

Computer Tools for Nuclear Physics

Initial state of collision within Glauber model

Krzysztof Piasecki



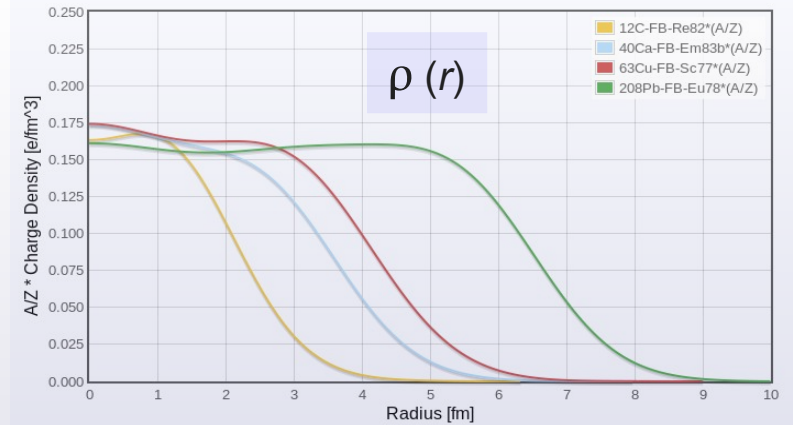
Basics of Glauber Model

- Glauber model: Relativistic Heavy-Ion collision
 - ▶ made of sum of independent NN collisions
 - ▶ density profile is diffused

Simplest density model:
Fermi distribution

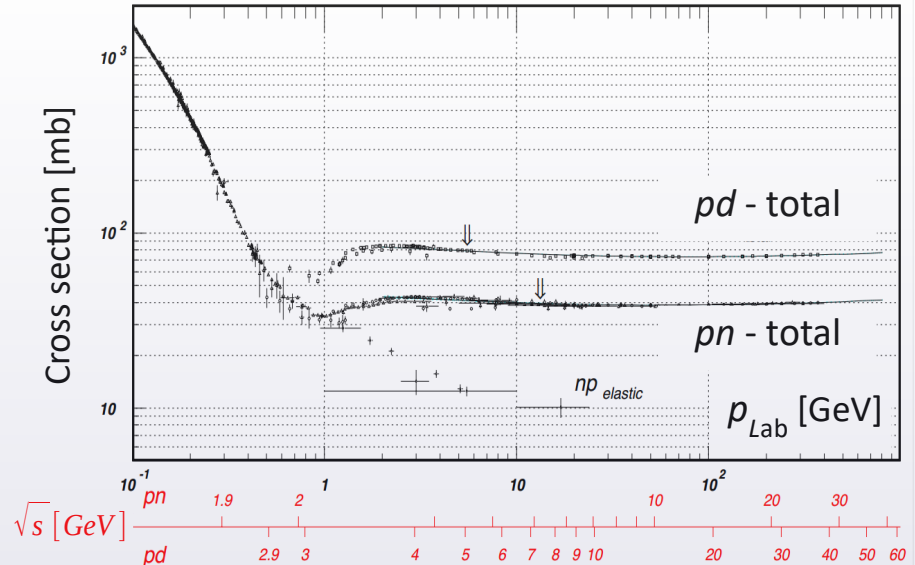
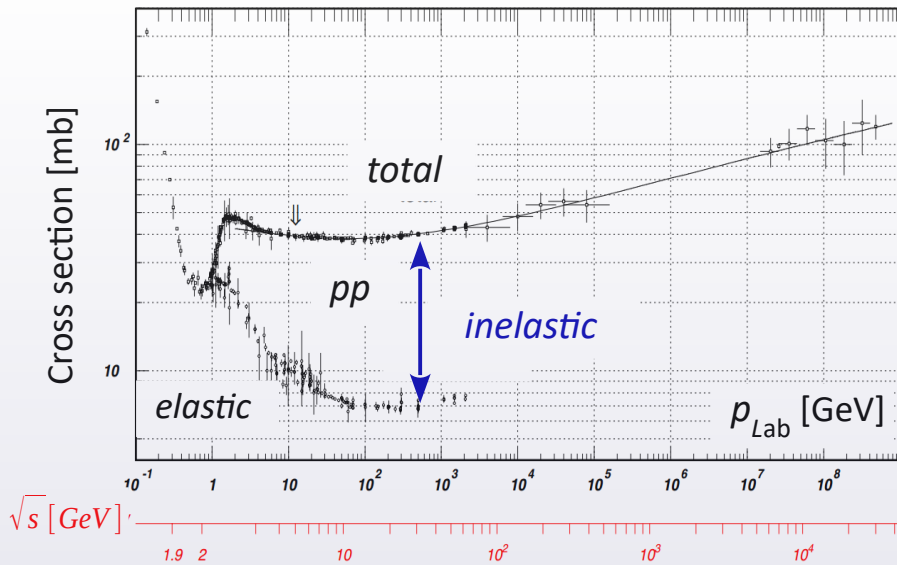
$$\rho(r) = \frac{\rho_0}{\exp\left(\frac{r-r_0}{d}\right) + 1}$$

- Cross section for NN collision is not constant.



faculty.virginia.edu/ncd

journals.aps.org/prd/abstract/10.1103/PhysRevD.86.010001



$\sigma_{\text{NN, inelastic}}$: swing + wide plateau $\sigma_{\text{NN}} \approx 30 \dots 50 \text{ mb}$

Glauber Model: optical approach

- **Introductory paper:** *M. Miller et al. Ann.Rev.Nucl.Part.Sci. 57, 205 (2007)*
Talk: *J. Wilkinson Glauber modelling in hi-ener nucl. coll.*

Optical

continuous $\rho(r)$ distribution
 Analytical formulae,
 Integrals are probed numerically.

Glauber Model

Monte Carlo

Granulation of A+B nucleons.
 Their centers are positioned
 according to $\rho(r)$.

Step 1: average chance for 1 NN collision = ?

- 3D \rightarrow 2D. **Thickness function T_A** [fm^{-2}]:

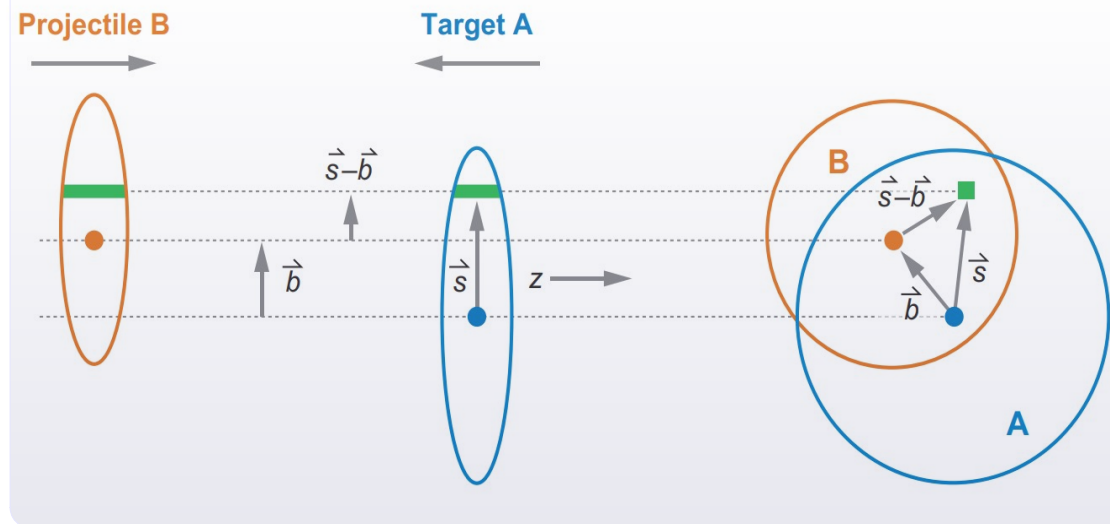
$$T_A(\vec{s}) = \int \rho_T(\sqrt{\vec{s}^2 + z^2}) dz \quad \int \rho_T d^3r = 1$$

(chance of finding 1 nucleon per 1 fm^2 of \perp area)

- **Overlap function T_{AB} :**

$$T_{AB}(\vec{b}) = \int d^2s T_A(\vec{s}) T_B(\vec{s} - \vec{b})$$

(chance that at fixed b ,
 a nucleon from A meets a nucleon of B) [fm^{-2}]



$$\Rightarrow T_{AB}(b) \cdot \sigma_{\text{NN,inel.}} = \text{chance for a single NN collision}$$

Glauber Model: optical approach

- We set up two nuclei at the parameter b . Nucleons move forward and „try to collide”, with success or failure.

Average probability of *single* NN collision is: $T_{AB}(b) \cdot \sigma_{NN}$.

Number of NN collisions = number of „successes” in $A \cdot B$ attempts \leftrightarrow follows the binomial distribution:

$$P(n, b) = \binom{AB}{n} [T_{AB} \sigma_{NN,inel}]^n [1 - T_{AB} \sigma_{NN,inel}]^{AB-n}$$

- Total cross section for nucleus-nucleus (AA) collision. The AA collision occurs if at least 1 NN collision is done:

$$P(\geq 1 \text{ NN collision}) = 1 - P(0 \text{ NN collisions}) \quad \Rightarrow \quad \sigma_{AB,inel} = 2\pi \int_0^\infty \left\{ 1 - [1 - T_{AB}(b) \sigma_{NN,inel}]^{AB} \right\} b db$$

- Number of „binary” NN collisions:

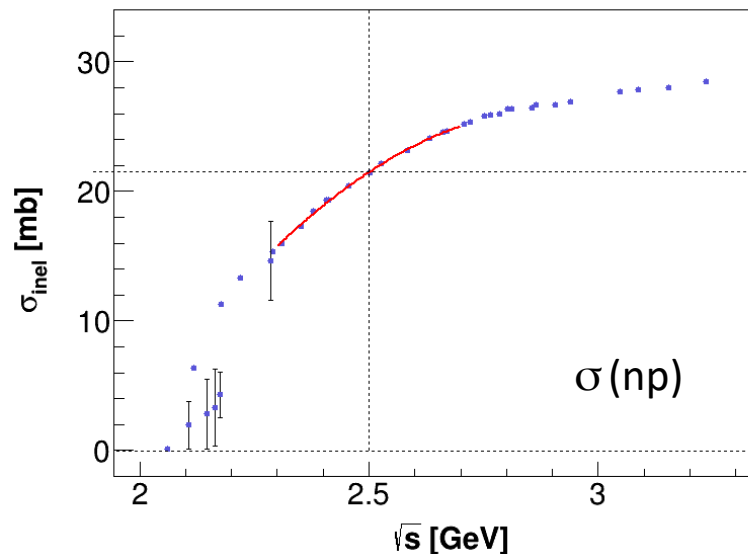
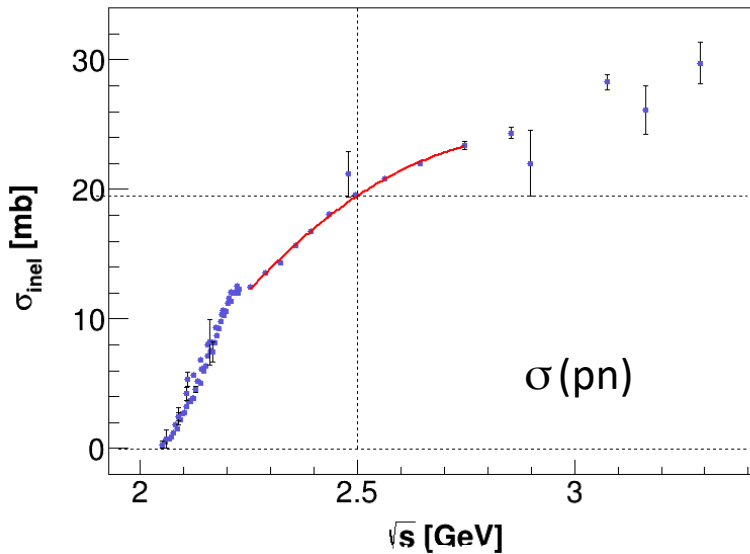
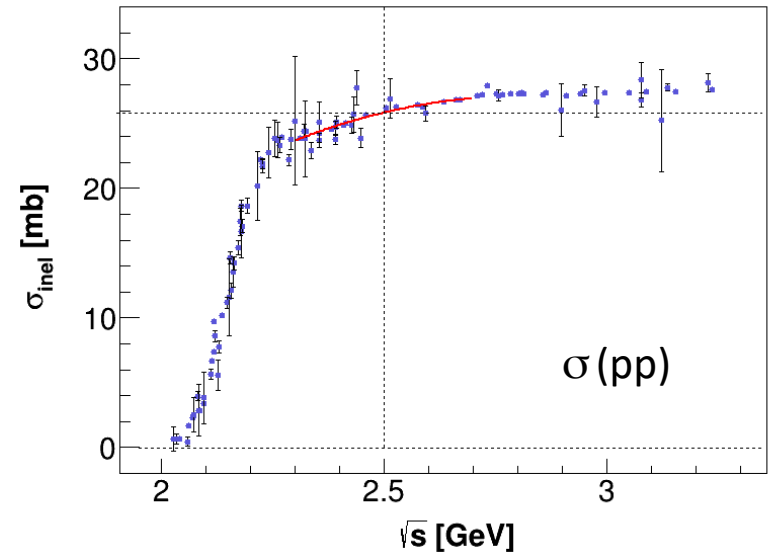
$$N_{coll}(b) = \sum_{n=1}^{AB} n P(n, b) = \dots = AB \cdot T_{AB}(b) \sigma_{NN,inel}$$

- Number of participant nucleons:

$$N_{part}(b) = A \int T_A(\vec{s}) \left\{ 1 - [1 - T_B(\vec{s} - \vec{b}) \sigma_{NN,inel}]^B \right\} d^2s + \\ B \int T_B(\vec{s} - \vec{b}) \left\{ 1 - [1 - T_A(\vec{s}) \sigma_{NN,inel}]^A \right\} d^2s$$

Inelastic NN cross sections

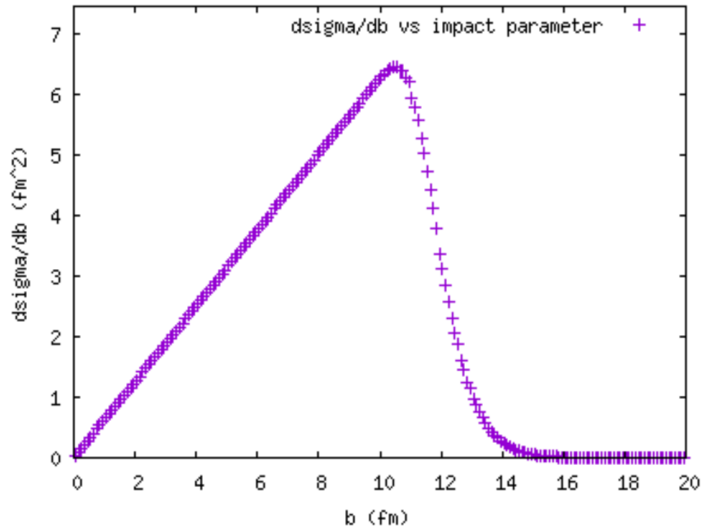
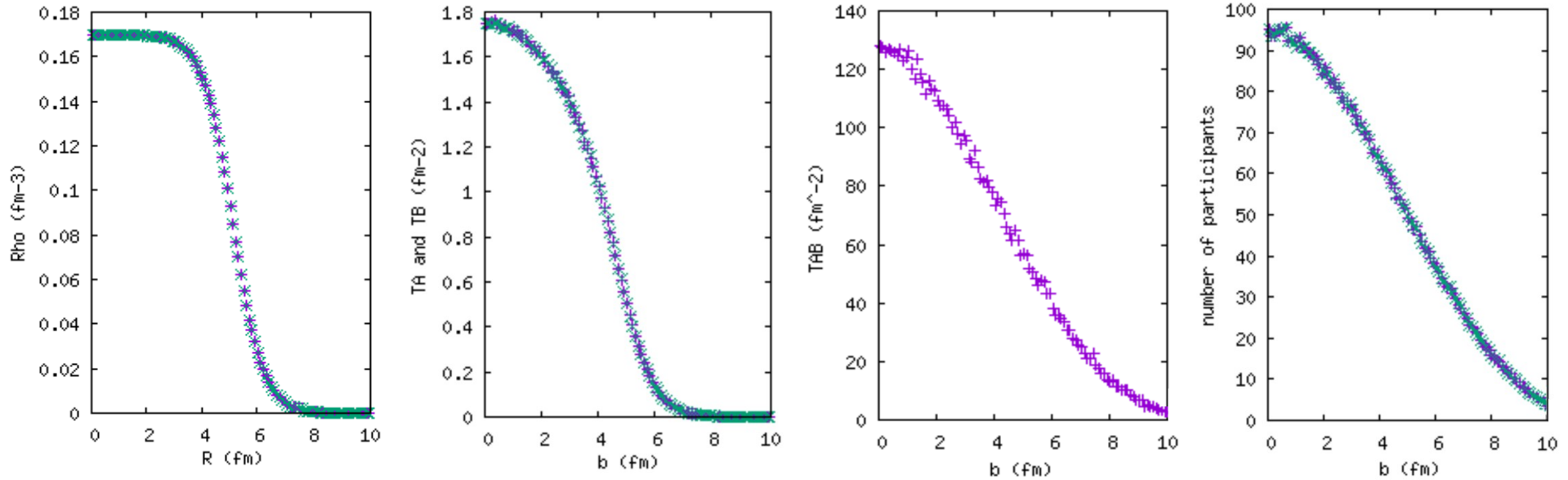
- ① $\sigma(pp)$ is different from $\sigma(pn)$ and $\sigma(np)$
 - ② Assumption: isospin symmetry [$\sigma_{nm} = \sigma_{pp}$]
- $$\sigma_{NN} = \frac{Z_p Z_t \sigma_{pp} + N_p N_t \sigma_{nn} + (Z_p N_t + N_p Z_t) \sigma_{np}}{A_p A_t}$$
- ③ Experimentally, $\sigma(pn)$ is not the same as $\sigma(np)$
 - ④ $\sigma(np)$ at low \sqrt{s} and $\sigma(pn)$ at higher \sqrt{s} are rare
- [③, ④] contribute to systematic errors



- Refs. :
- PDG
 - B. Kardan's Ms. th.

Glauber Model: optical approach

- "Overlap": optical Glauber model online [\[click here\]](#). E.g. for $^{108}\text{Ag} + ^{108}\text{Ag}$ collision (let's assume $\sigma_{\text{NN,inel.}} = 25 \text{ mb}$)



- Let's check how settings can influence $\langle A_{\text{part}} \rangle$ and $\langle N_{\text{coll}} \rangle$:

	Uniform balls	Fermi distrib. $\sigma_{\text{NN}} = 20 \text{ mb}$	Fermi distrib. $\sigma_{\text{NN}} = 30 \text{ mb}$
$\langle A_{\text{part}} \rangle_b$	55	44.5	48.5
$\langle N_{\text{coll}} \rangle_b$	92	65	76

Glauber Monte Carlo: TGlauberMC

- TGlauberMC: ROOT-based simulator of initial conditions of AA collision within the Glauber MC approach.
Authors: from PHOBOS @ RHIC

Homepage: tglaubermc.hepforge.org
Download: www.hepforge.org/downloads/tglaubermc
Prerequisites: Root ≥ 4

Main papers: [No. 1 @ 2008] [No. 2 @ 2019] [No. 3 @ 2019]
User guide: Best description in paper 2.

Applicability: (authors:) $\sqrt{s_{NN}} \in [200 \text{ GeV} \dots 10 \text{ TeV}]$, but used also for $\sqrt{s_{NN}} = 2.5 \text{ GeV}$.

- Installing TGlauberMC on your account of NPD's training computer:

```
$ mkdir tglaubermc ; cd tglaubermc
$ cp -p ~kpiasecki/soft/TGlauberMC/tglaubermc_install.sh .
$ ./tglaubermc_install.sh
```

- Each time you open a new Terminal, do `cd tglaubermc` and then :

```
$ . ./tglaubermc_start.sh
```

 (note: dot at the beginning)

Now if you just launch Root in this directory, you have all the TGlauberMC functions connected.

Glauber Monte Carlo: TGlauberMC

- Currently implemented nuclei: p d t ^3He C O Al Si S Ar Ca Ni Cu Xe W Au Pb U

Basic form of parametrization of nuclear shape:

$$\rho(r) = \rho_0 \cdot \frac{1 + w\left(\frac{r}{R}\right)^2}{1 + \exp\left[\frac{r-r_0}{a}\right]}$$

Shape deformation is an option for Al, Si, Cu, Xe, Au, U :

$$\rho(r) = \rho_0 \cdot \frac{1}{1 + \exp\left[\frac{r - R(1 + \beta_2 Y_{20} + \beta_4 Y_{40})}{a}\right]}$$

Possible nuclei and profiles can be checked or updated in the `TGlaucNucleus::Lookup` method.

- First, b parameter is pulled randomly from Δ distribution. Centers of nuclei are defined as $[-b/2, 0, 0]$ and $[b/2, 0, 0]$ within frame as shown on this Figure.

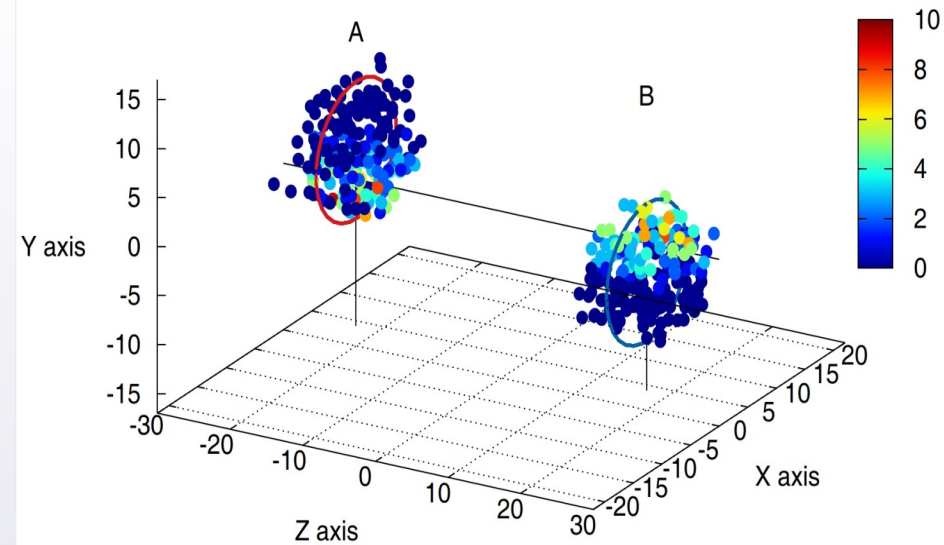
Positions of nucleons are pulled from shape distributions. A minimum distance between N-N centroids is required ($d_{\min} = 0.4$ fm by default).

- Deformed nuclei are rotated randomly by φ and ϑ .

Nucleons are 'traversing' head-on ('eikonal' approximation) and undergo collisions independently.

- A single NN collisions is counted if the distance d between centroids is less than:

$$d < \sqrt{\frac{\sigma_{NN}}{\pi}}$$



Glauber Monte Carlo: TGlauberMC

- Code offers a few classes and macros.
Class `TGlaucNucleon` represents a single nucleon: position, type (p/n), origin (nucleus A/B), No. of collisions,...
Class `TGlauberMC` is the main simulation manager.

- ```
root[1] TGlauberMC gmc ("Pb", "Pb", 30.)
```

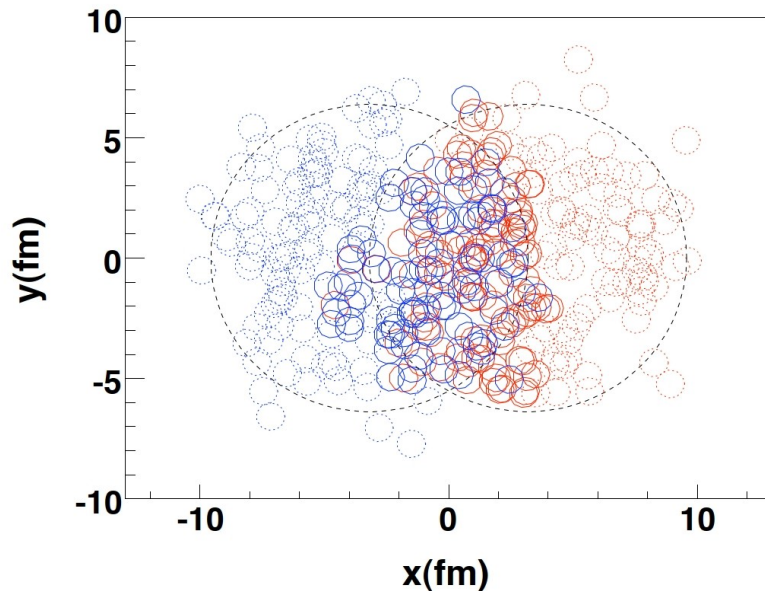
 Initialize Pb+Pb collision with  $\sigma_{NN} = 30$  mb  

```
root[2] gmc.NextEvent (5.)
```

 Simulate 1 collision at  $b = 5$  fm. For -1, random  $b$ .  

```
root[3] gmc.Draw ()
```

 Visualize of the collision (wounded/participants)



- Btw. you can apply range of  $b$  pars:

```
gmc.SetBmin (4.);
gmc.SetBmax (7.);
```

Get (inelastic) collision cross section based on this event:

```
root[4] gmc.GetTotXSect ()
12.57
```

Getting No. of participants and collisions:

```
root[5] cout << gmc.GetNpart() << '\t' <<
gmc.GetNcoll() << endl;
370 786
```

Get array of nucleons:

```
root[6] gmc.GetNucleons ()
```

Simulate e.g. 1000 collisions at once.

```
root[7] gmc.Run (1000)
```

Retrieve event-wise ntuple (can be saved in Root file)

```
root[8] TNtuple* ntu = gmc.GetNtuple ()
```

# Glauber Monte Carlo: TGlauberMC

- **Macro** `runAndSaveNtuple` simulates and stores the resulting event-wise ntuple in file:

```
root [0] runAndSaveNtuple (100, "Ni", "Ni", 25.) simulate 100 Ni+Ni collisions with $\sigma_{NN} = 25$ mb
```

- Variables available in the ntuple include:

|                      |                                            |                          |                                                   |
|----------------------|--------------------------------------------|--------------------------|---------------------------------------------------|
| <code>Npart</code>   | No. of participants                        | <code>AreaA</code>       | Area defined by "and" of participants             |
| <code>Ncoll</code>   | No. of collisions                          | <code>AreaO</code>       | Area defined by "or" of participants              |
| <code>Nhard</code>   | No. of hard-core collisions                | <code>MeanX</code>       | $\langle x \rangle$ of wounded nucleons           |
| <code>B</code>       | Impact parameter                           | <code>MeanY</code>       | $\langle y \rangle$ of wounded nucleons           |
| <code>BNN</code>     | Average NN impact parameter                | <code>MeanX2</code>      | $\langle x^2 \rangle$ of wounded nucleons         |
| <code>Ncollpp</code> | No. of pp collisions                       | <code>MeanY2</code>      | $\langle y^2 \rangle$ of wounded nucleons         |
| <code>Ncollpn</code> | No. of pn collisions                       | <code>MeanXY</code>      | $\langle xy \rangle$ of wounded nucleons          |
| <code>Ncollnn</code> | No. of nn collisions                       | <code>MeanXSystem</code> | $\langle x \rangle$ of all nucleons               |
| <code>VarX</code>    | Variance of X of wounded nucleons          | <code>MeanYSystem</code> | $\langle y \rangle$ of all nucleons               |
| <code>VarY</code>    | Variance of Y of wounded nucleons          | <code>MeanXA</code>      | $\langle x \rangle$ of nucleons in nucleus A      |
| <code>VarXY</code>   | Covar. between X and Y of wounded nucleons | <code>MeanYA</code>      | $\langle y \rangle$ of nucleons in nucleus A      |
| <code>NpartA</code>  | No. of wounded nucleons in nucleus A       | <code>MeanXB</code>      | $\langle x \rangle$ of nucleons in nucleus B      |
| <code>NpartB</code>  | No. of wounded nucleons in nucleus B       | <code>MeanYB</code>      | $\langle y \rangle$ of nucleons in nucleus B      |
| <code>Npart0</code>  | No. of singly-wounded nucleons             | <code>PhiA</code>        | $\phi$ angle nucleus A (applied if deformed)      |
| <code>AreaW</code>   | Area defined by width of participants      | <code>ThetaA</code>      | $\vartheta$ angle nucleus A (applied if deformed) |
| <code>PsiN</code>    | Event plane angle of n-th harmonic         | <code>PhiB</code>        | $\phi$ angle nucleus B (applied if deformed)      |
| <code>EccN</code>    | Participant eccentricity for n-th harmonic | <code>ThetaB</code>      | $\vartheta$ angle nucleus B (applied if deformed) |

- **Caution:** `PsiN` and `EccN` are constructed from weighted positions of wounded nucleons.

# Geometry of wounded nucleons

- Centroids of nuclei are initially fixed to the XZ plane (see Fig. on p. 8) .

Ideally:

- area occupied by wounded nucleons looks like a vertical almond ( $\sim$ ellipse)
- angle  $\psi$  between the longer axis and X axis:  $\psi_{EP} = 90^\circ$
- shape has given eccentricity  $\varepsilon$

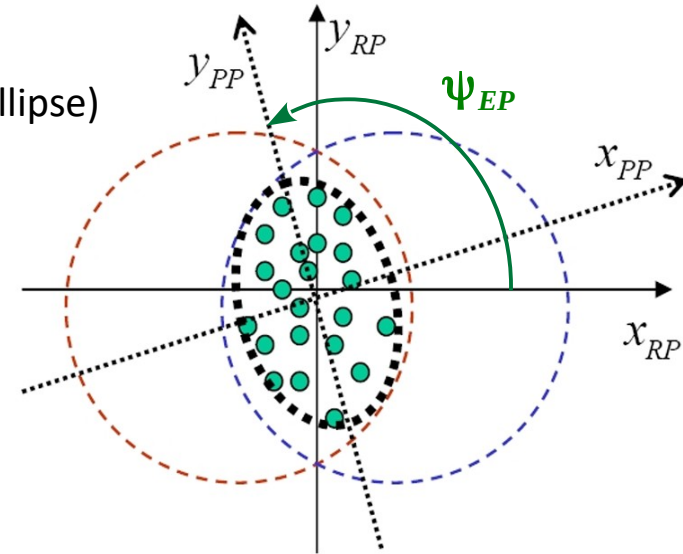
However, in an event, positions of nucleons are pulled randomly.

- shape deviates from an “ideal” almond
- angle  $\psi_{EP}$  differs from  $90^\circ$ .  
A de-facto plane is called “**event plane**” or “**participant plane**”
- shape has given **eccentricity  $\varepsilon$**  (deviated from ideal  $\varepsilon$ )

$\psi_{EP}$  is found from positions  $(x_i, y_i)$  by:

$$\psi_{EP} \equiv \text{atan} \left( \frac{\sum y_i}{\sum x_i} \right)$$

$\varepsilon$  is also found from wounded nucleons:

$$\varepsilon \equiv \frac{\sqrt{\langle r^2 \cos(2\varphi) \rangle^2 + \langle r^2 \sin(2\varphi) \rangle^2}}{\langle r^2 \rangle}$$


[S. Voloshin et al.]

- For more refined analyses of shape and fluctuations,  $\psi$  and  $\varepsilon$  of higher harmonics is available.

$$\varepsilon_n \equiv \frac{\sqrt{\langle r^n \cos(n\varphi) \rangle^2 + \langle r^n \sin(n\varphi) \rangle^2}}{\langle r^n \rangle}$$

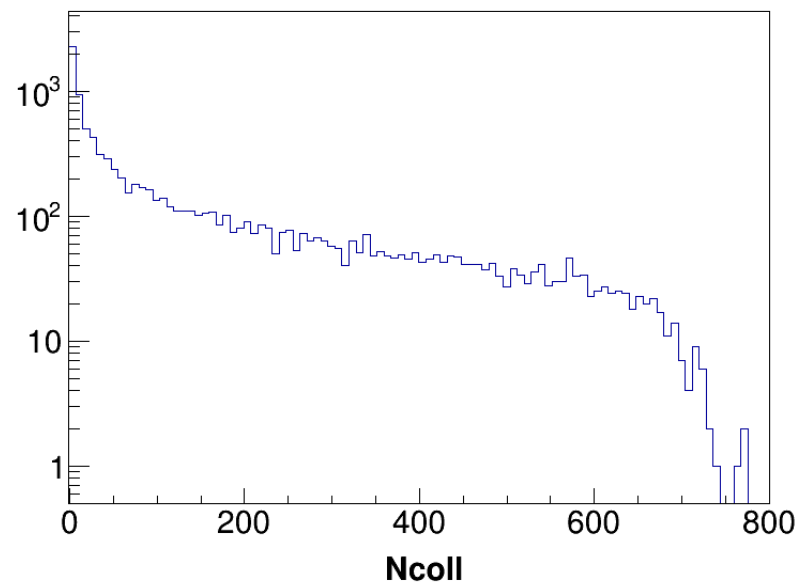
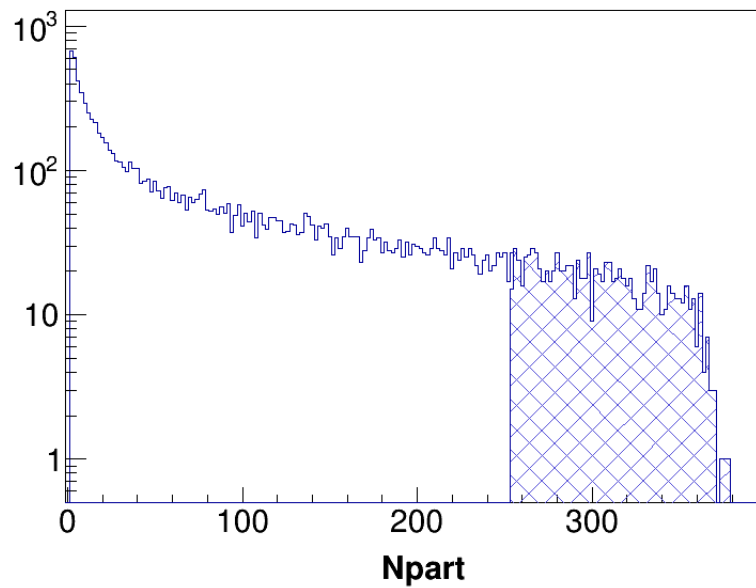
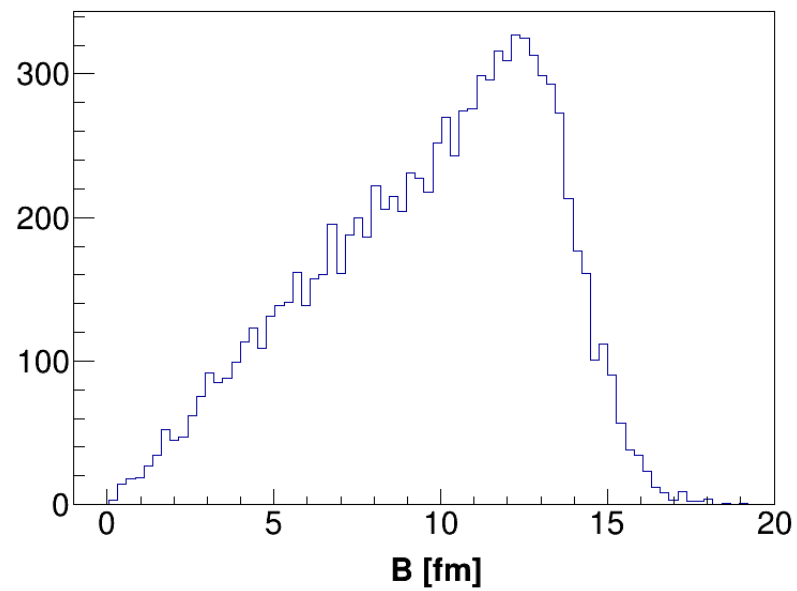
$$\psi_n \equiv \frac{1}{n} \text{atan} \left( \frac{\sum r^n \cos(n\varphi)}{\sum r^n \sin(n\varphi)} \right)$$



[Source: M. Stefaniak]

# Glauber Monte Carlo: TGlauberMC

- Simulation: Au+Au,  $\sigma_{NN} = 23.7$  mb .



# Glauber Monte Carlo: TGlauberMC

- Let's focus on `gmc.GetNucleons ()`. It returns the address to the array of `TGlaunucleon` objects.
- What does a single `TGlaunucleon` object contain?
  - ▶ **position**: You can `SetXYZ` and `GetX/Y/Z`, but also `RotateXYZ (_3D)` it. Also it knows if it `IsInNucleusA/B`.
  - ▶ **type**: `IsProton`, `IsNeutron`, `Get/SetType`. Also energy, but it has no effect.
  - ▶ **collision status**: if it `IsWounded` and how badly: `GetNcoll`. Alternatively, if it `IsSpectator`. You can also "make" it `Collide` or heal fully: `Reset`.

# Glauber Monte Carlo: TGlauberMC

- Looping over nucleons in an event.

```
R__LOAD_LIBRARY (libMathMore)
R__LOAD_LIBRARY (runGlauber_v3.2_C)

int nucloop () {
 TGlauberMC gmc ("Pb" , "Pb" , 25.);
 TH1F hNucColl ("hnucoll", "", 13, -0.5, 12.5);
 TObjArray* nucArray ;
 TGlaucNucleon* nuc ;

 for (int iEvent = 0; iEvent < 100 ; iEvent++)
 {
 gmc.NextEvent (-1);
 nucArray = gmc.GetNucleons ();
 for (int iNuc = 0; iNuc < nucArray->GetEntries(); iNuc++)
 {
 nuc = (TGlaucNucleon*) nucArray->At (iNuc) ;
 hNucColl->Fill (nuc->GetNColl());
 }
 }
 hNucColl->Draw ();
 return 0;
}
```

• Create the simulator of Pb+Pb collision @  $\sigma_{NN} = 25$  mb

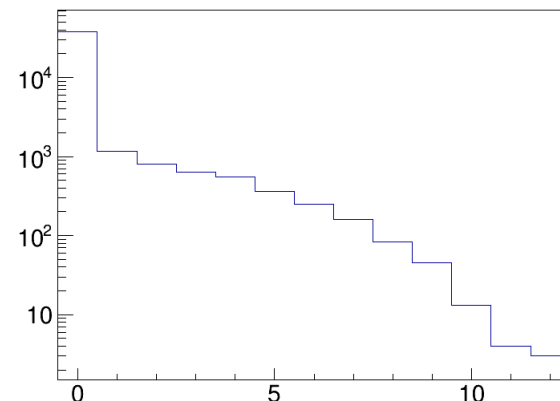
• Loop over events

• Simulate 1 coll. at random  $b$

• Get array of nucleons

• Loop over nucleons

• Get given nucleon



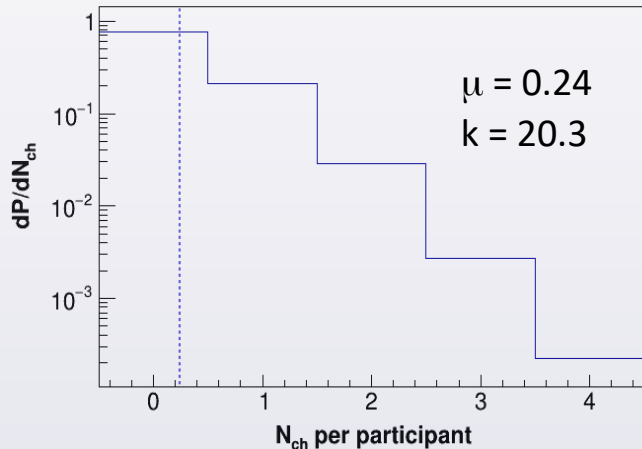
# Glauber Model vs experimental data

- Obtained  $A_{\text{part}}$  distribution is not yet a place of comparison with experiment:
  1. usually only charged particles are measured (protons)
  2. detector acceptance is limited  $\rightarrow$  will measure only part of particles
  3. even if particle gets inside: setup has some inefficiency of track measurement (efficiency  $\equiv \varepsilon$ ).
- $N_{\text{part}}$  distribution from TGlauberMC simulation has to be projected to  $N_{\text{ch}}$  measured particles, Accounting for fluctuations. Each participant is associated with „ $P$  of creating registered particle”.

- One usually applies the **negative binomial distribution, NBD** with mean  $\mu$  and width parameter  $k$ :

$$P_{\mu,k}(n) = \frac{\Gamma(n+k)}{\Gamma(n+1)\Gamma(k)} \cdot \frac{(\mu/k)^n}{(\mu/k+1)^{n+k}}$$

Example for Au+Au @ 1.2A GeV (HADES):



- Plot shows the distrib. of multiplicity of charged particles.

- Glauber MC
- Experiment, "minimum bias"
- Experiment "central trigger"

Good agreement between model and experiment.

Experimentally one defines the „**centrality classes**” on this plot.

