

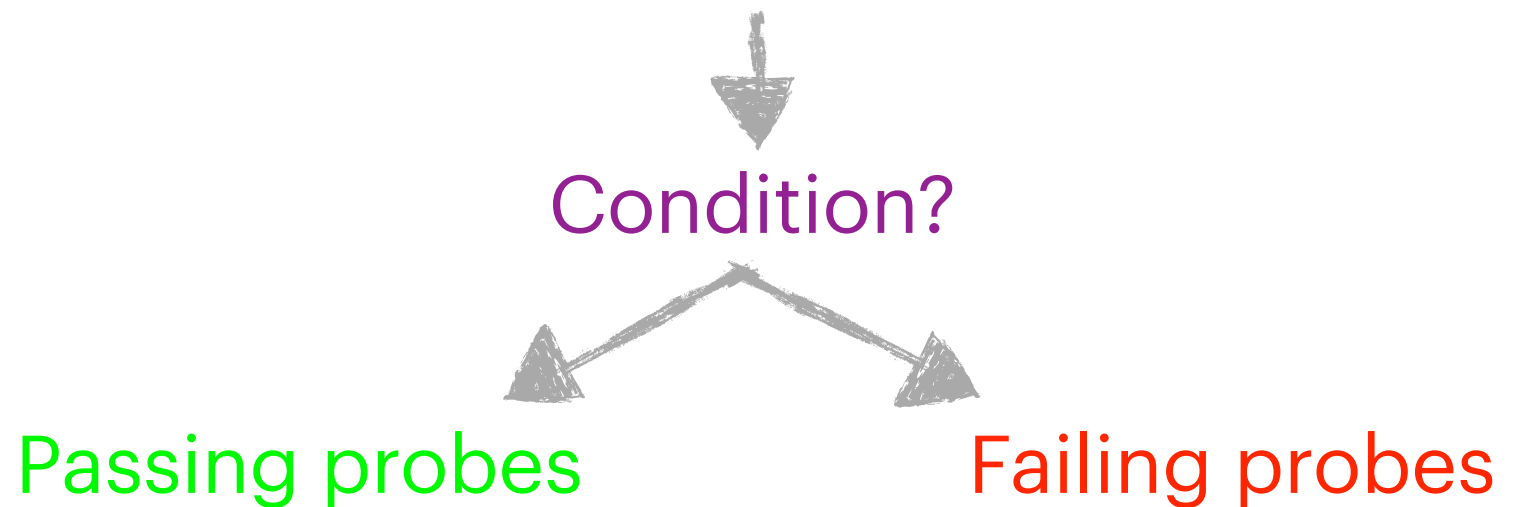
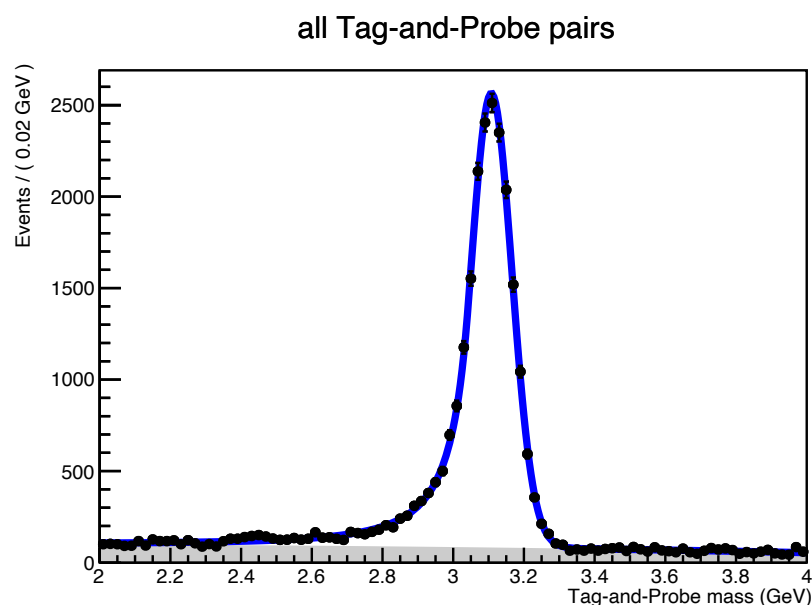
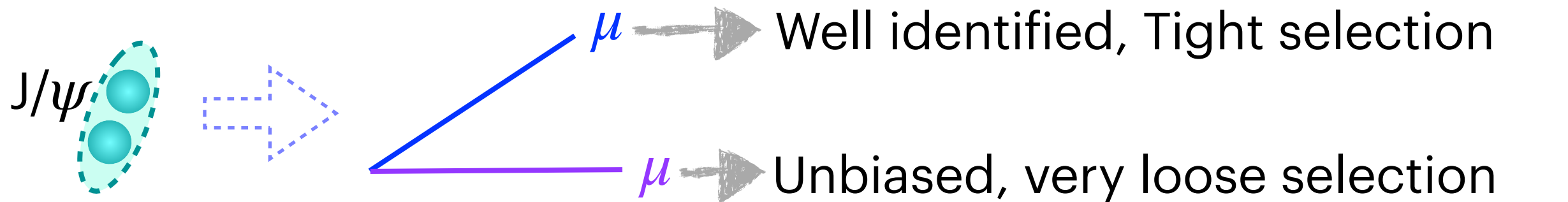
# Tag-and-Probe efficiency study for muons in Alice

O2 tutorial

Batoul DIAB - 08/11/2022

# The Tag-and-Probe method

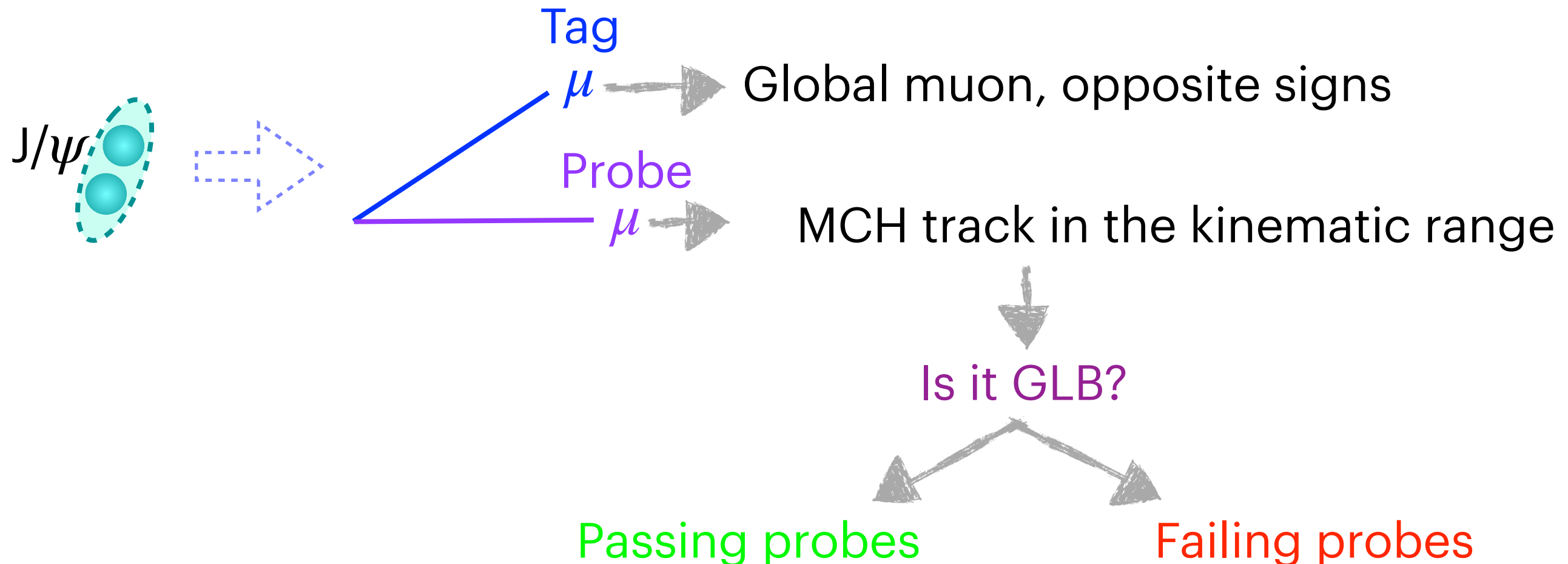
- Tag-and-Probe (T&P) is a data-driven efficiency calculation technique
- Simulations are not ideal → need data calibration
- based on the decays of known resonances, e.g.  $J/\psi$



$$\varepsilon = \frac{\text{probes passing condition}}{\text{all probes}}$$

# The starting point

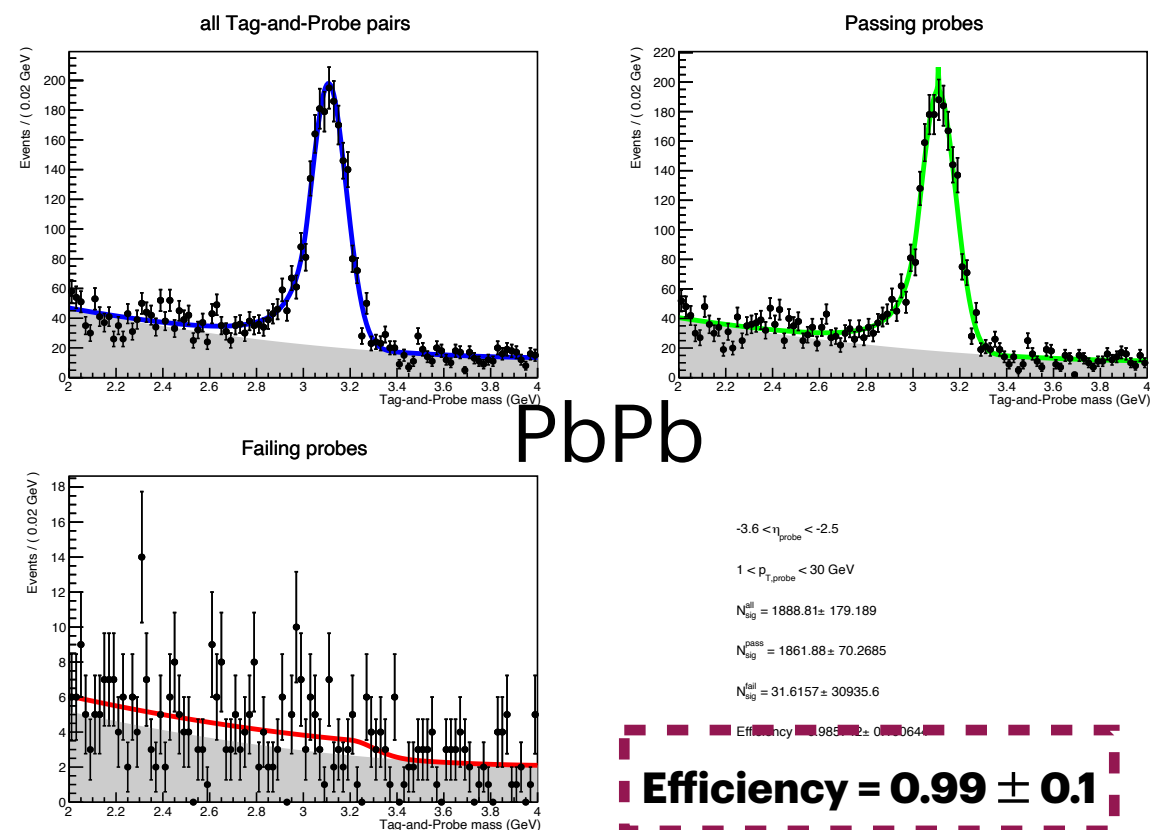
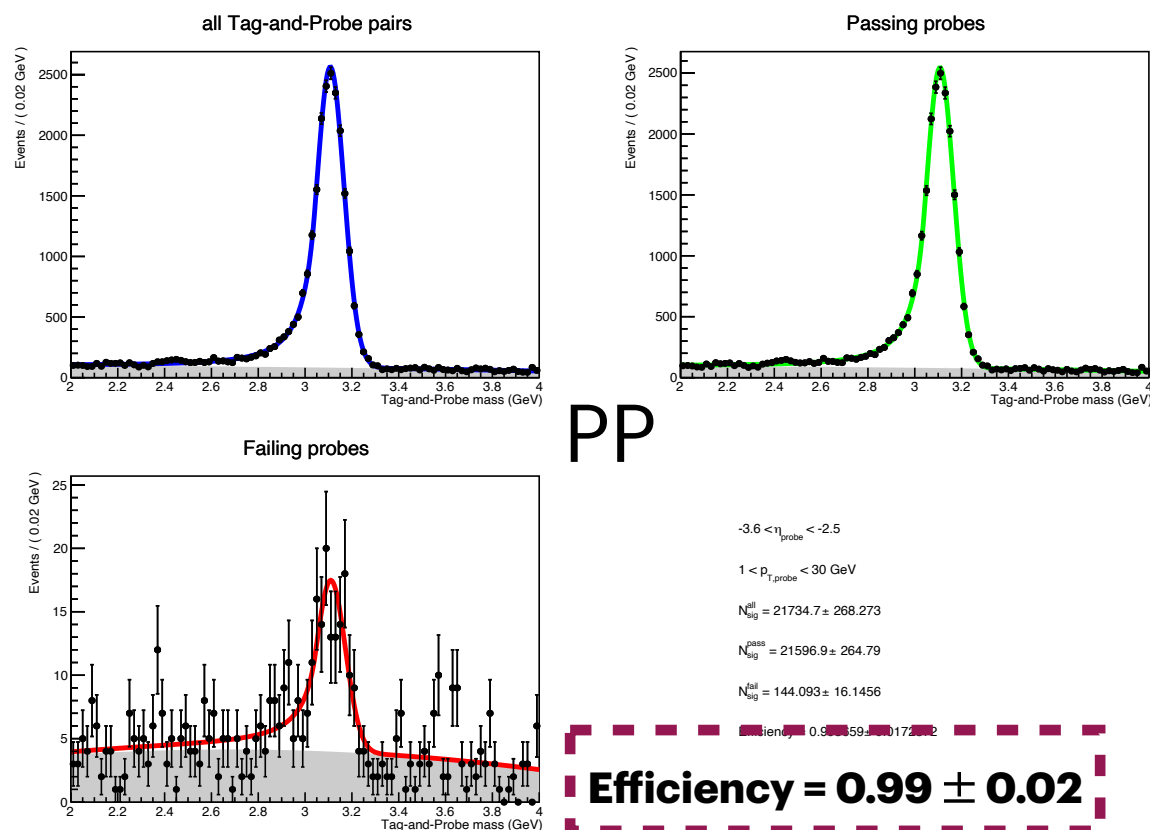
- For now we only have MC simulation
- Using the nonprompt  $J/\psi$  simulations in pp ([LHC21i3g](#)) and PbPb ([LHC22b2a](#))



- All kinematic quantities ( $p_T$ ,  $\eta$ ,  $M$ ...) are taken from the MCH tracks
- No additional criteria are applied on the quality of the tracks (e.g.  $\chi^2$ )
- In PbPb there is no centrality spectrum correction. Centrality is flat here which does not realistically represent data

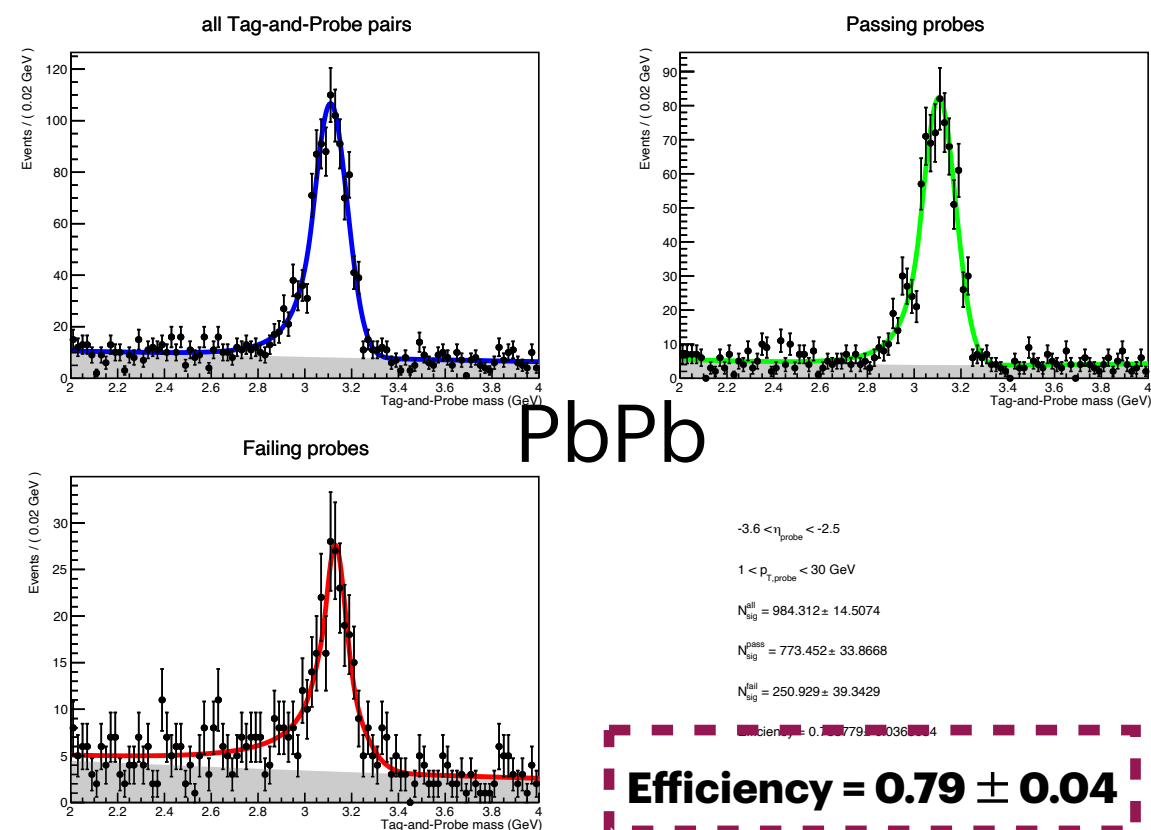
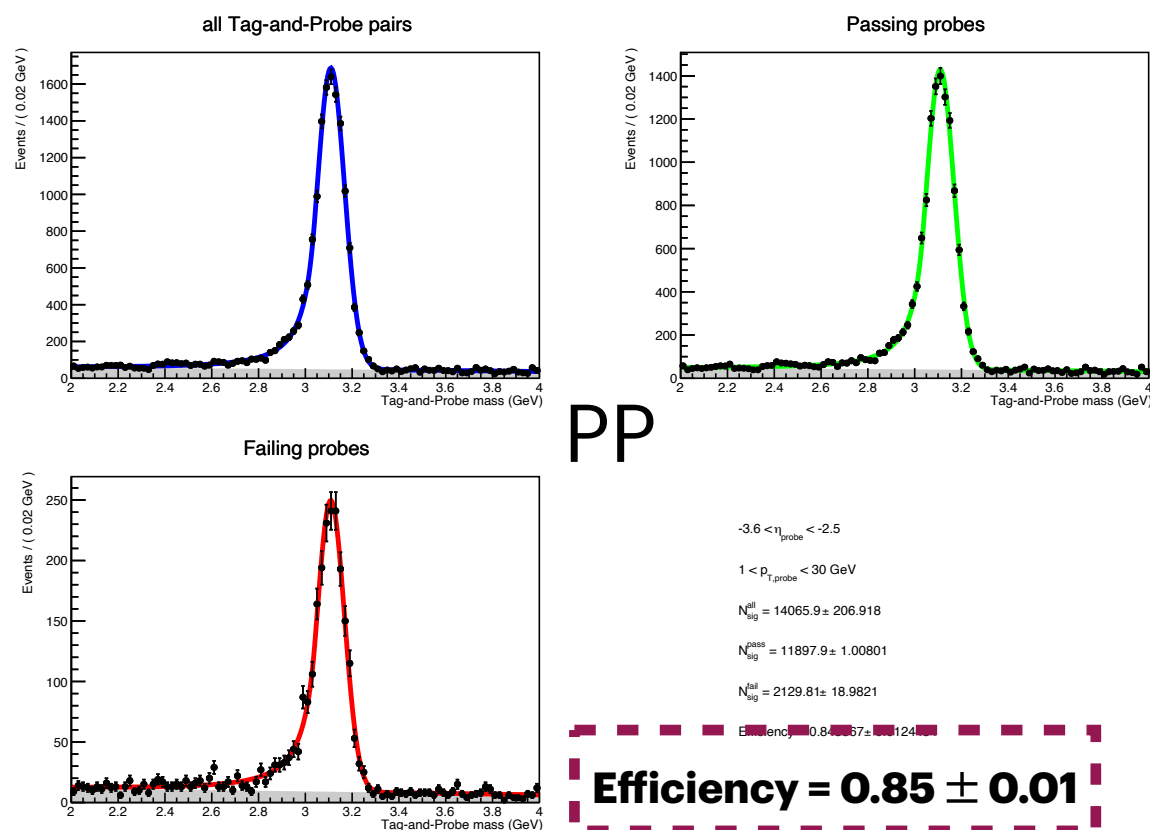
# Tag-and-probe fits

- Fits on the invariant mass of the T&P pairs, done with the rooFit package
- The fits are done for three categories: **all probes**, **passing probes** and **failing probes** but only **all** and **passing** fits are taken into account in the efficiency calculations
- For the signal: two Crystal Ball functions and for the background: 1st order Chebychev polynomial
- The fits definitely need more work but they give a good estimate for now
- An unrealistic efficiency of 0.99 for both pp and PbPb



# Tag-and-probe fits

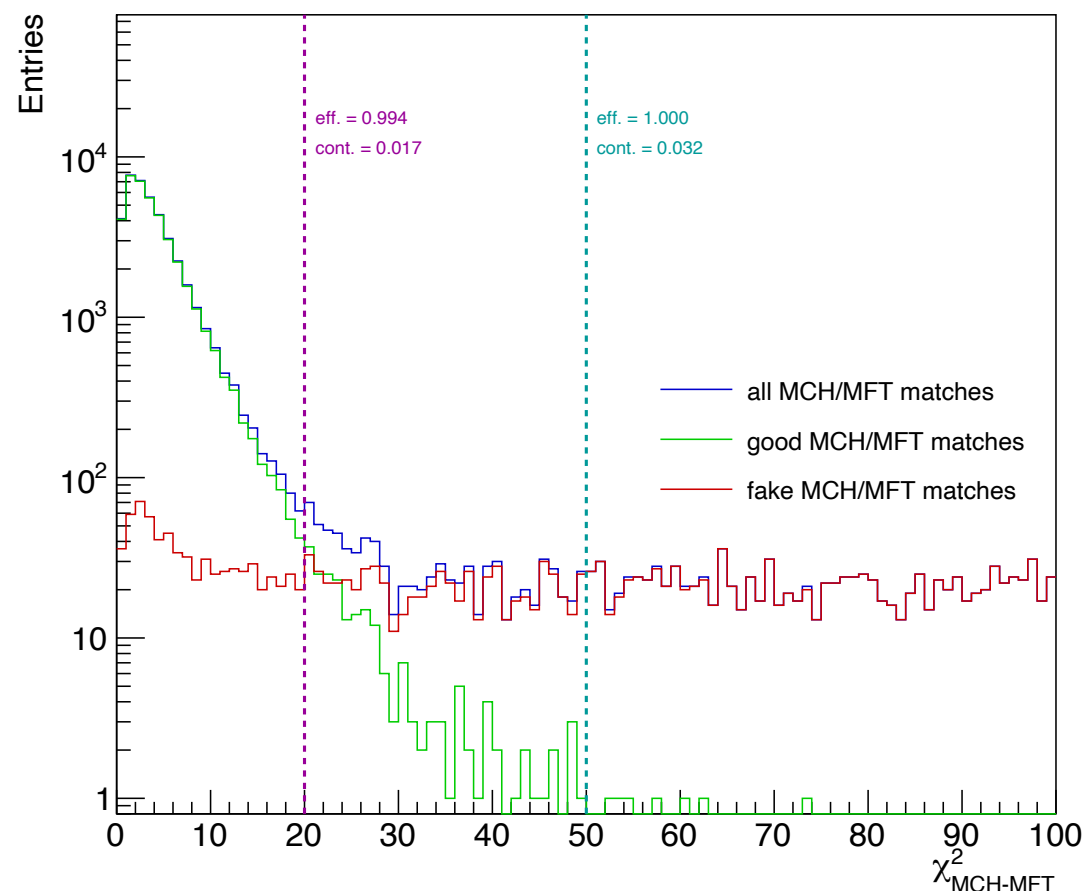
- It is not enough for the MCH to have a match in the MFT. It needs to be the right match
- In MC this information is available (🛑 will not be available in data)
- In these fits the match is required to be the correct one for both the tag and the passing probes
- A decrease in the efficiency from 0.99 to 0.85 in pp and 0.79 in PbPb (would even be lower in PbPb when the centrality distribution is corrected)
- The main challenge is not the inefficiency itself but the mismatching



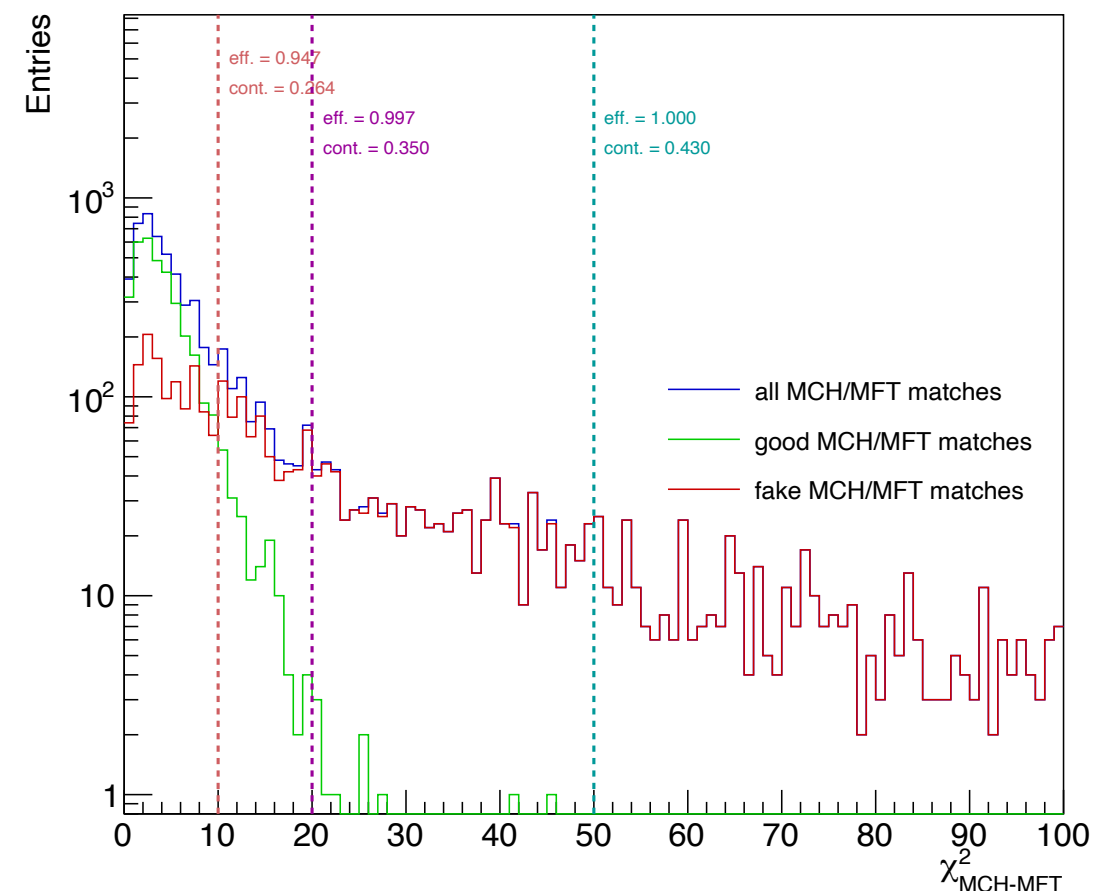
# Challenges for fake match rejection

- Need a discriminator that exist in data as well as MC
- Let's start by looking at the  $\chi^2$  of the MCH/MFT match in MC for good and bad matches and look for a pattern
- In PP: big peak of good matches at low  $\chi^2$ , almost flat distribution of bad matches
- Harder to separate in PbPb
- With a simple cut on the  $\chi^2$  we need to know the inefficiency and the contamination
- Not a good solution especially in PbPb
- Machine learning will be used to reject fake matches

PP,  $1 < p_T < 20$  GeV,  $-3.6 < \eta < -2.5$

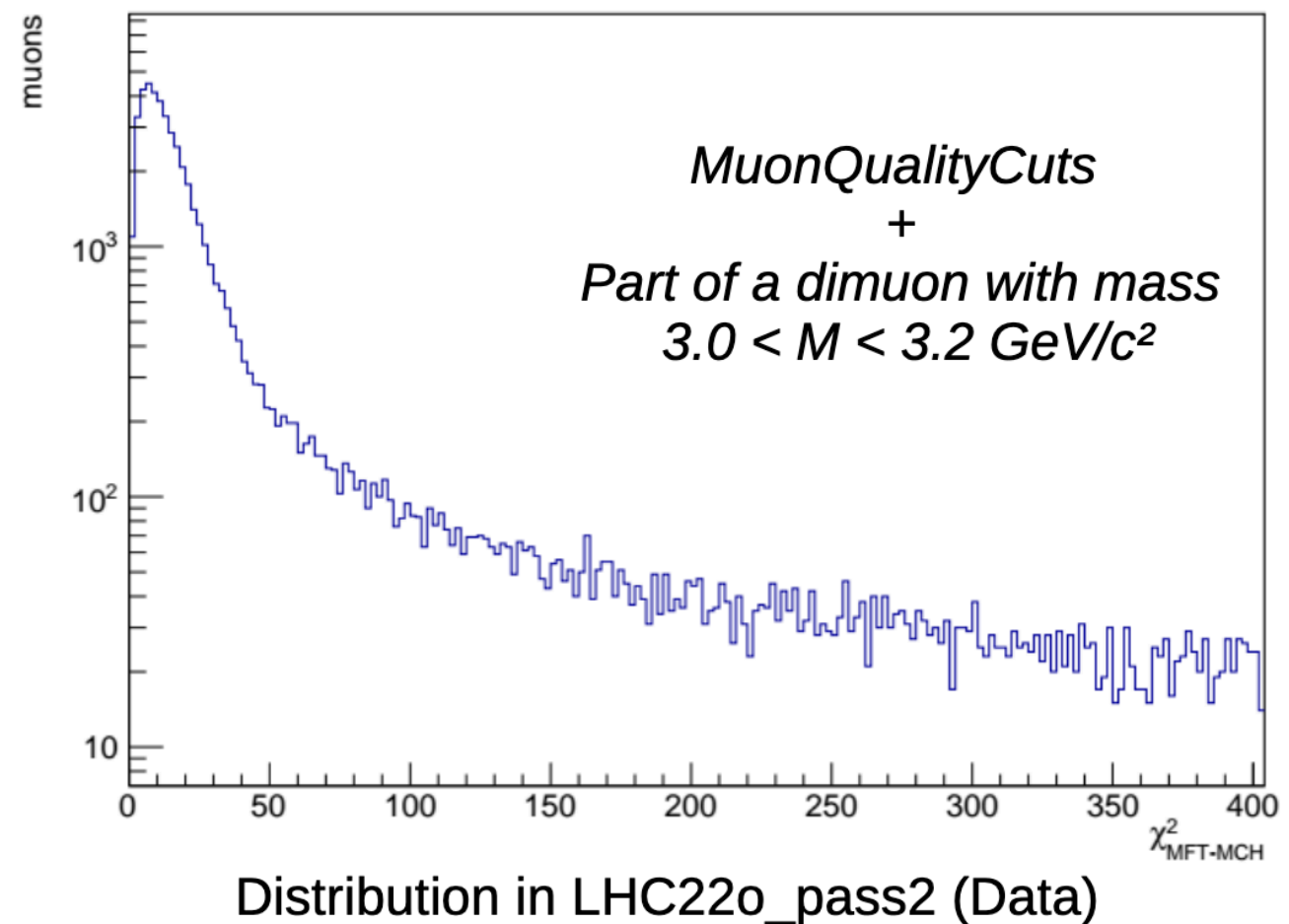
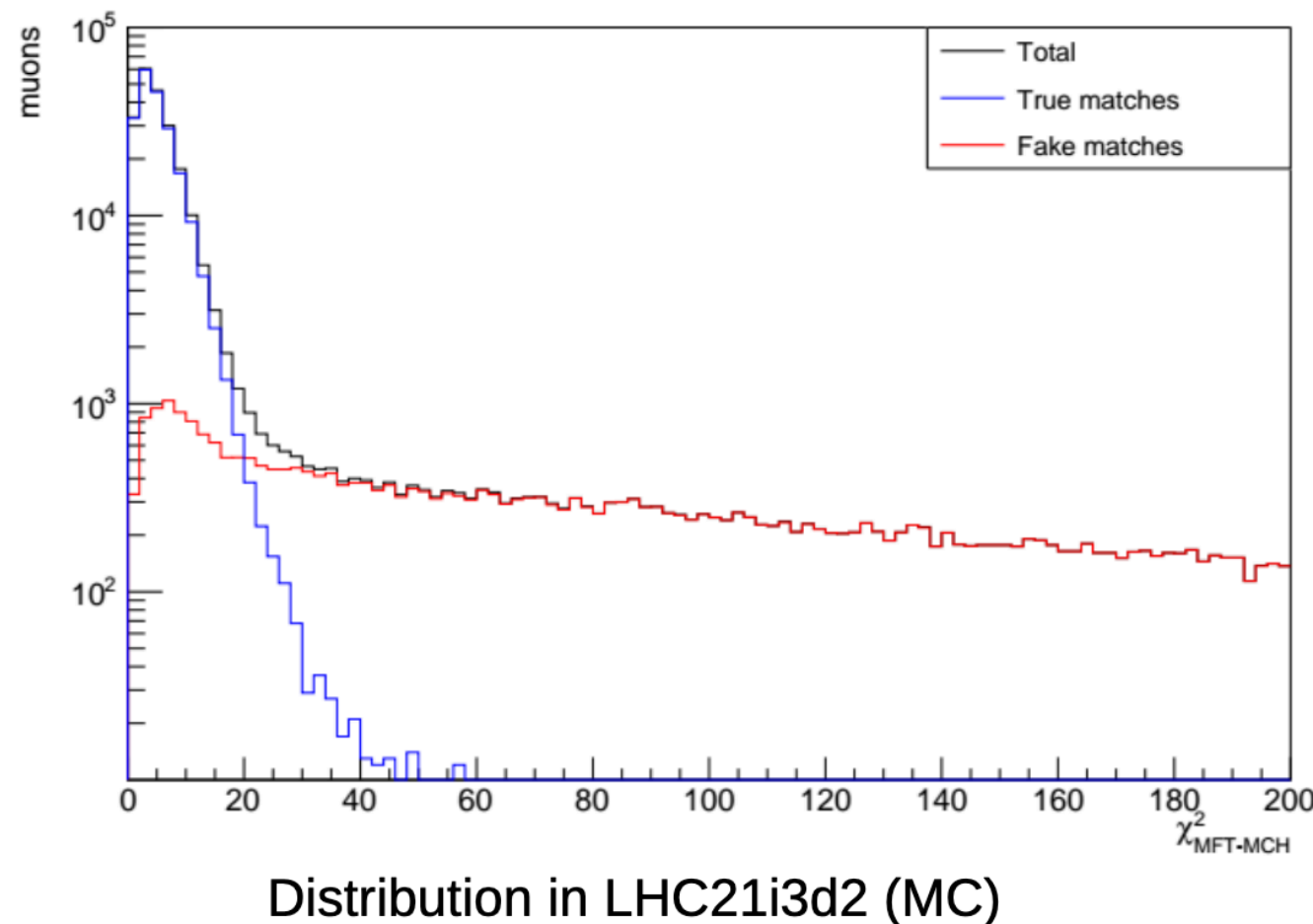


PbPb, Cent. 0-100,  $1 < p_T < 20$  GeV,  $-3.6 < \eta < -2.5$



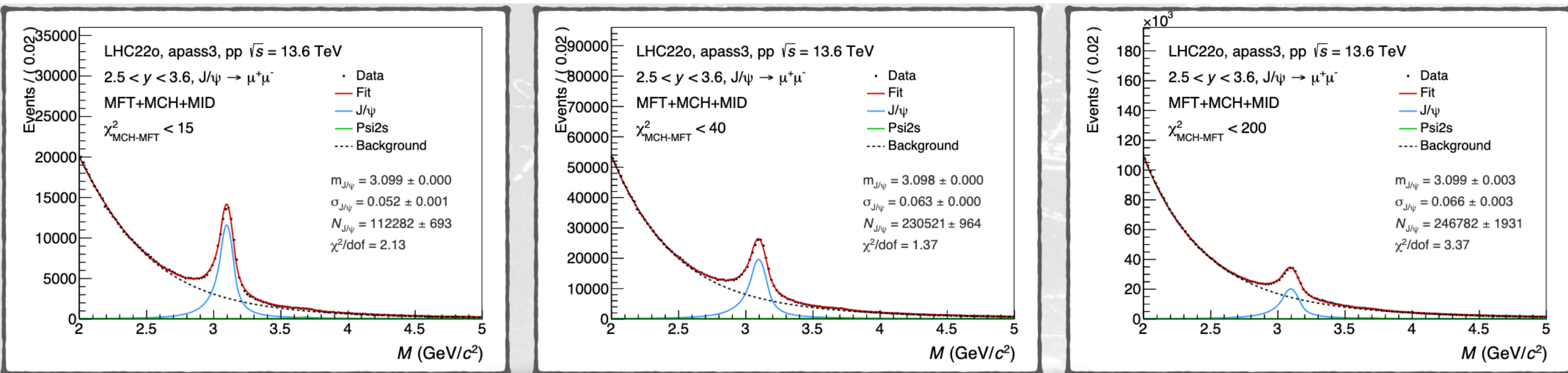
# $\chi^2$ matching for $J/\psi$ analyses

- In the  $J/\psi$  mass window, the  $\chi^2$  distribution starts resembling the MC shape: peak + background
- Different widths of the distributions: might be closer with a better MC
- Two tasks: chose a cut on  $\chi^2$  and estimate the efficiency + contamination



# Using the peak to chose a $\chi^2$ cut

- We use fits on the global dimuon invariant mass using different  $\chi^2$  cuts and compare the results
- These fits are done in apass3 but we check for each apass
- An overall improvement was actually observed for apass3 when compared to apass2
- We also check different run conditions: the IR has an impact on the performance

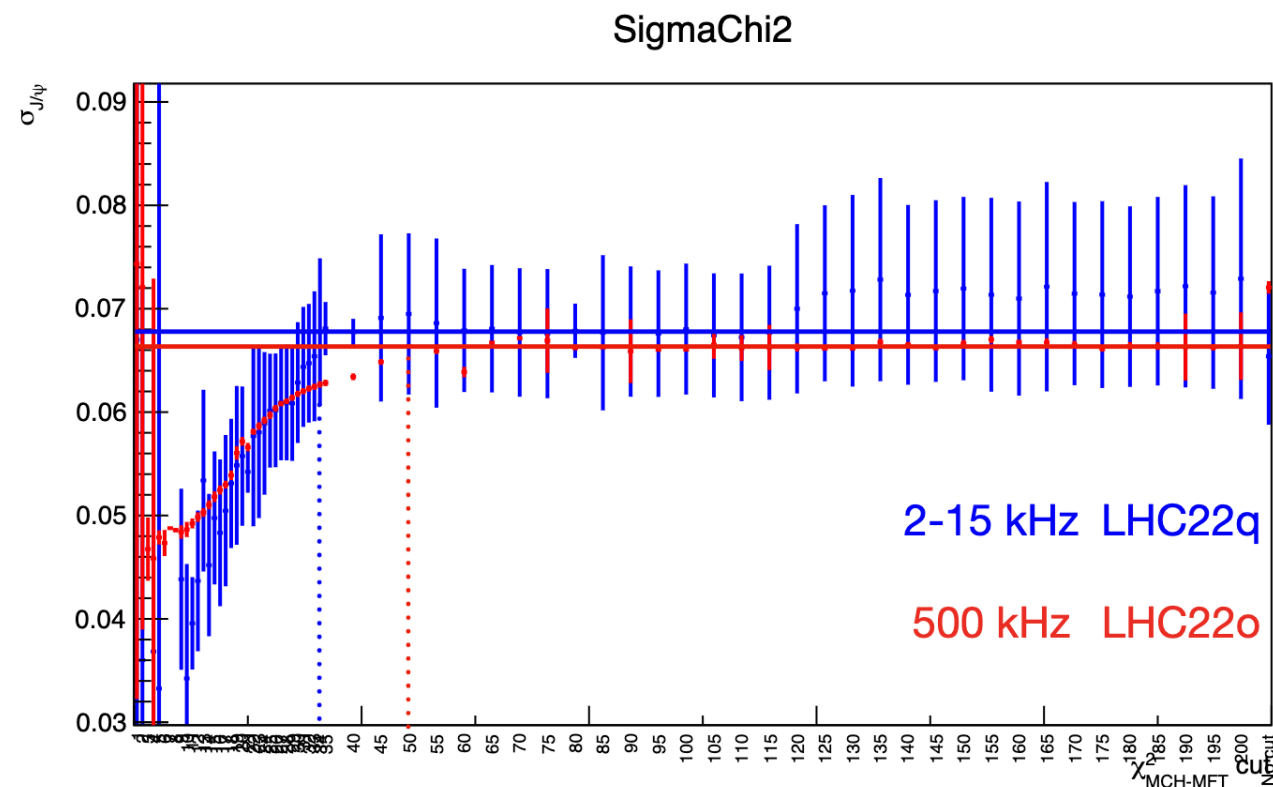


By N. Bizé

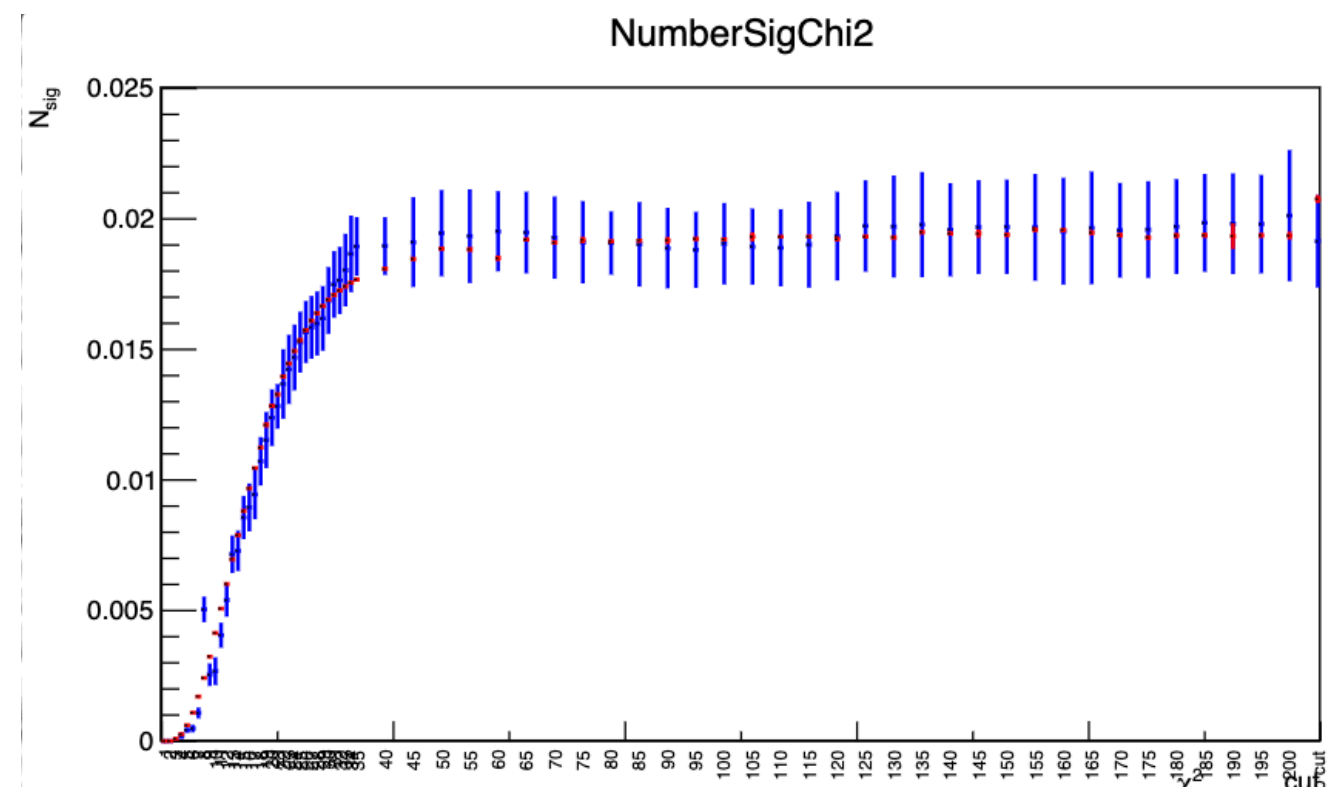


# Using the peak to chose a $\chi^2$ cut

- The resolution and the signal reach a plateau after a certain cut
- For both cases **LHC22q (low IR)** reaches the plateau faster than **LHC22o (high IR)**
  - 22q:  $\chi^2 < 35$
  - 22o:  $\chi^2 < 50$
- We still need to estimate the efficiency and the leftover contamination

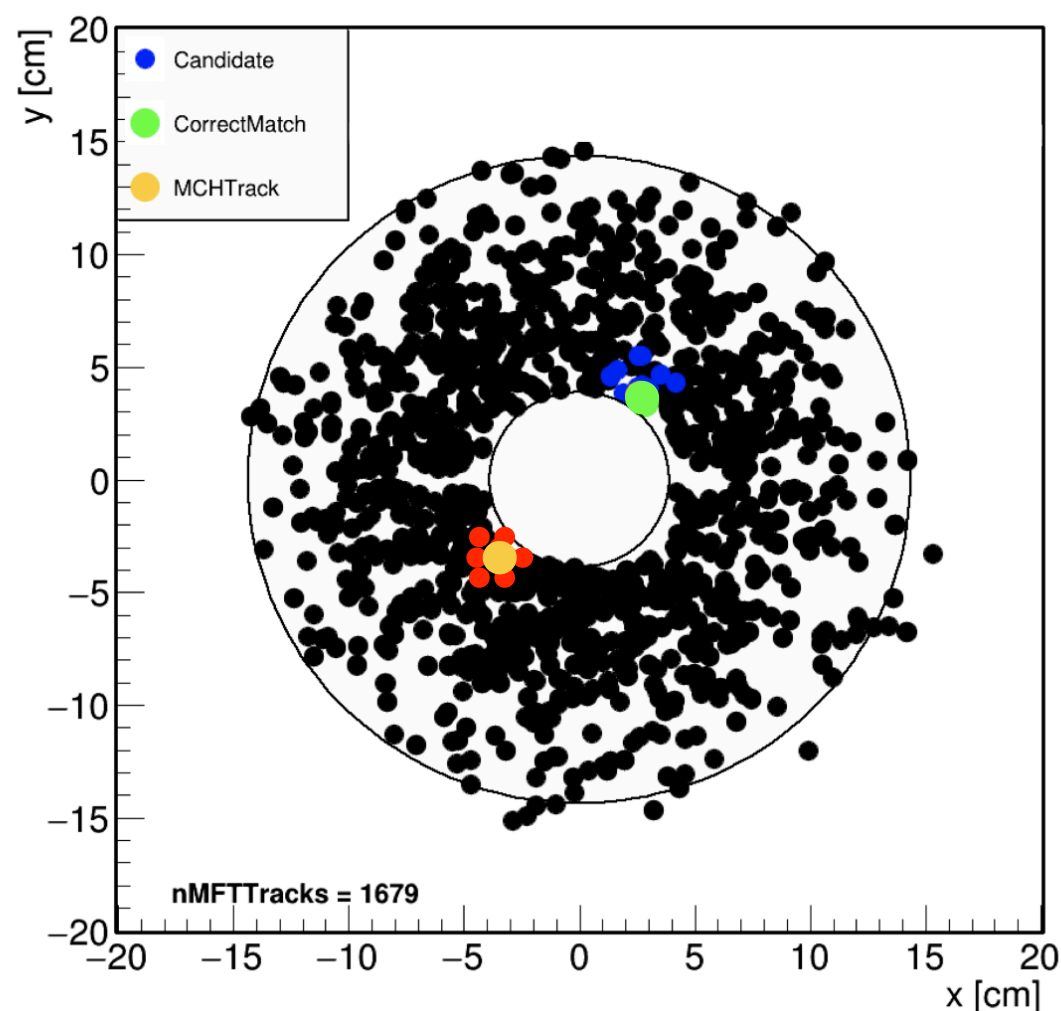


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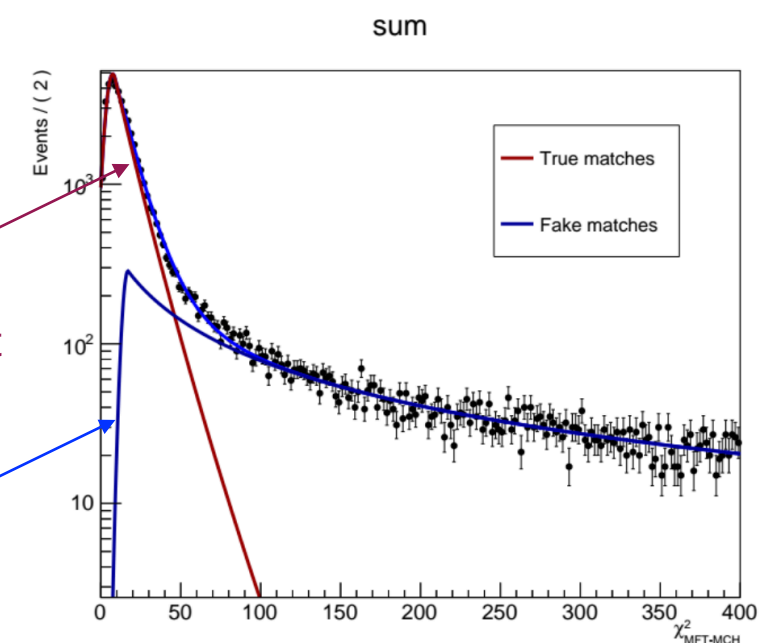
# New idea to get the efficiency

- A similar strategy can be used for the MFT-MCH mismatch shape
- Take the MCH and MFT candidates in our data, and rotate the MCH track (a  $\phi$  rotation should be good) and apply the exact matching procedure as normal
- Since all the matches are going to be fake, this will give us the shape of the fake matches
- From this shape we build a model (fit or template) of the fake matches and then use the total distribution in data to fit and get the good matches as well
- This will give the efficiency and contamination of our matching
- Can also be used in PbPb for the Machine Learning training data



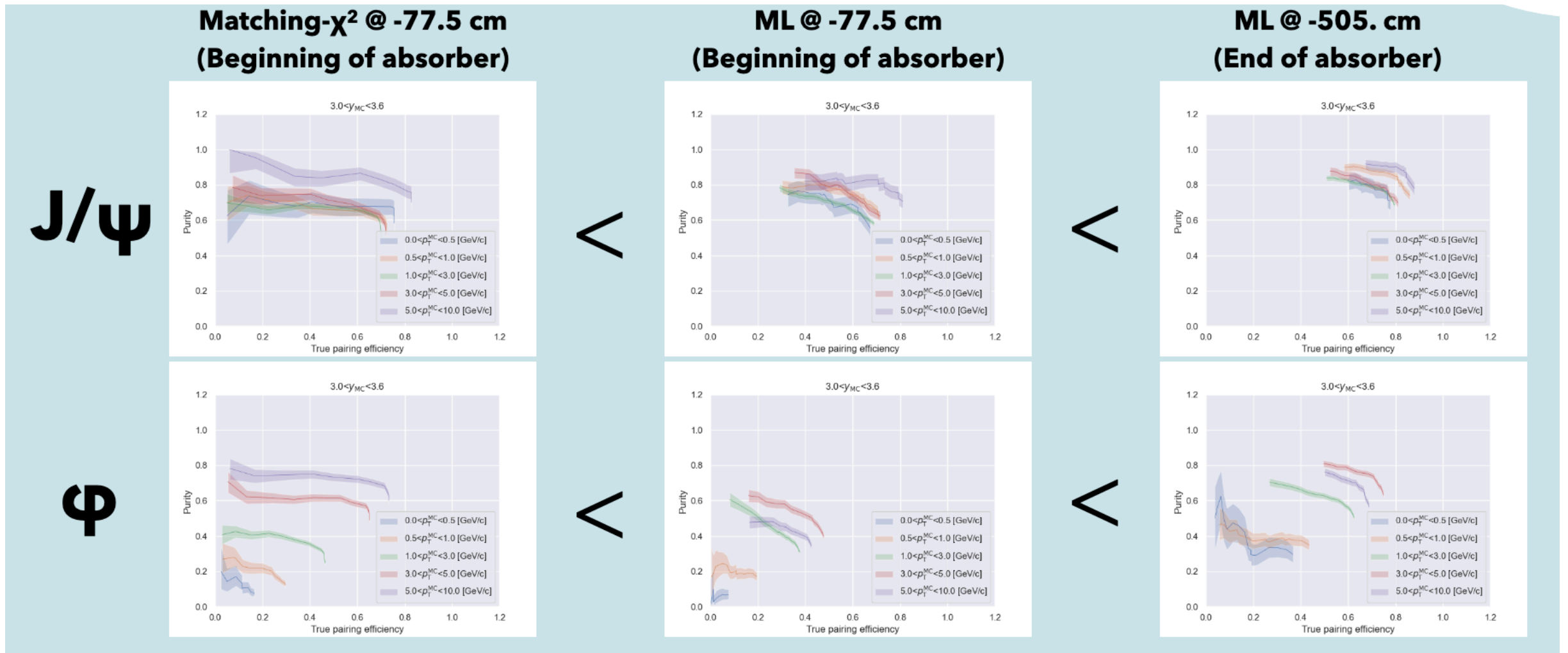
Get this shape from the fit

Get this shape from the rotation method

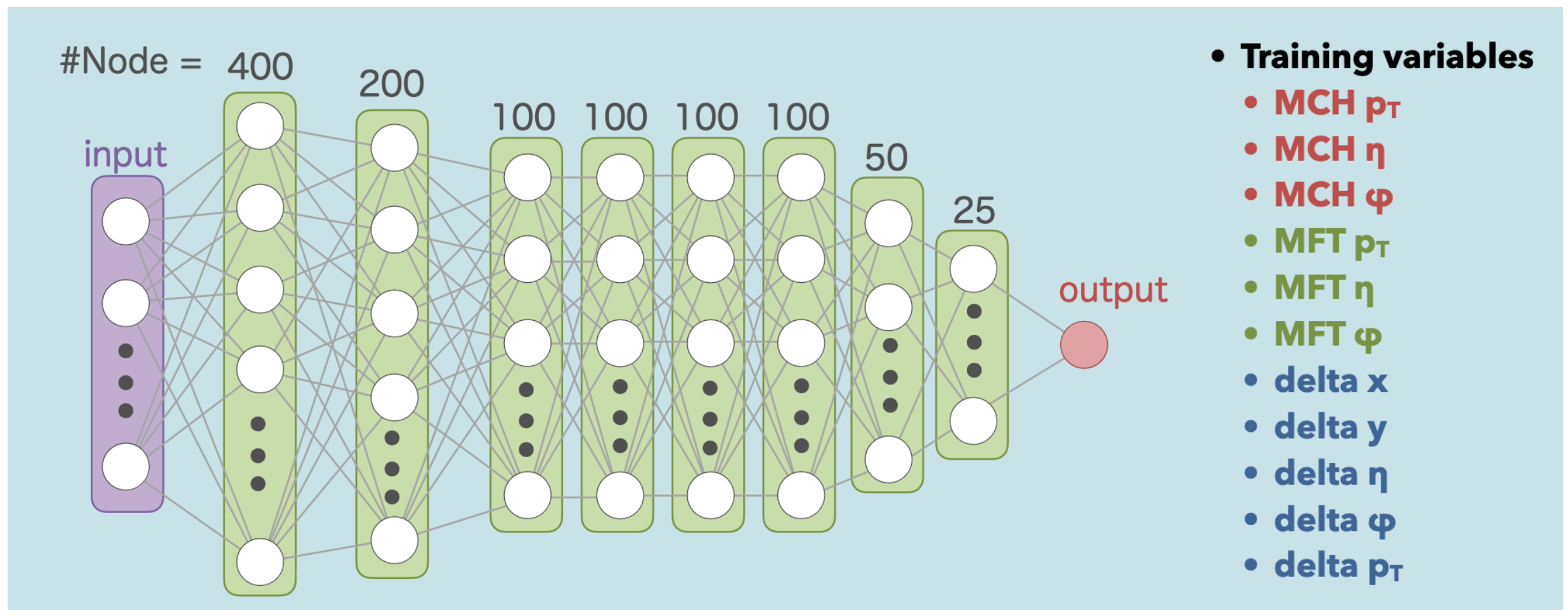


# Using ML for MFT-MCH matching

- The  $\chi^2$  performance is not good enough for low mass particles like  $\phi$
- We explored other options like using machine learning to improve the matching
- Important for low mass and can improve the  $J/\psi$  matching
- Different options are explored using AO2D



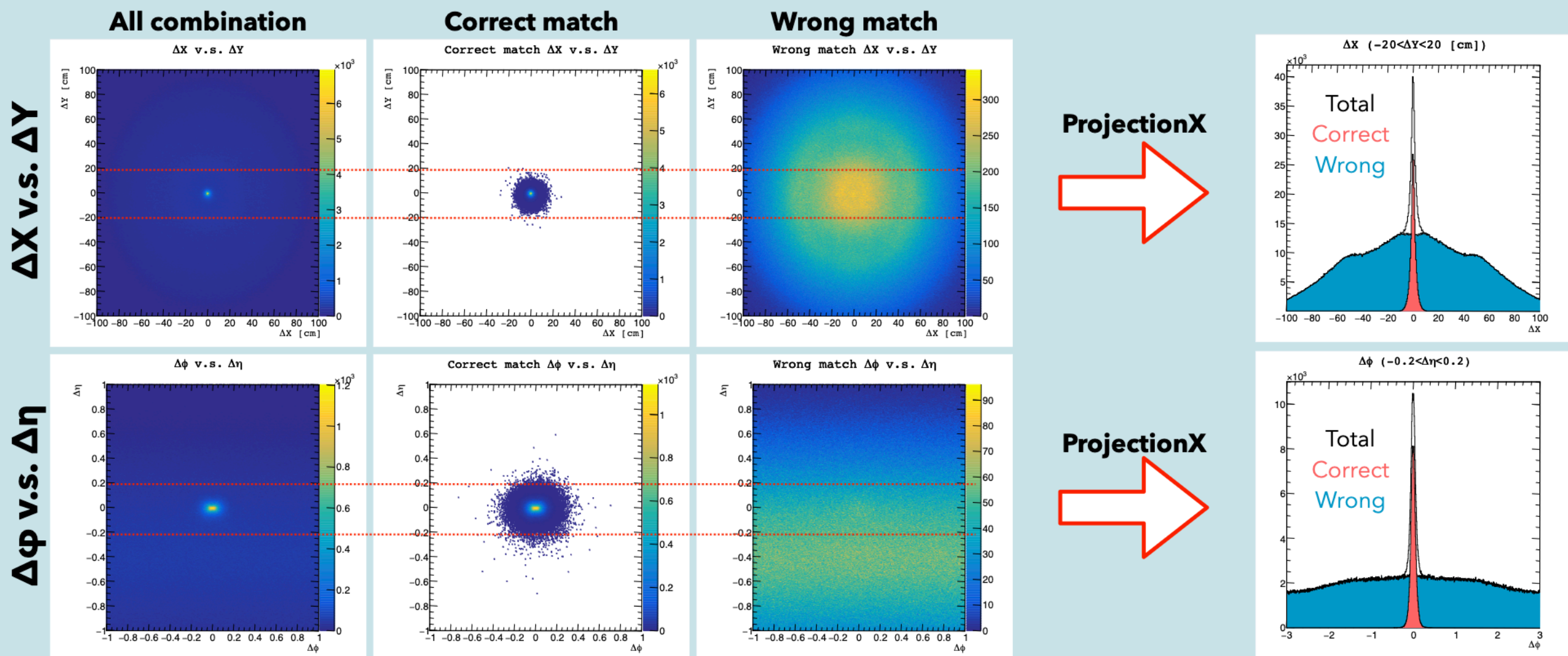
# Deep Neural Network model





# Matching plane choice

- The preliminary results show a good performance in MC
- The  $\Delta x$ ,  $\Delta y$  and  $\Delta\phi$  distributions show a peak of correct matches with a background of fake matches
- A similar shape is also seen in data but more tests need to be done
- Similar to  $\chi^2$ , we need to find a data driven method to estimate the leftover background: opposite sign matching



# Workflow for ML matching

## Purpose

MFT-MCH matching by machine learning based on the AO2D.root adaptable to the DQ framework

## New workflow: o2-analysis-mftmchmatchingml

- Match mfttrack and fwdtrack (MuonStandaloneTrack) by machine learning
- Create new table “FwdTracksML”

## New process function in table-maker: processMuonMLOnly

- Use FwdTracksML instead of FwdTracks to create reducedAOD

# o2-analysis-mftmchmatchingml

1. Select mfttrack and fwdtrack (MuonStandaloneTrack) from AO2D
2. Download onnx file from CCDB and get matching score for the selected track pair
3. Fill a new table named “**FwdTracksML**” if the track pair is true
  - The structure of table is same as FwdTracks
  - All tracks have TrackType=0 (GlobalMuonTrack)
  - Momentum is recalculated

$$\begin{aligned} \bullet \quad p_x &= p_{\text{MCH}} \sin\theta_{\text{MFT}} \cos\varphi_{\text{MFT}} \\ \bullet \quad p_y &= p_{\text{MCH}} \sin\theta_{\text{MFT}} \sin\varphi_{\text{MFT}} \\ \bullet \quad p_z &= p_{\text{MCH}} \cos\varphi_{\text{MFT}} \end{aligned}$$

## Configurables

```
Configurable<std::string> cfgCCDBURL{"ccdb-url", "http://ccdb-test.cern.ch:8080", "URL of the CCDB repository"};
Configurable<std::string> cfgModelDir{"ccdb-path", "Users/m/mooya/models", "base path to the ONNX models"};
Configurable<std::string> cfgModelName{"ccdb-file", "model_LHC22o.onnx", "name of ONNX model file"};
Configurable<float> cfgThrScore{"threshold-score", 0.5, "Threshold value for matching score"};
Configurable<int> cfgColWindow{"collision-window", 1, "Search window (collision ID) for MFT track"};
Configurable<float> cfgXYWindow{"XY-window", 3, "Search window (delta XY) for MFT track"};
```

# New process function in table-maker

```
void processMuonMLOnly(MyEvents::iterator const& collision, aod::BCsWithTimestamps const& bcs,  
                      soa::Filtered<aod::FwdTracksML> const& tracksMuon)  
{  
    fullSkimming<gkEventFillMap, 0u, gkMuonFillMap>(collision, bcs, nullptr, tracksMuon, nullptr, nullptr);  
}
```

- fullSkimming use FwdTracksML instead of FwdTracks
  - After reduced AOD is created, usage is the same as the case using FwdTracks
- 
- The effectiveness of the ML tool has not been verified with real data yet.
  - The DQ software team has pointed out that this strategy needs improvement because it completely replaces the existing fwdtrack, resulting in the loss of relevant information.



# Summary

- The Tag-and-Probe method will be used for the first time for muons in ALICE
- The focus will be the matching between the muon chambers and muon forward tracker
- The work started but still very preliminary
- Many challenges will be faced but they don't only concern Tag-and-Probe: MCH-MFT matching
- pp options:  $\chi^2$  matching and ML matching for low mass
- PbPb options: only ML
- Fake match rate estimation: track rotation and opposite sign matching

**Thank you**