Quick tutorial on MC-Glauber based centrality determination procedure in MPD (NICA)

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March 2021

1 Glauber Monte Carlo

1.1 Installation

The latest version of Glauber Monte Carlo (MC-Glauber) package is available here: https://tglaubermc.hepforge.org/downloads/

To download it on the cluster one can use wget command:

```
wget https://tglaubermc.hepforge.org/downloads/?f=TGlauberMC-3.2.tar.gz
```

To unpack and clean up the downloaded archive one can use the following command:

```
tar -xf *.tar.gz
rm *.tar.gz
```

The main ROOT script is runlauber_v3.2.C.

1.2 Usage

Firstly, one need to set up ROOT environment:

```
source /opt/fairsoft/mpd/new/bin/thisroot.sh
```

Use following commands to run Glauber Monte Carlo:

```
root -1
.L runlauber_v3.2.C
runAndSaveNtuple(Nev, sysA, sysB, signn)
.q
```

Here Nev denotes number of events, sysA and sysB – colliding nuclei, and signn – inelastic nucleon-nucleon cross section which defines energy of the colliding nuclei.

In general, it is reasonable to generate $(5-20)\cdot 10^6$ events to have enough statistics for the centrality determination procedure. The ratio between MC-Glauber and data statistics is 10/1, so for the centrality determination based on $5\cdot 10^5$ events from the data one must have at least $5\cdot 10^6$ MC-Glauber events.

The first colliding systems on the MPD experiment is planned to be Bi+Bi and Au+Au. Those nuclei must be manually set up in the runlauber_v3.2.C (under the line number 1172):

```
else if (TString(name) == "Au3")
    {fN = 197; fR = 6.5541; fA = 0.523; fW = 0; fF = 1; fZ=79;}
else if (TString(name) == "Bi")
    {fN = 209; fR = 6.75; fA = 0.468; fW = 0; fF = 1; fZ=83;}
```

Here fN denotes mass number, fR – radius of the nucleus, fA – skin thickness of the nucleus, fW – deformation parameter of the nucleus, fF – type of the nuclear density function (F = 1 means 3-parameter Fermi function) and fZ – is the atomic number.

The last parameter signn is set accordingly to the system energy $(\sqrt{s_{NN}})$.

$\sqrt{s_{NN}}$, GeV	4.5	7.7	9.5	11
σ_{NN}^{inel} , mb	29.3	29.7	30.8	31.2

After adding those modification in the code, one can generate MC-Glauber data. For example, to generate $5 \cdot 10^6$ Au+Au events at $\sqrt{s_{NN}} = 11$ GeV, one can use the following command:

```
root -1
.L runlauber_v3.2.C
runAndSaveNtuple(5000000, "Au3", "Au3", 31.2)
.q
```

MC-Glauber will then write gmc-Au3Au3-snn31.2--md0.4-nd-1.0-rc1-smax99.0.root file with TNtuple nt_Au3_Au3 in it. This file will be used in the centrality procedure.

2 Centrality Framework

2.1 Installation

To download Centrality framework from the git one can use git clone command and install the project:

```
git clone https://github.com/FlowNICA/CentralityFramework.git
cd CentralityFramework/Framework/centrality-master/
mkdir build/
cd build/
cmake ...
make
```

2.2 Usage

2.2.1 Preparing the data

To begin centrality determination procedure one needs multiplicity distribution from the data. Template reader CentralityFramework/Readers/MpdDstReader.C can be modified and used to fill histogram with multiplicity distribution from MpdDst file format. Generally, the framework was tested on multiplicity distributions with the following track selection criteria:

- $p_T > 0.15 \text{ GeV/c}$
- $|\eta| < 0.5$
- $N_{hits} > 16$ (for reconstructed data only)
- DCA < 0.5 cm.

2.3 Submitting centrality determination jobs

2.3.1 Configuring config.txt.template

Once the input files from MC-Glauber and data are ready, one can start the procedure. Template of the main script is stored in CentralityFramework/scripts/template/ directory. First, change config.txt.template: put full path to the MC-Glauber file in the first line, name of the TNtuple in the second line (nt_Au3_Au3 or nt_Bi_Bi) and full path to the file with multiplicity distribution from the previous step in the third line. If the name of the multiplicity histogram was modified it should be set in the line 4 of the config.txt.template.

There are 4 parametrizations for fit function available. They set number of ancestors N_a as a function of N_{part} and N_{coll} :

• Default : $N_a = fN_{part} + (1 - f)N_{coll}$

- PSD : $N_a = f N_{part}$
- Npart : $N_a = (N_{part})^f$
- Ncoll: $N_a = (N_{coll})^f$
- STAR : $N_a = \frac{(1-f)}{2} N_{part} + f N_{coll}$.

One can change parametrization in line 14 of config.txt.template.

2.3.2 Configuring parameter.list

Next step is to configure parameter.list file. It has 200 lines containing parameters configuration for each job in the following pattern:

```
f_min:f_max:k_min:k_max:Mult_min:Mult_max
```

Those parameters then parsed and placed into copy of config..txt.template for each job. Parameters f, k are set within ranges 0 < f < 1, 0 < k < 100 accordingly and do not require any changes. So the only change that is required are Mult_min:Mult_max which define multiplicity range within which the fit procedure is applied. To change those values one can use sed command:

```
sed -i "s/20:360/Mult_min:Mult_max/" parameter.list
```

Here Mult_min and Mult_max are chosen based on the multiplicity distribution: Mult_min generally is set to 10 or 20 and Mult_max is set close to the values of maximum of the multiplicity distribution.

2.3.3 Configuring and running start.sh

First, one should change the paths to the temporary directory of their choosing: lines 4, 12, 13 of the start.sh. Then, in line 25, full path to the framework core:

```
CentralityFramework/Framework/centrality-master/
```

In line 32, one can set short and usable name for this centrality determination run. Generally, it contains collision system and beam energy under investigation with additional optional short information. After all necessary modifications are done, one can start the procedure:

```
qsub start.sh
```

In most cases, it takes up to 2-3 hours for all jobs to finish. Results will be stored in the output directory OUT alongside start.sh script.

2.4 Final steps of the centrality determination analysis

After the fitting from the previous step is complete one can finish the analysis.

2.4.1 Finding best fit

First of all, one has to merge all fit results:

```
cd CentralityFramework/scripts/template/OUT/COMMIT/jobid/file/root/
hadd -k -f -j 20 fit_merged.root fit/*.root
```

Then use CentralityFramework/Framework/Chi2.C macro:

```
root -l -b -q Chi2.C'("...../fit_merged.root");
```

The macro may process for several minutes and result with a line:

```
f = 0.34+/-0.122 mu = 0.221908+/-0.197214
k = 41+/-8.243 chi2 = 0.991703+/-0.0813411
```

One has to find the corresponding file in $glauber_qa/directory$. To do so, find the line in parameter.list which contains found f and k parameters - let's say line number N. The resulting best fit then is $glauber_qa/glauber_qa_jobid_N.root$. Chi2.C macro also will result with $Fit_Errors_RPC.root$ file which contains quality of the fit information.

2.4.2 Dividing multiplicity into centrality cuts

This procedure is available once the file with optimal fit is found (see previous step). In file HistoCut.C one should modify the lines 2 and 3 with full path to the file glauber_qa/glauber_qa_jobid_N.root from the previous step. If the name of the multiplicity histogram was changed, it needs to be modified as well in line 5. Once the correct path to the file is set, simply run the macro:

```
root -l -b -q HistoCut.C'(10)'
```

Number of the centrality classes is an argument of the macro. The argument is set 10 which produces 10 centrality classes within 0-100% range (0-10%, 10-20%, ..., 90-100%). Resulting file HistoCutResult.root contains multiplicity distributions for each centrality cuts for both fitted multiplicity function (CentralityClass_Fit) and multiplicity distribution (CentralityClass) as well as the TTree Borders. This TTree contains the following information about each centrality class:

- Ncc number of entry
- MinPercent minimum value of centrality in the given centrality class
- MaxPercent maximum value of centrality in the given centrality class
- MinBorder lower cut on multiplicity for the given centrality class
- MaxBorder upper cut on multiplicity for the given centrality class.

2.4.3 Mapping parameters from MC-Glauber with centrality classes

Similarly to the previous step, in file CentralityClasses.C one has to modify lines 2 and 3 with full path to the HistoCutResult.root and glauber_qa/glauber_qa_jobid_N.root files correspondingly. After that, run the macro:

```
root -l -b -q CentralityClasses.C'(10)'
```

The argument here is the same as for HistoCut.C macro: total number of centrality classes within 0-100% centrality range. Resulting file FINAL.root contains impact parameter (B_VS_CentralityClass), number of participants (Npart_VS_CentralityClass), number of binary nucleon-nucleon collisions (Ncoll_VS_CentralityClass) distributions for each centrality class and centrality dependence of their mean values (B_average_VS_Centrality, Npart_average_VS_Centrality, Ncoll_average_VS_Centrality). Additionally, FINAL.root contains TTree Result which is a direct copy of TTree Borders from the previous step.

2.5 Using centrality classes provided from the framework in the analysis

As was mentioned above, files HistoCutResult.root and FINAL.root have all needed information. Use macro printFinal.C to display this information in a simple and readable way:

```
root -l -b -q printFinal.C'("path-to-FINAL.root");
```

This will print out all needed information for each centrality class. This macro also can save output information in several formats: latex, csv tables and C++ code.

Example of printFinal.C saving in latex table:

```
root -l -b -q printFinal.C'("path-to-FINAL.root","./example.tex")'
```

Example of printFinal.C saving in csv table (compatible with LibreOffice and MS Excel):

```
root -l -b -q printFinal.C'("path-to-FINAL.root","./example.csv")'
```

Example of printFinal.C saving in C++ code:

```
root -l -b -q printFinal.C'("path-to-FINAL.root","./example.C")'
```

After printFinal.C generates output C++ code, one can use Float_t GetCentMult(Int_t) as a function which returns centrality percent (for example, for 0-10% centrality class, the function will return 15.) based on input multiplicity value.