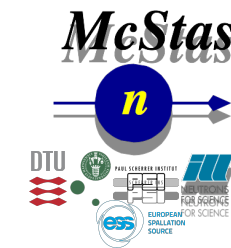


Practical J, McStas samples



MDANSE 2018

Simulation of Inelastic
Neutron Scattering
using McStas and
material dynamics models

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Agenda

- *Structure*
 - *PowderN*
 - *Single_crystal*
 - *Union*
- *Small angle scattering*
- *Connections to other pieces of software*



PowderN exercise

In this exercise we will try to put two powder samples together in a few ways and compare the results.

- 1) Two samples in the same spot, and stochastically choose between them*
- 2) Two samples stacked vertically*
- 3) Two samples where one is behind the other.*





PowderN exercise

Let's use the *PSI_DMC* instrument as a starting point. We will now make the simulation randomly choose between two powders.

- 1) Add another powder in the same spot as the one already there.
- 2) Add **double r**; inside the **DECLARE** section of the instrument file.
- 3) Add an Arm in front of the first one, and add to it an **EXTEND**-block. Add the following code in it: **r=rand01()** ;
- 4) Now add the following before the AT on the two powders.
WHEN (r<0.5) and **WHEN (r>0.5)** respectively.
- 5) Run the instrument again – Do you get what you expect?
- 6) What would you change to make the mixing factor $\neq 0.5$?





PowderN exercise

Let's change this to have two samples on top of each other.

- 1) Make a new copy of the instrument (or remove the edits you did before, leaving the second Powder sample in place).*
- 2) Change the y-position and size of the samples to be $\pm \text{height}/2.0$ and $\text{height}/2.0$ respectively*
- 3) Add the statement **GROUP sample** after the AT at both samples. (N.b. sample is a name chosen arbitrarily. It has to be different than the component names though.)*
- 4) Run a simulation – is there any difference to the previous result? Why/Why not?*





PowderN exercise

Move samples around such that one is in front of the other.

- 1) Run a simulation – Do you still see the signatures of both samples?*
- 2) Do you remember why this can be?*
- 3) How can we get around this?*



Intermission:

A quick trick to remove the direct beam

- If your monitor also can be hit by the direct beam, “swamping” the signal, you can do this:*

*Add the following code **just after** your sample code:*

EXTEND

```
% {  
    if (!SCATTERED) {ABSORB; }  
% }
```

- This will terminate **all** rays which the sample-code has not flagged as scattered. Bear in mind the McStas definition of scattered includes many things (guide-wall reflections etc.)*



SANS models

- Get a new instrument: We'll use the ISIS_SANS2d
- We'll explore some of the SANS-options that exist in McStas.
- The SANS2D-model does not include a sample, nor detector at present, so we'll add that in. Add Ls2d and RR to your **DEFINE INSTRUMENT** line in the top of the file to be able to scan them.

1) Add a PSD_monitor to replicate the detector:

```
COMPONENT detector_front = PSD_monitor(
    nx = 200, ny = 200, filename = "PSD.dat",
    xwidth = 1, yheight=1, restore_neutron=1)
AT(0,0,Ls2d) RELATIVE sample.
```

2) Add a sample – First we'll use the the standard SANS_spheres comp:

```
COMPONENT sample = SANS_spheres(
    xwidth=0.01, yheight=0.01, zdepth=0.001,
    focus_xw=1, focus_yh=1, target_z=Ls2d,
    R=RR, Delta_rho=0.6)
AT (0, 0, 0.2) RELATIVE psd4
```

Now run a simulation to see what you get.





SANS models

- Now we shall explore the possibilities that SasView_models can give us. Change the sample to a SasView_models.comp. Keep the target parameters but change the last line to:
`model_index=51, model_pars={1,7,50}`
- The model_index indicates which of a large number of models scattering functions) the sample is using. You can use mcdoc to find out which, and what the parameters mean. Try to generate a picture similar to the one before.
- Try a couple of the other SasView_models.
`ex. model_index=23, model_pars={50,0.2}`





SANS models

- *2D SANS images are all fine – but wouldn't it be better if we could let the computer do the radial integration for us?*

*recipe: Insert another monitor – for instance a **Monitor_nD** with **options="radius ..."***

*or an instance of **SANSQ_monitor***

- *Remember to use **mcdoc** – it's your friend!*





Union exercise

- Start the McStas GUI (`"mcgui"`) and `"Open"` (a copy) of `"Union_test_absorption_image.instr"`
- Run the instrument as is – to get a starting pt.

This is an image of a sample stick inside a cryostat.

The compilation step takes a bit of time so be patient.

- Open the instrument file and, by looking at the definition of the Al-material, change the cryostat drum material to be made of Iron. (Tip: take a peek at the data file `"Fe.laz"`. It resides inside `"/usr/share/mcstas-2.4.1/data"`)
- Try to add a sample that diffracts strongly (NaCaAlF?) and see if you can catch some powder diffraction on a detector. You may have to add a new monitor with a different size and shape to see it.

