

Working with PHQMD short introduction

30.09.2024, Susanne GläBel

Run PHQMD: steps

- 1) Run PHQMD code
- 2) Apply stabilisation routine
- 3) Convert PHQMD-output into one file (detector-input)



Parton-Hadron-Quantum-Molecular Dynamics

= n-body microscopic transport approach for the description of heavy-ion dynamics with dynamical cluster formation

PHSD



+ QMD

+ MST

Relativistic considerations

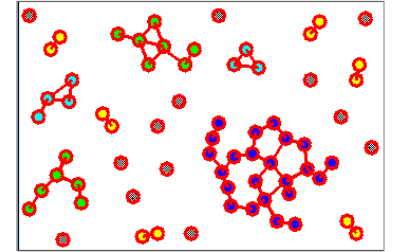
but: mean-field potentials
=> correlations are smeared out

Correlations between baryons

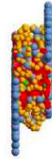
n-body transport approach
=> formation of clusters due to potential interactions

Cluster recognition

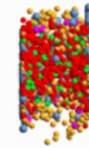
search for accumulations of particles in coordinate space



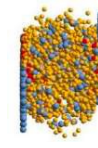
Initial A+A collisions



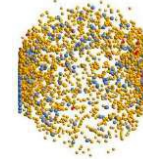
Formation of QGP



Partonic phase



Hadronization



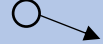
Hadronic phase



QMD

Initialization nuclei

propagation of baryons



local $\epsilon > \epsilon_c$: dissolution of pre-hadrons

propagation of partons

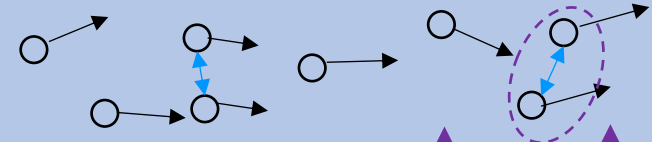
interactions of partons



interactions of hadrons



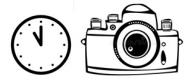
propagation of mesons



PHSD

Primary collisions
pre-hadronic states

MST



PHQMD Cluster recognition

Minimum Spanning Tree (MST)

Cluster formation due to attractive potential between nucleons in **QMD**

Cluster recognition criterion: distance of nuclei

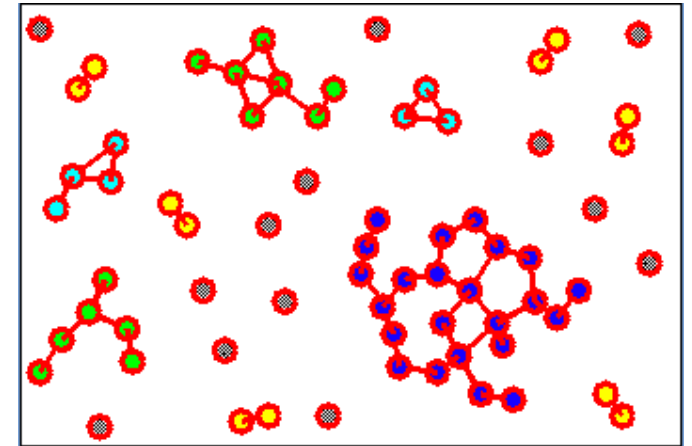
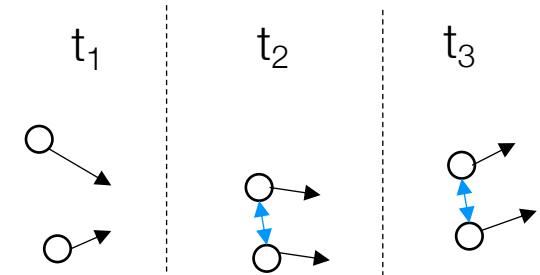
Algorithm: search for accumulations of particles in coordinate space

1. Two particles i & j are bound if:

$$|r_i - r_j| < 4.0 \text{ fm}$$

2. Particle is bound to cluster if bound with at least one particle of cluster

cut on binding energy is applied during conversion





Parton-Hadron-Quantum-Molecular Dynamics

= n-body microscopic transport approach for the description of heavy-ion dynamics with dynamical cluster formation

PHSD



+ QMD

+ MST

Relativistic considerations

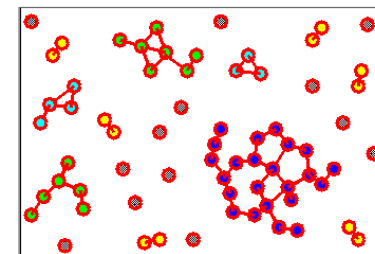
but: mean-field potentials
=> correlations are smeared out

Correlations between baryons

n-body transport approach (Skyrme potential)
=> formation of clusters due to potential interactions

Cluster recognition

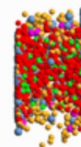
search for accumulations of particles in coordinate space



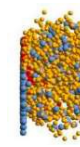
Initial A+A collisions



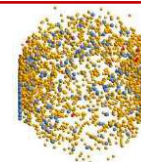
Formation of QGP



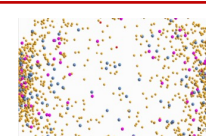
Partonic phase



Hadronization



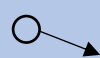
Hadronic phase



QMD

Initialization nuclei

propagation of baryons



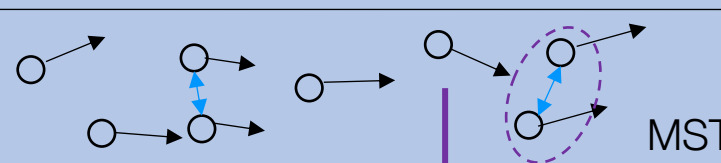
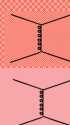
local $\epsilon > \epsilon_c$: dissolution of pre-hadrons

propagation of partons

interactions of partons

interactions of hadrons

propagation of mesons



MST

PHSD

Primary collisions
pre-hadronic states

PHQMD OUTPUT:

Clusters in fort.791 & Anticlusters in fort.781

Hadrons in phsd.dat

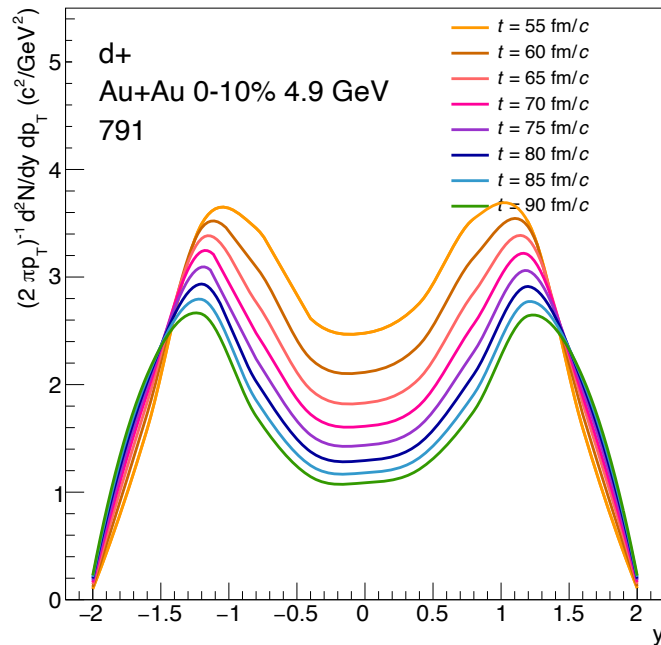
Run PHQMD: steps

- 1) Run PHQMD code
- 2) Apply stabilisation routine
- 3) Convert PHQMD-output into one file (detector-input)

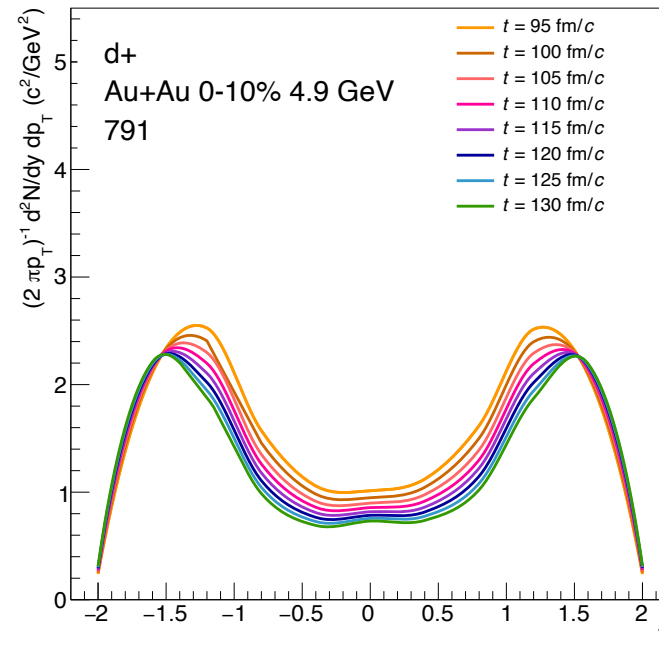
MST clusters for different times during the HIC

Deuteron y-distribution fort.791

$t = 55 \text{ fm/c} - 90 \text{ fm/c}$



$t = 95 \text{ fm/c} - 130 \text{ fm/c}$



Decreasing of cluster multiplicity:

- at early times: baryons collide with other baryons & escape cluster
- at all times: artificial dissolving of clusters, because:

QMD is a semiclassical model:

Skyrme potential is not relativistic

→ different times in computational frame and centre of mass frame of clusters

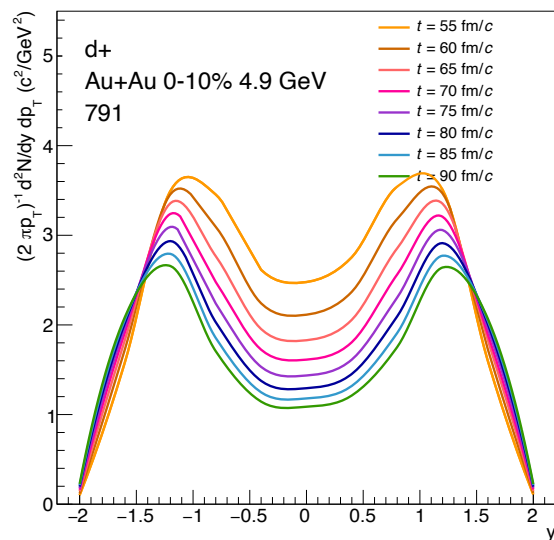
→ numerical artifact when calculating potentials for propagation

→ Stabilisation routine to overcome artificial dissolving

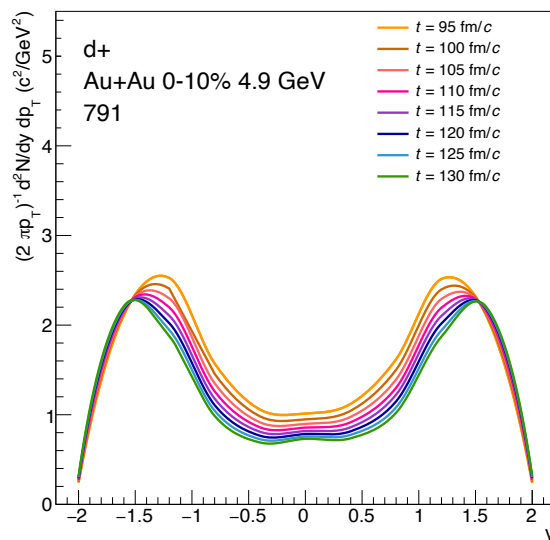
Stabilization of cluster output

Before stabilisation

$t = 55 \text{ fm/c} - 90 \text{ fm/c}$

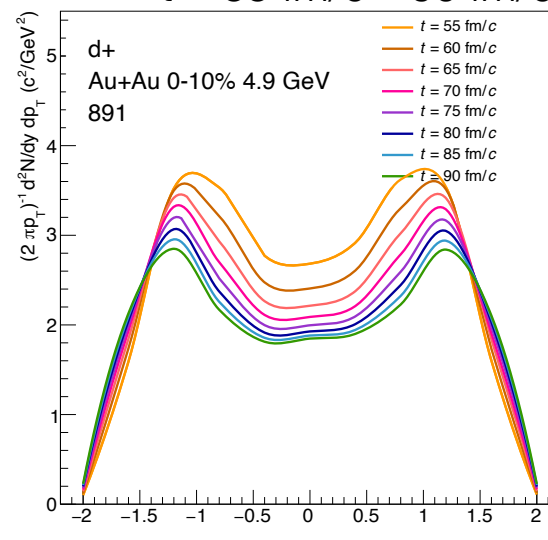


$t = 95 \text{ fm/c} - 130 \text{ fm/c}$

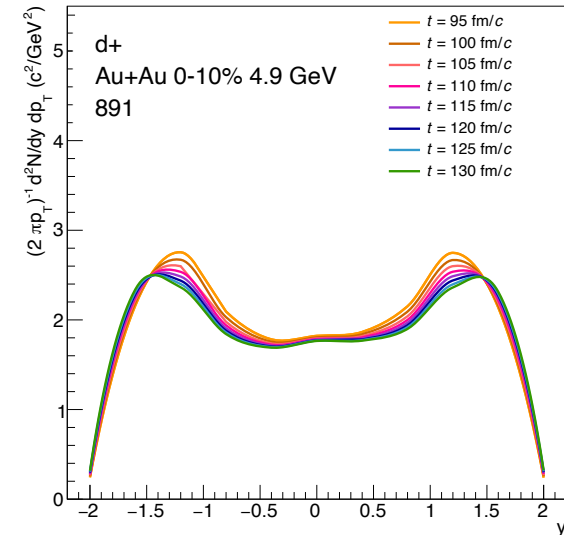


With stabilisation procedure (aMST)

$t = 55 \text{ fm/c} - 90 \text{ fm/c}$



$t = 95 \text{ fm/c} - 130 \text{ fm/c}$



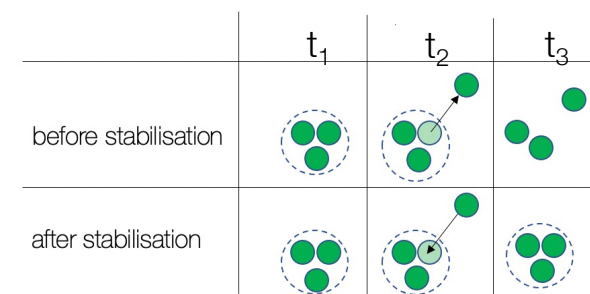
Stabilisation procedure Advanced MST (aMST):

Consider a cluster as stable after all baryons in the cluster are frozen

- 1) Check if dissolving of a cluster was artificial, i.e. if all cluster-baryons had their last collision.
- 2) Restore cluster.

fort.791 &
fort.781 ↓

fort.891 &
fort.881



Run PHQMD: steps

- 1) Run PHQMD code
- 2) Apply stabilisation routine
- 3) Convert PHQMD-output into one file (detector-input)

Convert PHQMD output into detector input

- A) Get hadrons from phsd.dat (& remove hadrons bound in clusters)
- B) Get clusters from fort.891 / 881

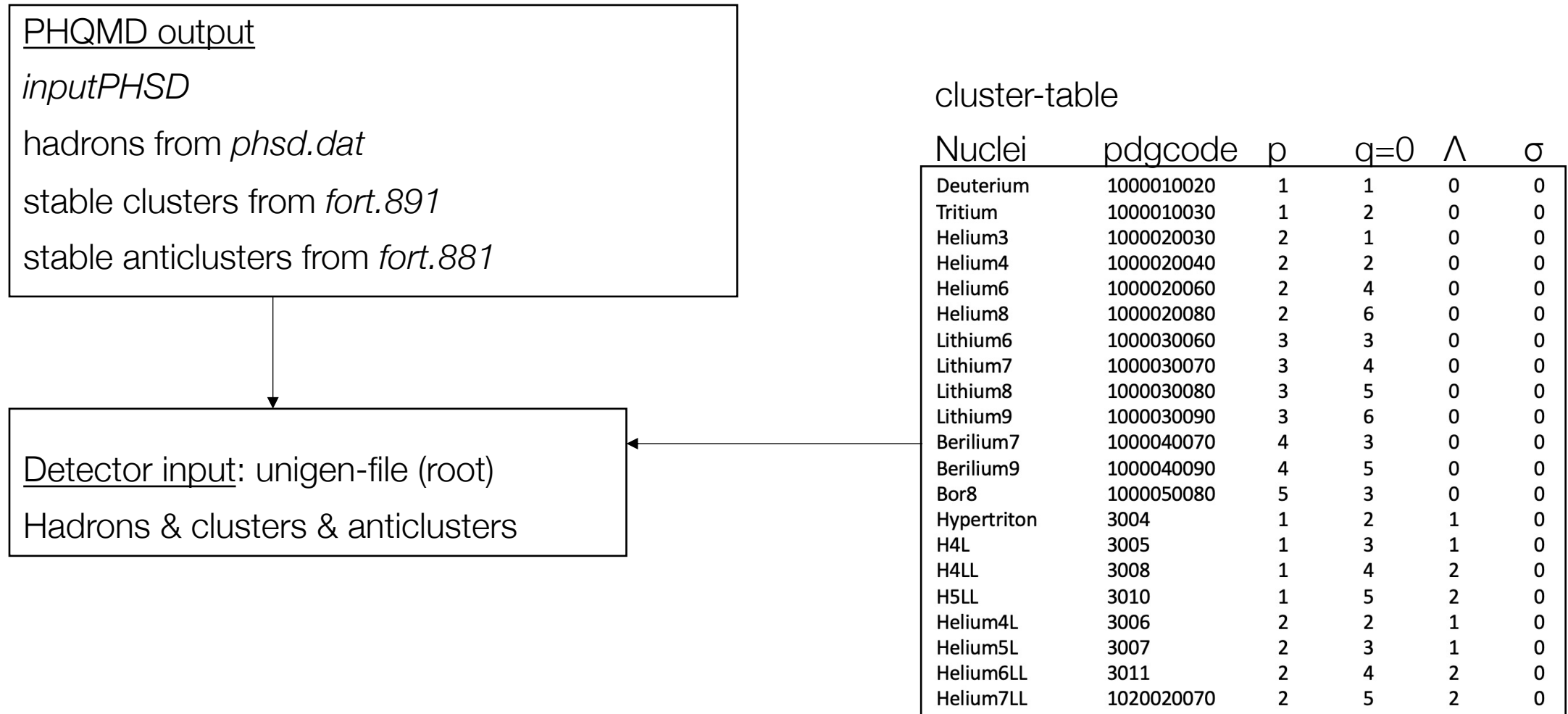
MST recognises clusters / anti-clusters independently of their physical existence

Conversion:

- 1) Check if cluster binding energy > 0
 - 2) Check if cluster is physical (cluster table $A < 9$)
& if not, count baryons as single baryons
- options for big clusters ($A > 7$):
- count independent of physical existence
 - don't count



Convert PHQMD output into detector input



The cluster_table.root can be modified by changing cluster_table.dat and by recreating it with write_clustertree.C.

UniGen output (root)

UEvent

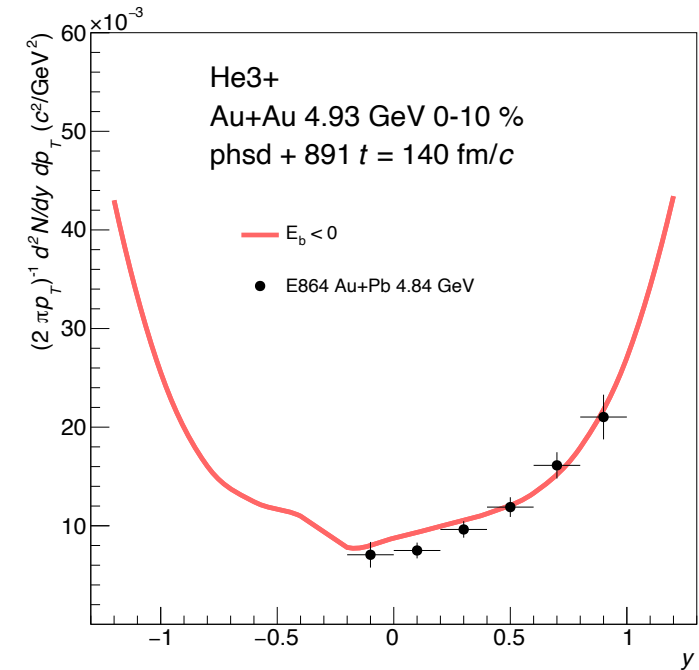
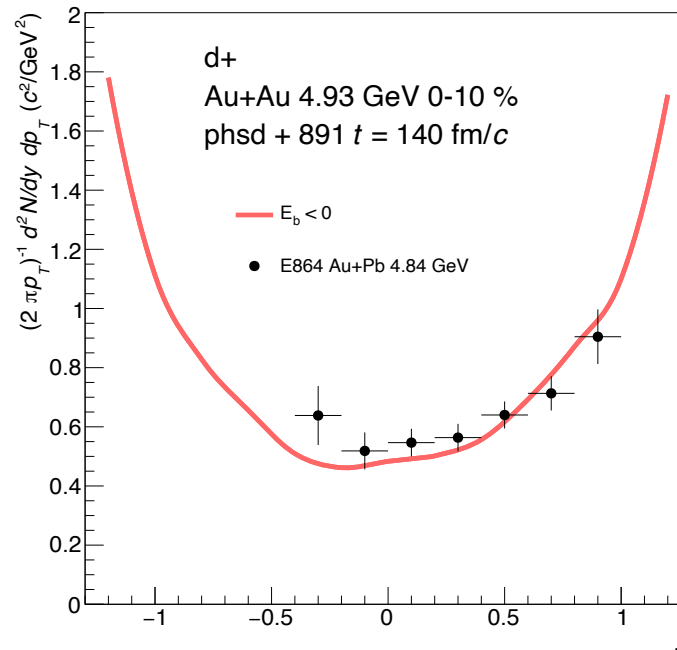
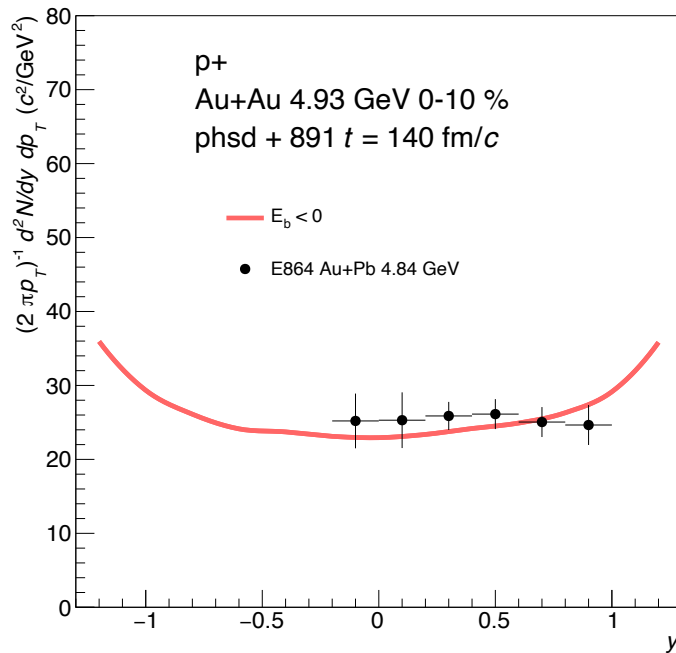
```
Int_t      fEventNr;      // Event number
Double_t   fB;            // Impact parameter (fm)
Double_t   fPhi;          // Reaction plane angle
Int_t      fNes;          // Number of event steps
Int_t      fStepNr;       // Event step number
Double_t   fStepT;        // Event step time
Int_t      fNpa;          // Number of particles
TString    fComment;      // Generator-specific information
TClonesArray* fParticles; // Array of particles
```

*available in actual version, other information can be added

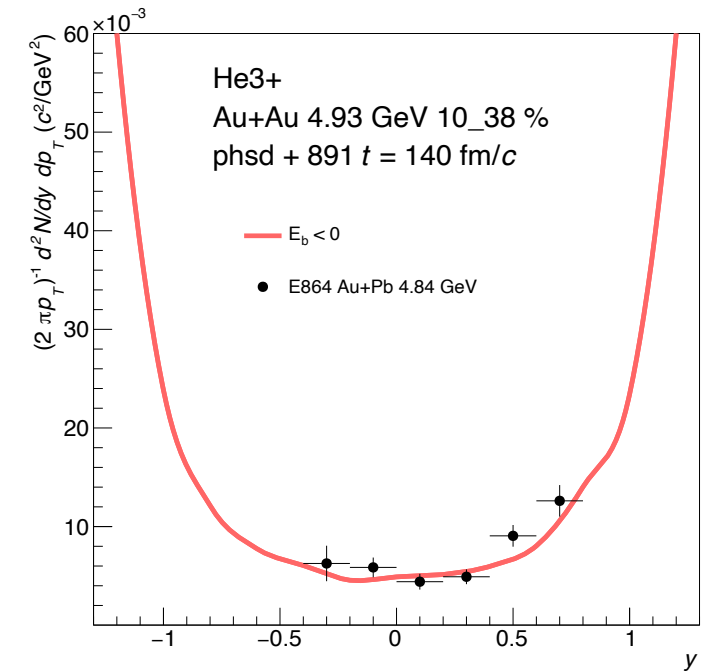
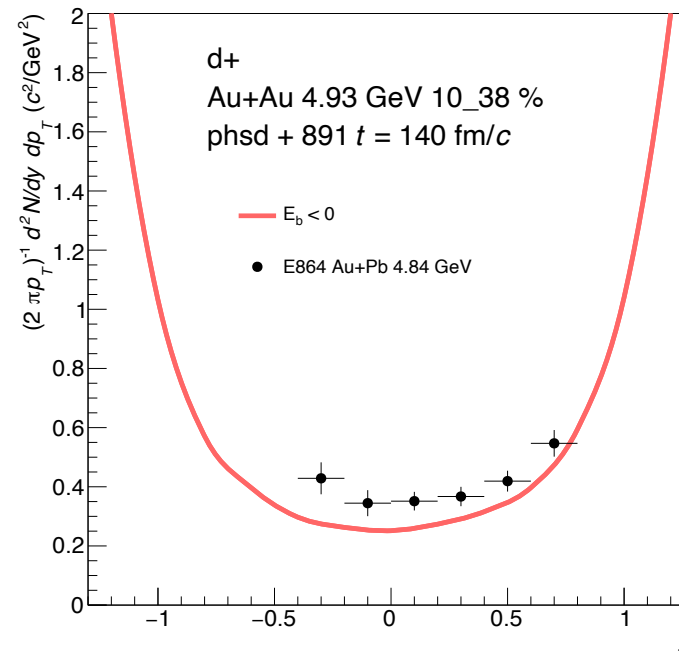
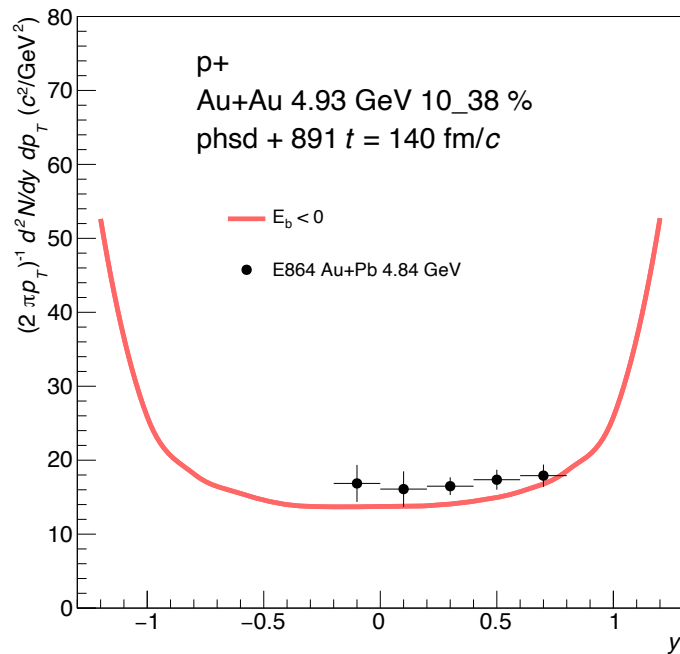
UParticle

```
Int_t      fIndex;        // index of this particle
Int_t      fPdg;          // PDG code
Int_t      fStatus;       // Status
Int_t      fParent;       // Index of parent
Int_t      fParentDecay;   // Parent decay index
Int_t      fMate;         // index of last collision partner
Int_t      fDecay;        // decay index (-1 if not decayed)
Int_t      fChild[2];     // index of first and last child
Double32_t fPx;           // px (GeV)
Double32_t fPy;           // py (GeV)
Double32_t fPz;           // pz (GeV)
Double32_t fE;            // Energy (GeV)
Double32_t fX;            // x (fm)
Double32_t fY;            // y (fm)
Double32_t fZ;            // z (fm)
Double32_t fT;            // t (fm)
Double32_t fWeight;       // weight
```

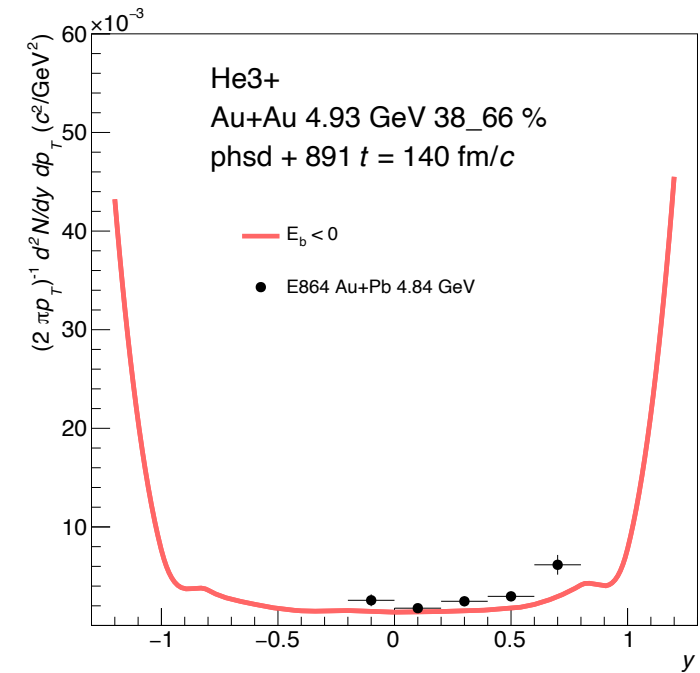
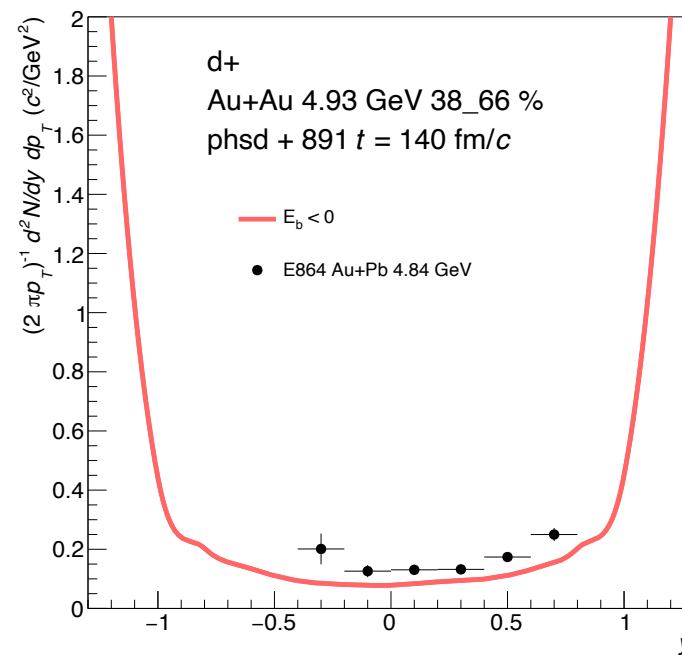
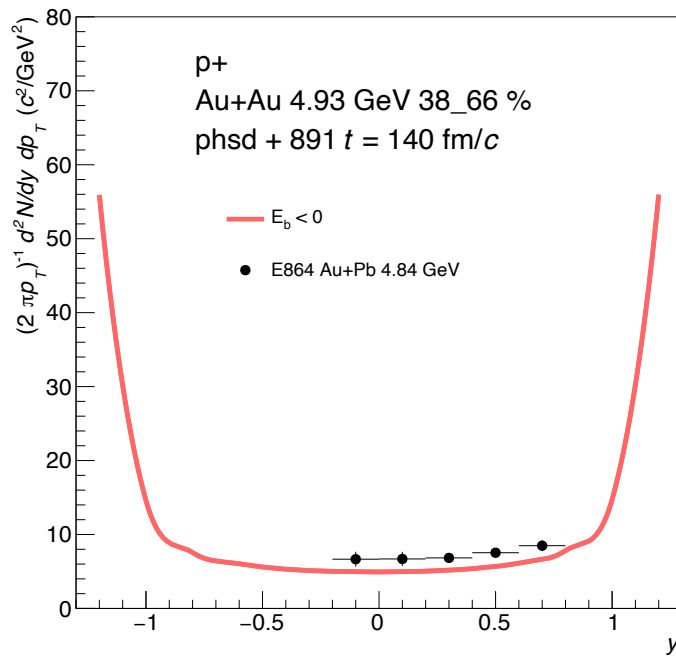
PHQMD Au-Au 4.93 GeV 0-10 % most central compared to experimental data



PHQMD Au-Au 4.93 GeV 10-38 % most central compared to experimental data



PHQMD Au-Au 4.93 GeV 38-66 % most central compared to experimental data



Run PHQMD, stabilisation & conversion at the batchfarm

All 3 steps* can be performed with:

. runPHQMD.sh

Only *runinfo.sh* needs to be modified.

The following files are needed (besides the PHQMD code):

- | | |
|-----------------------------------|--|
| - runinfo.sh | modification of specifications for run |
| - runPHQMD.sh | starts run on batch farm |
| - batch_runPHQMD.sh | executed on batch farm |
| - 791to891.exe | stabilisation routine |
| - convert_phqmd_detector_unigen.C | conversion PHQMD-output files into one root-file |
| - cluster_table.dat | contains information for definition of nuclei |

runinfo.sh

Only runinfo.sh needs to be modified:

- selection of steps (run PHQMD code, stabilisation, conversion)
- choose option for cluster with $A > 7$ (all or only physical clusters are counted)
- choose if only clusters OR clusters & anti-clusters are converted
- select input information for PHQMD, e.g system, energy, number of parallel events, impactparameter etc.
- location information

runinfo.sh – run options, locations

```
## Options for running PHQMD code, stabilisation & conversion #
```

```
## Steps:
```

```
export RunPhqmdCode=1      ## = 1: run PHQMD code – input: inputPHSD, output: phsd.dat (hadrons), fort.791 (cluster-baryons), fort.891 (anti-  
cluster-baryons)  
export RunStabilisation=1  ## = 1: run stabilisation routine – input: fort.791 & fort.781, output: fort.891 & fort.881  
export MakeDetectorInput=1 ## = 1: run conversion of PHQMD output into detector input – input: fort.891 & fort.881, output: unigen format
```

```
## Options for steps:
```

```
export ConvertAntiClusters=1 ## = 0: convert/stablise only clusters, = 1: convert clusters and anti-clusters  
export CountAllClusters=0    ## = 0: only physical clusters are counted (unphysical clusters are counted as single baronys),  
                             ## = 1: all clusters A > 7 are counted as clusters
```

```
## Delete & Merge
```

```
export DeleteFiles=0        ## = 1: deletes all output files, but the detector-input [dataset].phqmd.root  
export RunHadd=0            ## = 1: add 100 jobs into one file  
export RunHaddSim=0         ## = 1: add 10 jobs in one file for detector simulations
```

```
##### Directories and files #####
```

```
export LOCATION=/lustre  
export DIR=$LOCATION/cbm/users/$USER  
export PHQMDDIR=$DIR/phqmd  
export CBMROOT_DIR=$DIR/cbmsoft/cbmroot/build  
export OUTDIR=$DIR/mc/phqmd/$SYSTEM_"_$ENERGY"GeV_"$IMPACTPARAMETER_MAX"fm_Num"$NUM"xSub"$ISUBS
```

runinfo.sh – input parameter, sbatch parameter

```
## Parameters for sbatch #####
```

```
export firstJob=1
export lastJob=2
export array=$firstJob-$lastJob
export last_hadd=$(printf "%0.0f" "$(echo " $lastJob / 100 " | bc -l)")
export array_hadd=1-$last_hadd
export last_hadd_simcbm=$(printf "%0.0f" "$(echo " $lastJob / 10 " | bc -l)")
export array_hadd_simcbm=$firstJob-$last_hadd_simcbm
export time="1-23:59:59"
export partition="long"
```

```
## Parameters for inputPHSD (more parameters available, but should usually not be changed)
```

```
export SYSTEM=auau          ## specification below for auau, pbpb, aupb, aupt – other systems need to be added below
export ENERGY=4.93         ## center of mass energy – !for sqrt(s) > 7 GeV setting for kinetic deuterons in PHQMD code should be changed!
export ELAB=$(echo " $ENERGY * $ENERGY / (2 * 0.938) - (2 * 0.938)" | bc -l) ## Lab energy per nucleon (needed for PHQMD)
export NUM=100              ## number of parallel events (NUM=100 for 3 GeV & 4.9 GeV required)
export ISUBS=1              ## number of subsequent runs
export nEvents=$(echo " $NUM * $ISUBS " | bc -l)

export IMPACTPARAMETER_MIN=0.0 ## minimal impact parameter in fm
export IMPACTPARAMETER_MAX=5.0 ## maximum impact parameter in fm ! Max impact parameter must not be > 15.0 fm !

export IGLUE=1              ## =1 with partonic QGP phase (PHSD mode); =0 – HSD mode ! Needs to be set to = 0 for 3 GeV
export EOS=0                ## EoS; =0: hard EOS without M.D.I; =1 soft EoS
export ICLUSTER=1          ## enable or disable CLUSTER output
```

PHQMD settings- important remarks

Energy:	needs to be given in units of lab energy per nucleon
Impact parameter:	maximum must be set $b < 15.0$ fm
NUM:	number of parallel events (needed for mean-field in PHSD) must be: NUM = 100 for 3.0 & 4.9 GeV, NUM = 200 for lower energies
IGLUE	: =1 with partonic QGP phase (PHSD mode); =0 - HSD mode must be set to = 0 for 3 GeV

Please be aware: It is possible that jobs do not finish due to time limit.

For higher energies $\sqrt{s} > 7$ GeV: Setting for kinetic deuterons needs to be changed in the PHMQD code.

Options

- * Store PHQMD output files into root-format *EventPHQMD*

<https://github.com/SusanneGlaessel/EventPHQMD>

- * PHQMD with freeze-out time, position & momentum (available for hadrons):

change in PHQMD code & recompile: IFreezeOut=1 (in ini_param.f)

calculation of freeze-out time, position & momentum for clusters:

- freeze-out time of cluster: freeze-out-time of cluster-baryon with latest freeze-out-time
- freeze-out position & momentum: transport all cluster-baryons to cluster freeze-out-time

!!! with QMD (attractive potential) freeze-out-momentum $\neq$ final momentum

→ Format: *EventFreeze* to save freeze-out-momentum!!!

- * Write output from phsd.dat only (single hadrons)