 x 10 mm², 50 mm before the sample

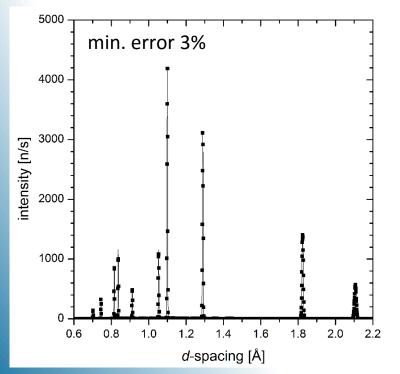
output radial collimator: 4 mm resolution (fwhm)

resolution mode: medium, $\Delta d/d=0.3\%$

detector coverage: $2\theta = 75^{\circ} - 105^{\circ}$, $\phi = \pm 15^{\circ}$

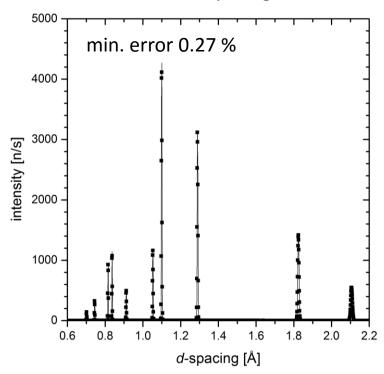
forward

trials: 5 x 10⁹, computing time: 12 h



reverse

trials: 5×10^8 , computing time: 4.2 h



Test with γ -Fe sample (Fe_Gamma.laz)

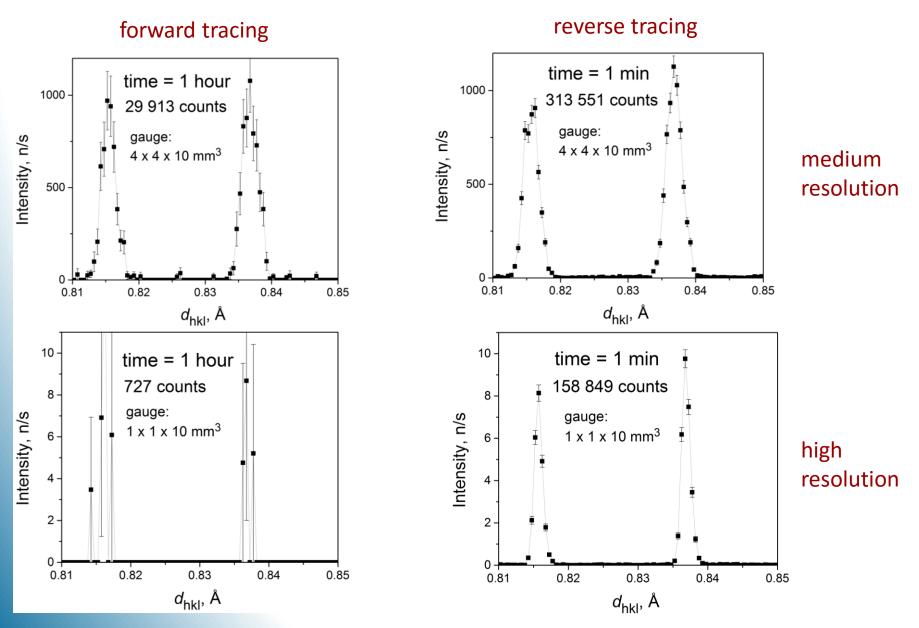
Simulation statistics (counts at the sample)

	forward	reverse	gain factor
Trials / 10 ⁶	5000	500	
Counts / 10 ⁶	0.18	24	
Efficiency [%]	0.0037	4.7	1300
Accuracy [%]	0.34	0.031	
Time [h]	11.95	4.15	
Time to 0.3 % accuracy [h] (*)	9.2	0.045	200

^(*) corresponds to about 20 000 counts in the strongest peak, using SPLIT 20

NOTE: The gain would be much higher for high resolution setup with small gauge volume of ~ 10 mm³ (typical for strain mapping experiments)

Test with γ -Fe sample (Fe_Gamma.laz)



Drawbacks of the present implementation in McStas

- no "user friendly" solution
- requires construction of an inverted primary spectrometer
- all source components need to be modified (this is sometimes quite easy)
- monitors within the primary spec. do not show true flux
 BUT: they <u>could</u> show "useful" neutrons = good for instrument optimization

Better solutions on the level of McStas compiler to be discussed ...

Advantages:

- ✓ gain about 2-3 orders of magnitude in CPU time
- ✓ speed independent of sample size!
- enables virtual experiments with small samples (e. g. strain scanning)
 - + optimization of primary spec. based on scattering data

Thank you for your attention