Lead-lead scattering

We describe the configuration file found in examples/

• expl6.optns - PbPb collision with hydro and hadronic cascade

The content of the file is as follows:

```
application hadron !hadron-hadron, hadron-nucleus, or nucleus-nucleus collision
set laproj 82
                !projectile atomic number
set maproj 207
                   !projectile mass number
set latarg 82
                   !target atomic number set matarg 207 !target mass number
set ecms 2760
                   !sqrt(s)_pp
set istmax 25
set iranphi 1
ftime on
!suppressed decays:
nodecays 110 20 2130 -2130 2230 -2230 1130 -1130 1330 -1330 2330 -2330 3331 -3331 end
                         !number of initial conditions used for hydro evolution
set ninicon 1
core full
                         !core/corona activated
hydro hlle
                         !hydro activated
eos x3ff
                          !eos activated
hacas full
                          !hadronic cascade activated
set nfull 2
                         !number of events
                    !number of freeze out events per hydro event
!certain printout every modsho events
! O=min bias
set modsho 100
set nfreeze 1
set centrality 0
fillTree4(C2)
```

The command **application hadron** specifies which kind reaction we simulate (**hadron** stands for all kinds of hadron-hadron scattering, but also hadron-nucleus or nucleus-nucleus),

Then we specify projectile and target, providing the atomic and mass number, and the energy given as center of mass energy (in GeV), using the **set** command.

Some more "technical settings" are needed: **set istmax 25** defines the maximal status value considered for storage, **set iranphi 1** is needed for internal use. In case of iranphi=1 the event will be rotated such that the impact parameter angle and the (n=2) event plane angle (based on string segments) coincide. Particles are rotated back at the end. Finally **ftime on** means that a non-zero string formation time is used (it should only be turned off for testing purposes).

The command **nodecays** allows us to block the decay of the particles π^0 , K_s , Λ , $\bar{\Lambda}$, $\Sigma^-, \bar{\Sigma}^ \Sigma^+, \bar{\Sigma}^+$, $\bar{\Xi}^0$, $\bar{\Xi}^0$, $\bar{\Xi}^-$, Ω^- , (see **src/KWt/idt.dt** for the id codes).

Then we activate all the hydro components (core full, hydro hlle, eos x3ff) and the hadronic re-scatterings simulated with UrQMD (hacas full).

To increase statistics, one might increase nfreeze.

The command fillTree4(C2) writes results into a ROOT file, where C2 means that the number of Pomerons is used as centrality variable. In case of nucleus-nucleus collisions, we usually use C1 for impact parameter as centrality variable.