

# «Nuclei» wagon

Update

# Outline

- Introduction
  - How it was
  - Some new ideas
- Implementation
- Documentation and code organisation
- Utilisation
- Summary

# Introduction

# How it was

*Do first, think later*

- JSON configuration file for the wagon 
- Documented C++ «wagon» code 
- Single Python post processing program:
  - Many subroutines in one place 
  - Fitting procedure parallelisation: hardcoded ranges, initial parameters  
  - Wagon configuration file used for the analysis  

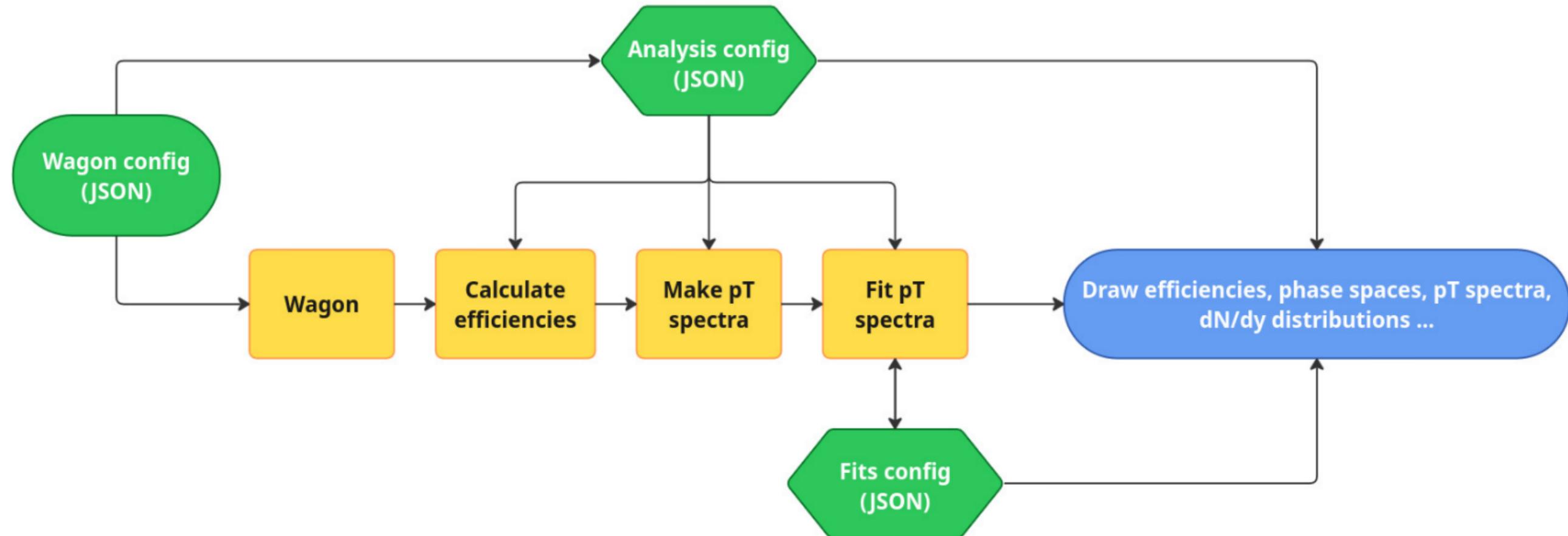
# Some new ideas

*Think first, do later*

- Config driven analysis — move as much as possible out of the code
- Rethink the architecture, but keep old good ideas
- Old code refactoring
- New approach for the code storage, integration and deployment
- Documentation: out of the code, extended

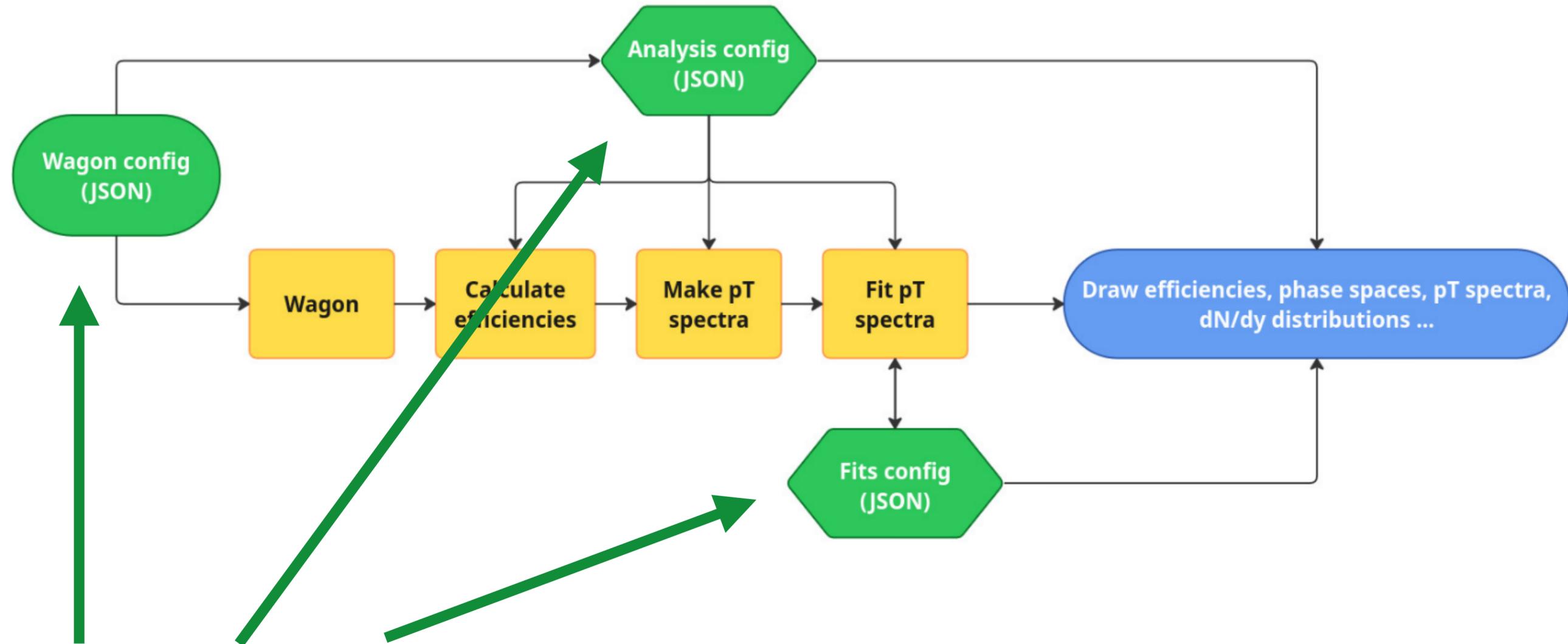
# Some new ideas

*How it should be*



# Some new ideas

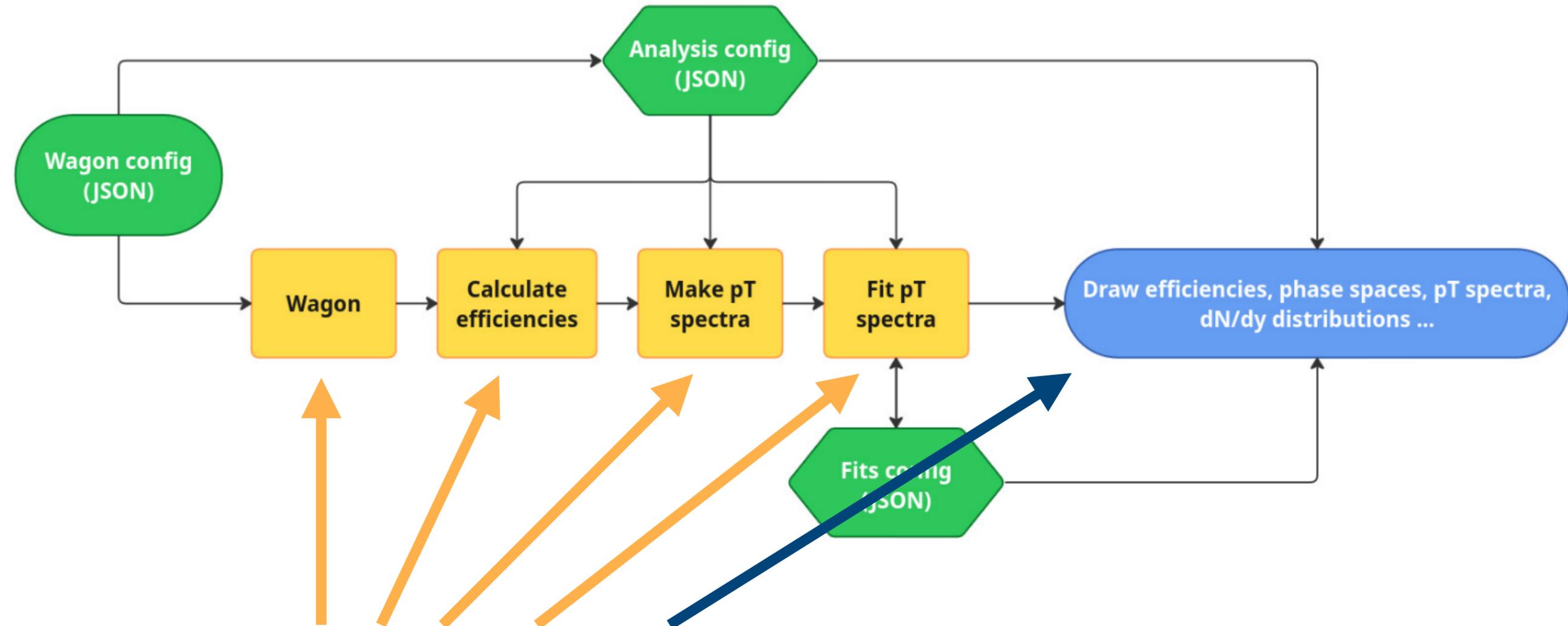
*How it should be*



**These parts are out of the code, configurable and could/must be modified**

# Some new ideas

*How it should be*



These parts are the code and must **NEVER** be modified

# Typical errors in MpdRoot wagons

*which could be easily solved by the configuration files*

- ✘ Hardcoded values such as particles PDG codes or PID IDs
- ✘ Hardcoded constants such as particle masses or even Pi
- ✘ Hardcoded centrality classes
- ✘ Hardcoded histograms and their bins, limits etc
- ✘ Hardcoded parameters and «magic numbers» without explanation

# Typical errors in MpdRoot wagons

*which could not be solved by the configuration files  
but must be solved somehow*

- ✘ Missing comments: purpose of the functions? Parameters? Return values?
- ✘ Lack of the documentation: «black box» type analysis — how to run it? How to use it?
- ✘ Obsolete commented-out code. Why it is here? Is it needed?
- ✘ Code duplication
- ✘ No post processing code: how to make distributions? How to get (pre-) final plots?

# Implementation

trying again to avoid common mistakes

# Implementation

*of the new ideas*

- JSON configuration file for the wagon



Left unchanged — it was a brilliant idea

```
{
  "Verbose": "1",
  "N_MPD_PID_Particles": "8",
  "make_MC": "1",
  "make_Efficiency": "1",
  "PID_mode": "2",
  "DCA_mode": "1",
  "TOF_mode": "1",
  "use_pt_corrections": "0",
  "pt_corrections_file": "pt_corrections.root",
  "Events": {
    "PrimaryVertexZ": "100",
    "Centrality": [[0, 20], [20, 40], [40, 80]]
  },
  "Tracks": {
    "NHits": "27",
    "NSigmaDCAx": "1",
    "NSigmaDCAy": "1",
    "NSigmaDCAz": "1",
    "LowPtCut": "0.05"
  },
  "PID": {
    "TPCSigma": "2",
    "TOFSigma": "2",
    "TOFDphiSigma": "3",
    "TOFDzSigma": "3"
  },
  "MpdPid": {
    "Energy": "9.2",
    "Coef": "1.0",
    "Generator": "PHQMD",
    "Tracking": "CFHM",
    "IniString": "pikaprdetrhe3he4",
  }
}
```

# Implementation

*of the new ideas*

- Documented C++ «wagon» code



Mostly unchanged — amazingly good architecture from the beginning

**MpdNuclei**

Main Page | Classes | Files

MpdNuclei

- Classes
- Class List
- MpdNuclei
- Class Index
- Class Hierarchy
- Class Members
- Files

### MpdNuclei Class Reference

#include <MpdNuclei.h>

Inheritance diagram for MpdNuclei:

```
graph TD; MpdNuclei --> MpdAnalysisTask; MpdNuclei --> MpdPid;
```

**Classes**

- struct centrality\_bin
- struct particle\_info

**Public Member Functions**

- MpdNuclei ()**  
Default constructor.
- MpdNuclei (const char \*name, const char \*outputName="taskName", const char \*settings\_filename="file.json")**
- ~MpdNuclei ()**  
Default destructor.
- void UserInit ()**
- void ProcessEvent (MpdAnalysisEvent &event)**
- void Finish ()**

**Private Types**

# Implementation

*of the new ideas*

## Changes in the wagon code

```
std::vector<std::vector<TH2F*>> hv__eff_tpc_numerator;  
std::vector<std::vector<TH2F*>> hv__eff_tpc_denominator;  
std::vector<std::vector<TH2F*>> hv__cont_sec_numerator;  
std::vector<std::vector<TH2F*>> hv__cont_sec_denominator;  
std::vector<std::vector<TH2F*>> hv__eff_pid_numerator;  
std::vector<std::vector<TH2F*>> hv__eff_pid_denominator;  
std::vector<std::vector<TH2F*>> hv__cont_pid_numerator;  
std::vector<std::vector<TH2F*>> hv__cont_pid_denominator;  
std::vector<std::vector<TH2F*>> hv__eff_tof_denominator;  
std::vector<std::vector<TH2F*>> hv__eff_pid_dedx_numerator;  
std::vector<std::vector<TH2F*>> hv__cont_pid_dedx_numerator;  
std::vector<std::vector<TH2F*>> hv__cont_pid_dedx_denominator;  
std::vector<std::vector<TH2F*>> hv__eff_pid_tpc_numerator;  
std::vector<std::vector<TH2F*>> hv__cont_pid_tpc_numerator;  
std::vector<std::vector<TH2F*>> hv__cont_pid_tpc_denominator;  
std::vector<std::vector<TH2F*>> hv__eff_pid_tof_numerator;  
std::vector<std::vector<TH2F*>> hv__cont_pid_tof_numerator;  
std::vector<std::vector<TH2F*>> hv__cont_pid_tof_denominator;
```



```
/// List of efficiencies counters  
enum EffCounters {  
    kMC_MCPID,          ///< 0: MC tracks, MC PID, mc y and pt  
    kRC_MCPID,          ///< 1: RC tracks, MC PID, mc y and pt  
    kRC_MCPID_R,       ///< 2: RC tracks, MC PID, reconstructed y and pt  
    kRC_MCPID_PRIM,    ///< 3: RC tracks, MC PID, Primary by MC, mc y and pt  
    kRC_MCPID_SEC,     ///< 4: RC tracks, MC PID, Secondary by MC, reconstructed y and pt  
    kRC_MCPID_TOF,     ///< 5: RC tracks, MC PID, TOF, reconstructed y and pt  
    kRC_RCPID_DEDX,    ///< 6: RC tracks, RC PID dEdx only, reconstructed y and pt  
    kRC_RCPID_DEDX_T,  ///< 7: RC tracks, RC PID dEdx only == PDG, reconstructed y and pt  
    kRC_RCPID_DEDX_F,  ///< 8: RC tracks, RC PID dEdx only != PDG, reconstructed y and pt  
    kRC_RCPID_COMB,    ///< 9: RC tracks, RC PID combined (has ToF), reconstructed y and pt  
    kRC_RCPID_COMB_T,  ///< 10: RC tracks, RC PID combined == PDG, reconstructed y and pt  
    kRC_RCPID_COMB_F,  ///< 11: RC tracks, RC PID combined != PDG, reconstructed y and pt  
    kRC_EVPID_TPC,     ///< 12: RC tracks, evPID TPC, reconstructed y and pt  
    kRC_EVPID_TPC_T,   ///< 13: RC tracks, evPID TPC == PDG, reconstructed y and pt  
    kRC_EVPID_TPC_F,   ///< 14: RC tracks, evPID TPC != PDG, reconstructed y and pt  
    kRC_EVPID_TOF,     ///< 15: RC tracks, evPID TOF, reconstructed y and pt  
    kRC_EVPID_TOF_T,   ///< 16: RC tracks, evPID TOF == PDG, reconstructed y and pt  
    kRC_EVPID_TOF_F,   ///< 17: RC tracks, evPID TOF != PDG, reconstructed y and pt  
    kNumCounters       ///< 18: Total number of counters  
};  
/** @name Efficiency histograms  
*/  
///@{  
    std::vector<std::vector<std::vector<TH2F*>>> hv__eff_counter =  
        std::vector<std::vector<std::vector<TH2F*>>>(kNumCounters); ///< Efficiencies histograms (2D pT-y phase-space).  
///@}
```

- New approach to the efficiencies counters: enum list, single object → easy to understand and to operate ✓
- New counters can be added to the enum list → automatic creation and initialisation of histograms ✓

# Implementation

*of the new ideas*

- Single Python post processing program



- Several simpler single-purpose Python programs



Analysis configuration file — the main part of the analysis

# Implementation

*New analysis configuration file*

Config driven analysis 

- System string, e.g. «Xe + W @ 2.5 A.GeV».
- Efficiencies (numerators, denominators — any combination of the wagon «efficiencies counters»)
- Paths to the wagon configuration file, each program output files, files with fits initial parameters.
- Analysis rapidity intervals, centrality bins, particle species to include.
- The pT ranges (for each particle in each rapidity interval in each centrality bin) to get reconstructed dN/dy point (and thus the ranges for the Blast-Wave extrapolations).
- More to come.

# Implementation

*New analysis configuration file*

Config driven analysis 

Although **everything** can be set in **the single configuration file for the whole analysis chain** the priority in all programs is given to the command line arguments.

**Flexibility is also a key.**

# Implementation

*Efficiencies calculation program* 

- This program calculates the efficiencies used for the spectra corrections.
- **Output:** the ROOT-file with the 2D phase-space histograms and rapidity slices.
- Rapidity slices and efficiencies are defined in the analysis configuration JSON file.
- Works automatically for the defined particles, centralities, rapidity intervals.

# Implementation

## *Efficiencies calculation program*

### Efficiencies defined in the program

```
# Efficiencies definition: numerator, denominator, name
efficiencies_types_example = [
  # TPC efficiency:
  # Numerator: MC_PID, RC tracks, Primary by MC.
  # Denominator: MC_PID, MC tracks.
  {'numerator': 'h_eff_counter_3', 'denominator': 'h_eff_counter_0', 'output': 'efficiency_tpc'},
  # Secondaries contamination:
  # Numerator: MC_PID, Secondary by MC, RC tracks.
  # Denominator: MC_PID, RC tracks.
  {'numerator': 'h_eff_counter_4', 'denominator': 'h_eff_counter_1', 'output': 'contamination_secondaries'},
  # ToF efficiency:
  # Numerator: MC_PID, has ToF, RC tracks.
  # Denominator: MC_PID, RC tracks.
  {'numerator': 'h_eff_counter_5', 'denominator': 'h_eff_counter_2', 'output': 'efficiency_tof'},
  # MpdPid 'Combined' PID efficiency:
  # Numerator: RC_PID, has ToF, RC tracks.
  # Denominator: MC_PID, has ToF, RC tracks.
  {'numerator': 'h_eff_counter_9', 'denominator': 'h_eff_counter_5', 'output': 'efficiency_pid'},
  # MpdPid 'Combined' PID purity:
  # Numerator: RC_PID == PDG, has ToF, RC tracks.
  # Denominator: RC_PID, has ToF, RC tracks.
  {'numerator': 'h_eff_counter_10', 'denominator': 'h_eff_counter_9', 'output': 'purity_pid'},
  # MpdPid 'Combined' PID contamination:
  # Numerator: RC_PID != PDG, has ToF, RC tracks.
  # Denominator: RC_PID, has ToF, RC tracks.
  {'numerator': 'h_eff_counter_11', 'denominator': 'h_eff_counter_9', 'output': 'contamination_pid'},
  # MpdPid dE/dx only PID efficiency:
  # Numerator: RC_PID, RC tracks.
  # Denominator: MC_PID, RC tracks.
  {'numerator': 'h_eff_counter_6', 'denominator': 'h_eff_counter_2', 'output': 'efficiency_pid_dedx'},
  # MpdPid dE/dx only PID purity:
  # Numerator: RC_PID == PDG, RC tracks.
  # Denominator: RC_PID, RC tracks.
  {'numerator': 'h_eff_counter_7', 'denominator': 'h_eff_counter_6', 'output': 'purity_pid_dedx'},
  # MpdPid dE/dx only PID contamination:
  # Numerator: RC_PID != PDG, RC tracks.
  # Denominator: RC_PID, RC tracks.
  {'numerator': 'h_eff_counter_8', 'denominator': 'h_eff_counter_6', 'output': 'contamination_pid_dedx'}
]
```

### Efficiencies defined in the configuration file

```
..
"Efficiencies": [
  {"numerator": "h_eff_counter_3", "denominator": "h_eff_counter_0", "output": "efficiency_tpc"},
  {"numerator": "h_eff_counter_4", "denominator": "h_eff_counter_1", "output": "contamination_secondaries"},
  {"numerator": "h_eff_counter_5", "denominator": "h_eff_counter_2", "output": "efficiency_tof"},
  {"numerator": "h_eff_counter_9", "denominator": "h_eff_counter_5", "output": "efficiency_pid"},
  {"numerator": "h_eff_counter_10", "denominator": "h_eff_counter_9", "output": "purity_pid"},
  {"numerator": "h_eff_counter_11", "denominator": "h_eff_counter_9", "output": "contamination_pid"},
  {"numerator": "h_eff_counter_6", "denominator": "h_eff_counter_2", "output": "efficiency_pid_dedx"},
  {"numerator": "h_eff_counter_7", "denominator": "h_eff_counter_6", "output": "purity_pid_dedx"},
  {"numerator": "h_eff_counter_8", "denominator": "h_eff_counter_6", "output": "contamination_pid_dedx"}
],
```

If efficiencies are not defined in the configuration file those defined in the efficiencies calculation program will be used!

# Implementation

*Transverse momentum spectra calculation program* 

- This program calculates the transverse momentum spectra and applies the efficiency and contamination corrections calculated on the previous step.
- **Output:** ROOT-file with the transverse momentum spectra sliced by the defined rapidity bins.
- Three types of the spectra are saved: MC, reconstructed and reconstructed after applying the efficiency and contamination corrections.
- Rapidity slices and efficiencies are defined in the analysis configuration JSON file.
- Works automatically for the defined particles and centralities.

# Implementation

*Transverse momentum spectra fitting program* 

- This program fits the transverse momentum spectra calculated with previous program.
- Currently there are only two fit functions:

- Thermal: 
$$\frac{d^2N}{dp_T dy} = \frac{dN/dy}{T(m_0 + T)} p_T \exp\left(-\frac{m_t - m_0}{T}\right)$$

- Blast-Wave: 
$$\frac{dN}{p_T dp_T} = C \int_0^{R_{max}} r dr m_T I_0\left(\frac{p_T \sinh \rho(r)}{T}\right) K_1\left(\frac{m_t \cosh \rho(r)}{T}\right)$$

- Initial parameters for the each fit are defined in the JSON files.

# Implementation

*Transverse momentum spectra fitting program* 

Thermal fit initial parameters file has the structure of the sublist of the fit parameters for the each:

- particle
  - centrality bin
  - rapidity region



```
therma_init.json 12.36 KIB
1 {
2   "pim": {
3     "0": [
4       [96.75, 0.1003, 0.139, 0.2, 1.5],
5       [99.27, 0.0977, 0.139, 0.2, 1.5],
6       [89.42, 0.1014, 0.139, 0.2, 1.5],
7       [85.47, 0.1030, 0.139, 0.2, 1.5],
8       [79.28, 0.0972, 0.139, 0.2, 1.5],
9       [74.14, 0.1021, 0.139, 0.2, 1.5],
10      [70.35, 0.1012, 0.139, 0.2, 1.5],
11      [65.70, 0.1008, 0.139, 0.2, 1.5],
12      [60.17, 0.0972, 0.139, 0.2, 1.5],
13      [53.84, 0.1030, 0.139, 0.2, 1.5]
14    ],
15  },
16 }
```

Parameters:  $dN/dy$ ,  $T$ ,  $m_0$  (fixed!),  $pt_{low}$ ,  $pt_{high}$ , where  $[pt_{low}, pt_{high}]$  is the fit range.

# Implementation

*Transverse momentum spectra fitting program* 

Blast-Wavefit initial parameters file has the structure same for the reconstructed and the Monte-Carlo data (also for the each particle, centrality bin, rapidity interval):

- `C`: Normalization constant. `m0`: Particle rest mass.
- `T`: Kinetic freeze-out temperature. `beta`: Average transverse flow velocity.
- `[pt\_low, pt\_high]`: Fit range.
- `R\_max`, `steps`: Source radius and the number of integration steps.
- `fit\_found`: Flag which indicates that fit was found (1) or not (0).

```
[  
 3333.33,  
 3.728,  
 0.15,  
 0.4,  
 1.0,  
 4.0,  
 20.0,  
 40,  
 1  
],
```

# Implementation

*Transverse momentum spectra fitting program* 

The Blast-Wave (BW) fitting procedure is quite complex and the fit quality is not very often «good».

- The fit is performed with the initial parameters.
- The values of the fit at the low and high transverse momentum bins are compared with the fitted spectra.
- If the difference in both regions is small — the flag `fit\_found` is set to 1, the initial parameters are considered as final.
- If the difference in any region is high — the the procedure randomly set the new parameters until "good" parameters are found or until 100 tries. If "good" parameters are found — they overwrite the initial parameters in the fit parameters file, the `fit\_found` is set to 1. Otherwise the `fit\_found` flag is still 0.
- On the next runs the program will use the fit parameters with the flag `fit\_found = 1` and skips the fitting procedure.

# Implementation

*Transverse momentum spectra fitting program* 

In this case the **config-driven approach** allows you to target the specific particle, centrality, rapidity interval without affecting other particles and/or regions and thus losing time to re-fit everything again and again.

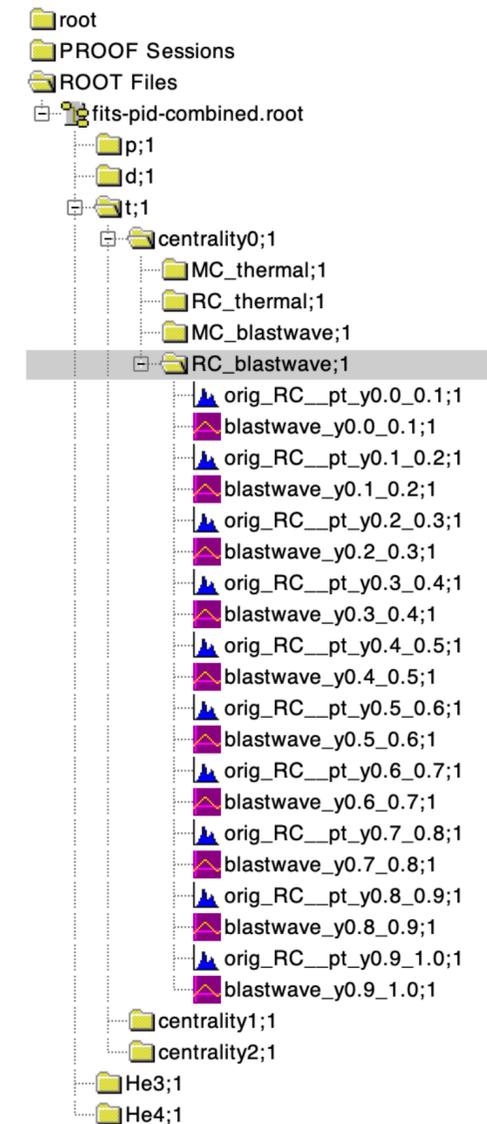
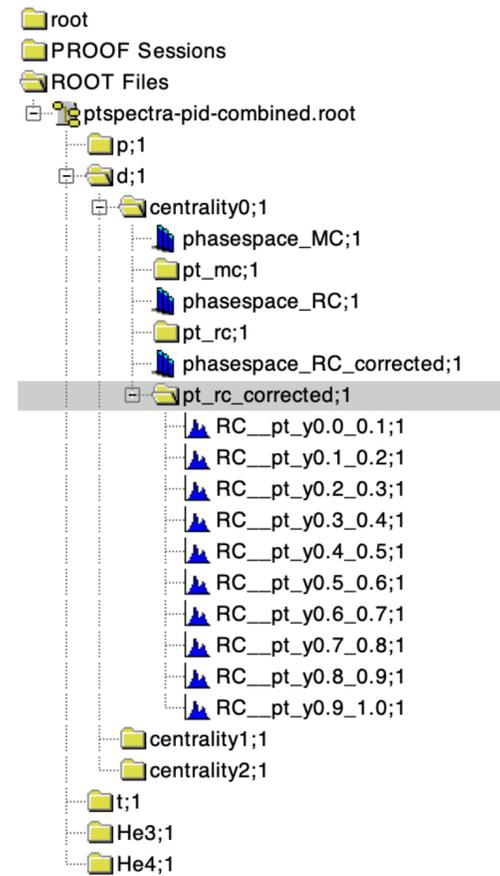
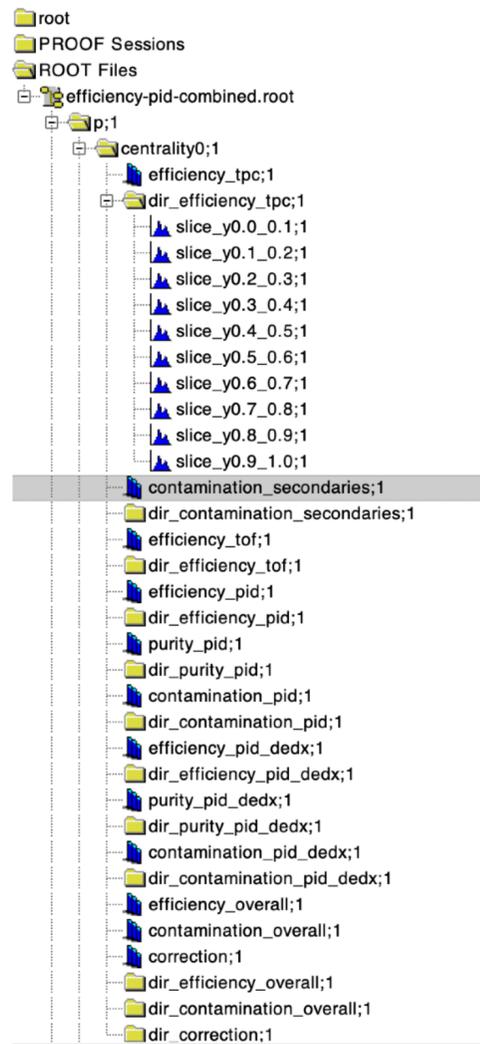
# Implementation

*Separate drawing programs* 

- Efficiency and contamination phase spaces and their slices in the defined rapidity bins.
- Particles phase spaces: Monte-Carlo, reconstructed and reconstructed corrected by the efficiencies and contaminations.
- Transverse momentum spectra: Monte-Carlo and reconstructed (corrected) with the Blast-Wave fits and ratios between the original Monte-Carlo spectra and the reconstructed data + Blast-Wave fits interpolation to the low and high transverse momentum regions.
- Particles  $dN/dy$  spectra (Monte-Carlo and reconstructed):
  - The integral of the corrected spectra within defined transverse momentum  $p_T$  bins.
  - The integral of the interpolation of the Blast-Wave fits to the low and high  $p_T$  regions.

# Output files

- Clarity, logical structure, self-describing names



# Summary for the first part

- Some old approach ideas works really great, however some modifications were be made.
- The new config-driven analysis approach sounds complex, but in reality is much easier to implement and then to perform than the standard «put everything in the program» method.
- The performance gain becomes even more pronounced when re-use analysis with the tuned fit parameters.

# Documentation

and code organisation

# Documentation

- Wagon C++ code is documented using the Doxygen engine and tons of comments.
- Analysis and drawing routines are documented with Doxygen-style comments instead of the Python docstring for the sake of unification. Tons of ordinary comments are still included.
- The main wagon C++ code and analysis routines are also supplemented by the documentation in Markdown (README.md files in the repository).
- New documentation dedicated repository organised.

# Documentation

New documentation repository: overall view

 README.md

---

## Some documentation for/from the PWG-2 group

---

### Getting started

---

- [First steps](#)

### Wagons

---

- [Nuclei \(documentation\)](#)
  - [Source code](#)

### Analysis

---

- [Nuclei](#)

**Created on**  
April 22, 2025

# Documentation

New documentation repository: first steps — newcomers are welcome

## 0. Install MpdRoot

---

Instructions here: <https://mpdroot.jinr.ru/running-mpdroot-on-the-cluster-nica-lhep-hybrilit/>

## 0.5. Load the helper environment

---

Load environment for the helper macros:

```
$ . /nica/mpd1/kireev/public/00_env.sh
```

### Important

**Please, note, this environment and helper macros are not official.**

You can look into their source code for some comments/description to get an idea how to run programs for the each described step.

**Be very careful while using these macros** and even maybe contact me in advance.

## 1. Run the Monte-Carlo simulations

---

You can run [PHQMD](#) and/or [UrQMD](#) models on the [NCX](#) batch farm:

**phqmd2batch** : run PHQMD on the cluster

- `-h, --help` show this help message and exit
- `--massta` Target mass (default: 208)

# Documentation

New documentation repository: dive deeper into the analysis wagons structure

Please, do not hesitate to contact me if you want to contribute and describe your analysis.

More documentation — better:

- It simplifies onboarding of the newcomers (and students).
- It give insights on what we are doing (pure code does not allow this) — exchange of methods and ideas, cross-checks etc.

## MpdNuclei wagon

---

MPDRoot wagon for the light nuclei analysis. Also applicable for the hadrons: pions, kaons, protons.

- [MpdNuclei wagon](#)
  - [Introduction](#)
  - [Dependencies](#)
    - [Branches](#)
    - [Wagons](#)
  - [Usage](#)
  - [Configuration](#)
    - [Global settings](#)
    - [Event settings](#)
    - [Track settings](#)
    - [PID settings](#)
    - [MpdPid settings](#)
    - [evPID settings](#)
    - [Particles settings](#)
    - [Example](#)
  - [Documentation](#)

## Introduction

---

The main idea of the 'Nuclei' wagon is to collect 2D phase-space hisograms ( $p_T, y$ ) for the defined particle species. Then the two-dimensional histogram can be 'sliced' by the  $y$  bins to get the transverse momentum  $p_T$  spectra in these slices. By integrating the spectra one can get the total number of particles in that rapidity bin (slice) and at the end -- rapidity density distribution ( $dN/dy$ ) by combining the numbers obtained in slices.

# Documentation

New documentation repository: analysis in details

Sometimes it is inconvenient to put everything is small details in the code — we can still describe what and how we are doing.

Please, do not hesitate to contact me if you want to contribute and describe your analysis.

More documentation — better:

- It simplifies onboarding of the newcomers (and students).
- It give insights on what we are doing (pure code does not allow this) — exchange of methods and ideas, cross-checks etc.

## Efficiencies in the 'Nuclei' wagon

### Common "event quality" conditions

These cuts are applied to the each event:

- Event has a primary vertex.
- Primary vertex Z-position is within some window (e.g.  $|V_Z| < 100$  cm).
- Event has a defined centrality by the TPC particles multiplicity.

### Common "track quality" cuts

For the MPD collider mode (used for Bi+Bi @ 9.2 GeV):

- $DCA < 3$  cm
- $N_{hits} > 20$
- The split track is rejected

For the MPD fixed-target mode (FXT, used for Xe+W @ 2.5 A.GeV):

- $DCA_{x,y,z} < 1\sigma$
- $N_{hits} > 24$
- The split track is rejected

### TPC efficiency

Numerator: reconstructed (actually, MC tracks, associated to the reconstructed) primary tracks which satisfy the reconstruction 'Track quality' condition. Denominator: all simulated Monte-Carlo tracks.

$$\epsilon_{TPC} = \frac{[\text{Primary(MC) = TRUE}] [\text{Identified(MC)}] [\text{Track quality}]}{[\text{Primary(MC) = TRUE}] [\text{Identified(MC)}]} \quad (1)$$

## Postprocessing for the "Nuclei" wagon

- Postprocessing for the "Nuclei" wagon
  - Efficiencies in the 'Nuclei' wagon
    - Common "event quality" conditions
    - Common "track quality" cuts
    - TPC efficiency
    - Secondaries contamination
    - ToF matching efficiency
    - PID efficiency (MpdPid class case)
    - Final corrections
  - Analysis configuration file
  - Analysis scripts
    - 0. nw\_common.py
    - 1. nw\_ana\_efficiencies.py
      - Usage
      - Options
    - 2. nw\_ana\_ptspectra.py
      - Usage
      - Options
    - 3. nw\_ana\_fits.py
      - Usage
      - Options
      - Fits
        - Thermal fit initial parameters file:
        - Blast-Wave fit initial parameters file:
    - 1. Thermal fit

# Code organisation

- The wagon and analysis development moved out of the MpdRoot tree:  
now it is easier to implement and check new ideas, refactor code, test the workflow etc.
- The new stand-alone repository is still publicly available.
- The 'nuclei' analysis directory is still in the MpdRoot tree:  
when a new «stable» version of wagon code or analysis routines become available it will be immediately synchronised with the MpdRoot 'dev' branch.
- Such procedure helps to maintain the development of the analysis active while reducing the commits and merge requests load on the MpdRoot repository: **win-win** approach.

# Summary for the second part

- Please, write **documentation**.
- Do not hesitate to put explanatory comments in the code:
  - Describe what, how and why you are doing.
  - Describe input parameters, returned value.
- This sounds annoying, waste of time and unnecessary, but in reality **is extremely important**.

# Utilisation

actually, this is the results section

# Utilisation

*End of the day*

## Request 37:

5M of PHQMD Xe+W min. bias events at  $T = 2.5$  GeV/n

## MPD

Sign Up Log In



### Request 37: General-purpose (hadrons, light nuclei), 5M PHQMD Xe+W ( $T = 2.5$ GeV/n, FXT)

Monte-Carlo productions



vkireyeu

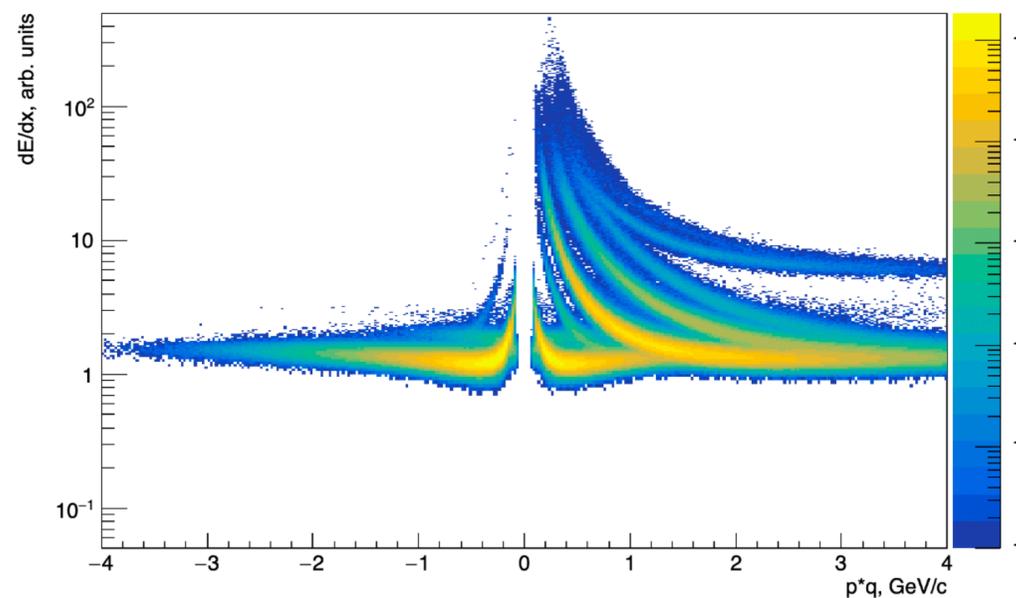
1 Apr 4

Apr 4

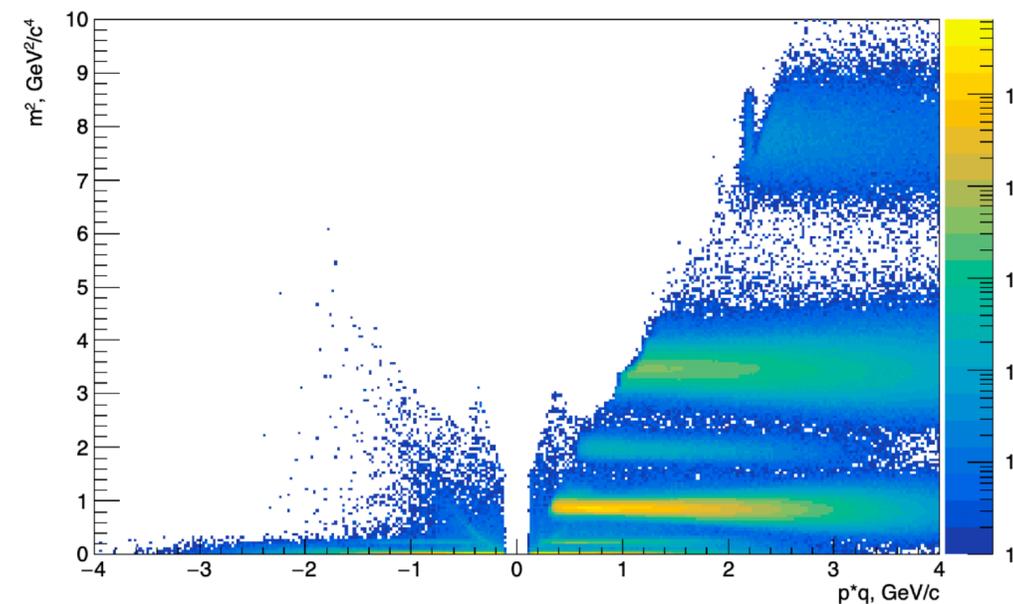
1 / 2  
Apr 4

This is a PWG2 general-purpose (hadrons and light nuclei spectra) simulation for min.bias Xe124 + W184 collisions.  
Beam kinetic energy: 2.5 A.GeV.  
Impact parameter range: 0 - 14.5 fm.  
Events: 5M  
ECAL excluded to speed-up the calculations.

dEdx vs P for all particles



$m^2$  vs P for all particles



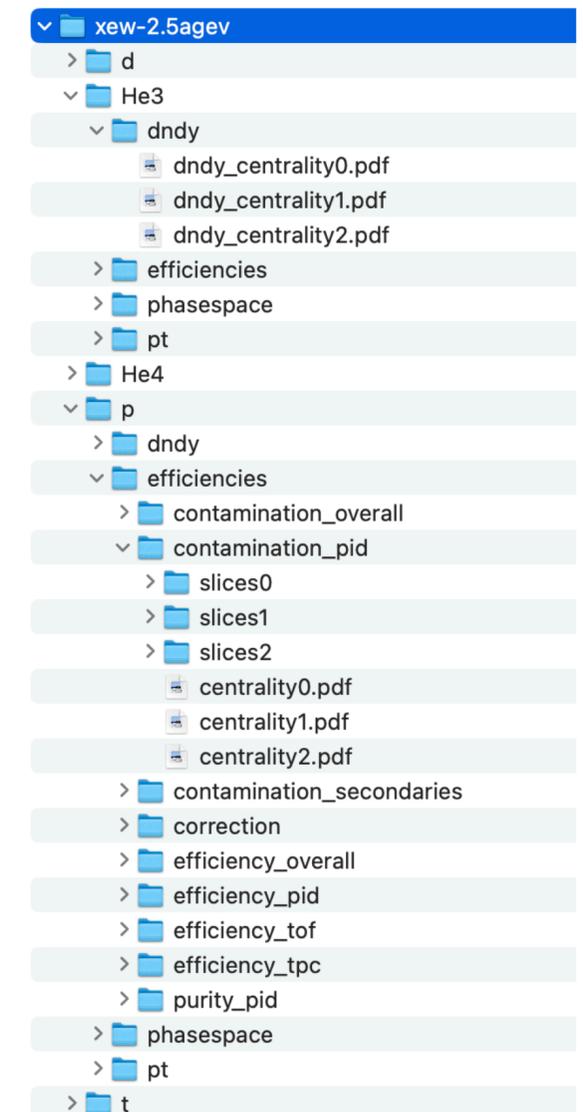
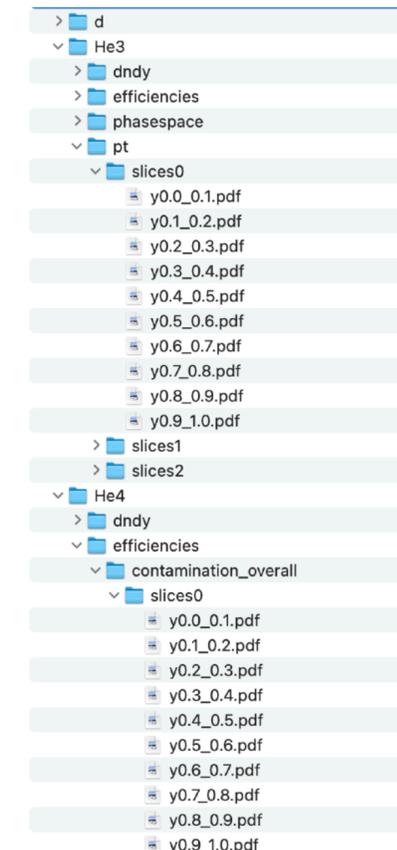
# Utilisation

*End of the day*

- After running the whole analysis chain you obtain **thousands of plots**.
- Here only few are presented to show the proof of work.

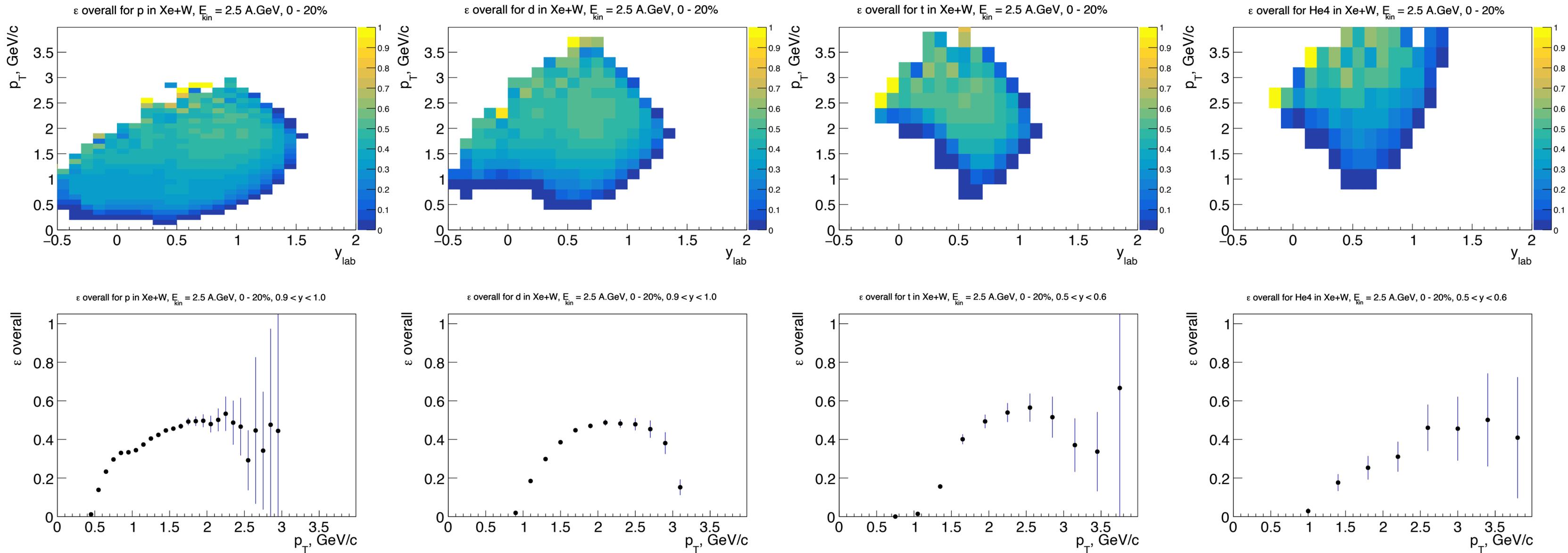
```
$ find . -name *.pdf | wc -l  
1695
```

339 plots per single specie



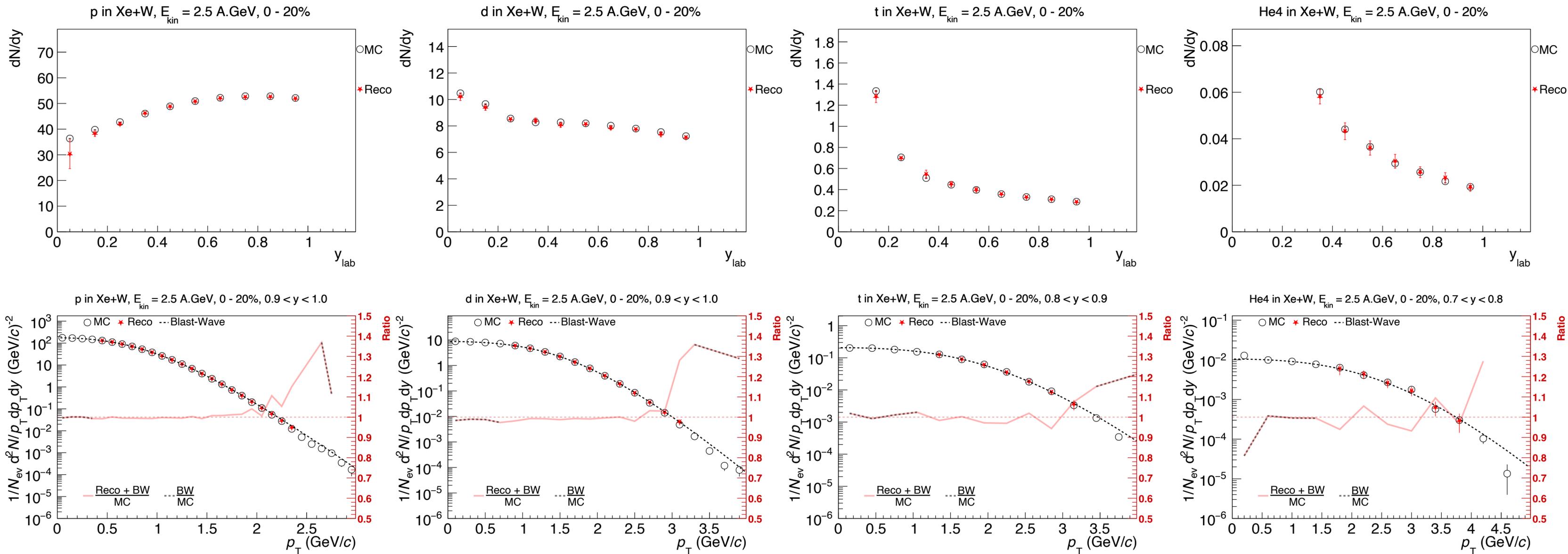
# Utilisation

## *Efficiencies*



# Utilisation

Spectra:  $p_T$ ,  $dN/dy$



# Utilisation

*Room for improvement*

- PID criteria: purity (contamination), efficiency, low transverse momentum part
- Blast-Wave fit parameters and form: low and high transverse momentum part,  $dN/dy$  points.

Sounds easy, but actually can take an indefinite amount of time.

**Good news: all machinery is set up.**

# Summary

this ends soon already

# Summary

- We must **think about the future** — students, newcomers — and finally start to write the code in a «good» way:
  - Avoid hardcoded and «magic» numbers.
  - Put explanatory comments in the code.
  - Add the documentation which covers the whole analysis.

**It does not take so much time as you can think — but saves your time in the future.**

# Summary

- The new **config-driven analysis** paradigm was used for the «nuclei» wagon update:
  - Easy to implement.
  - Easy to use.
  - It saves a lot of time during the post-processing (after the MpdRoot train) analysis.

**«Put everything in the black box» paradigm is obsolete and dangerous.**

# Summary

- Despite the right approach used within the «nuclei» wagon, there is room for improvement:
  - PID part in the wagon C++ part must be refactored (reduce the code).
  - More options must be added to the analysis configuration file (post-processing analysis).
  - Small refactor of the post-processing programs for the heavies usage of the configuration file — it is a must.

**~1 month to implement and test**

**Fully config-driven approach**