

glbreco_v1

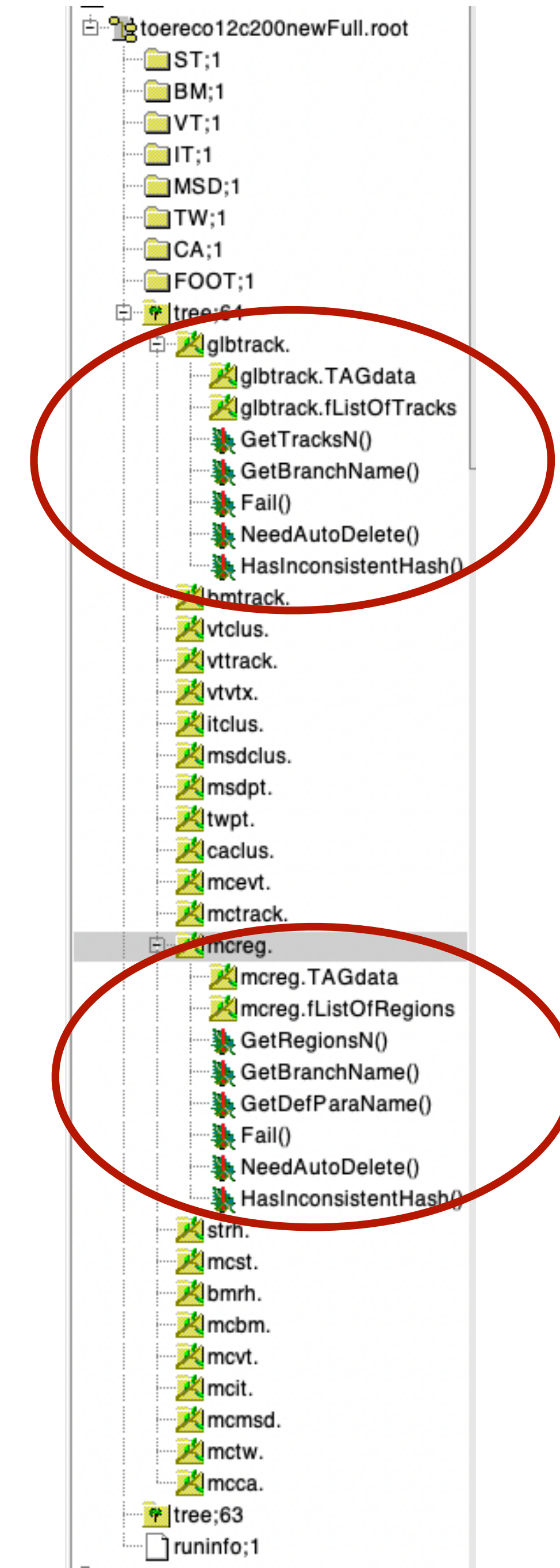
A general overview

Yun & Ilaria (for the moment)

Instructions

- Download the shoe code, checkout the **branch glbreco_v1** and do the usual build commands (cmake, source, make etc.)
- Get or create a root file with a ttree containing global tracks reconstructed from Genfit or Toe.
If it is a MC study, the including of crossing regions is highly recommended
- To launch the global analysis from builddir/Reconstruction/fullrec:

../bin/DecodeGlbAna -in toeorgenfitoutputfile.root -out
outputfilename -exp 12C_200 -run 1 -mc
(once we'll have the real data reconstruction, we'll omit the -mc
flag)
- Enjoy the output file with cross sections, systematic uncertainty
analysis and a first draft of a paper ready to be submitted... almost



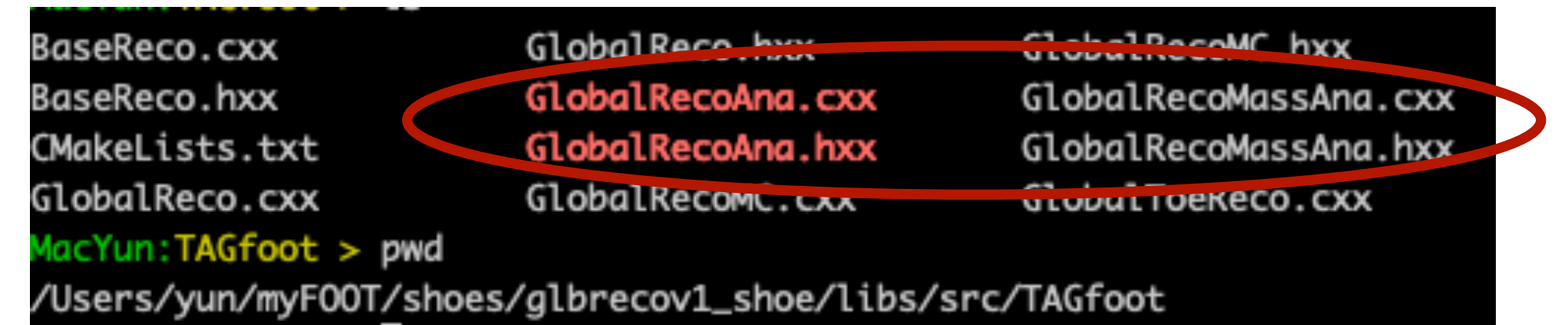
Overview

- The idea of a global analysis branch is to provide a tool with analysis methods and codes developed and verified by all the collaboration
- New analysis techniques can be easily added in the workflow.
- Once the cross section analysis has been done for a specific campaign, it can be easily adapted for all the other campaigns
- For the moment, all the analysis codes are in TAGfoot:
GlobalRecoAna.cxx/hxx and GlobalRecoMassAna.cxx/hxx
- The goal is the differential cross section:

$$\frac{d\sigma_i}{d\Omega}(\theta) = \frac{Y_i(\theta)}{N_C \times N_{TG} \times \Delta\Omega \times \epsilon_{trk}^i(\theta)}$$

$$\frac{d\sigma_i}{dE_{kin}}(E_{kin}) = \frac{Y_i(E_{kin})}{N_C \times N_{TG} \times \Delta E_{kin} \times \epsilon_{trk}^i(E_{kin})}$$

- For this presentation: no vtx pileup, triggered simulation, TW match

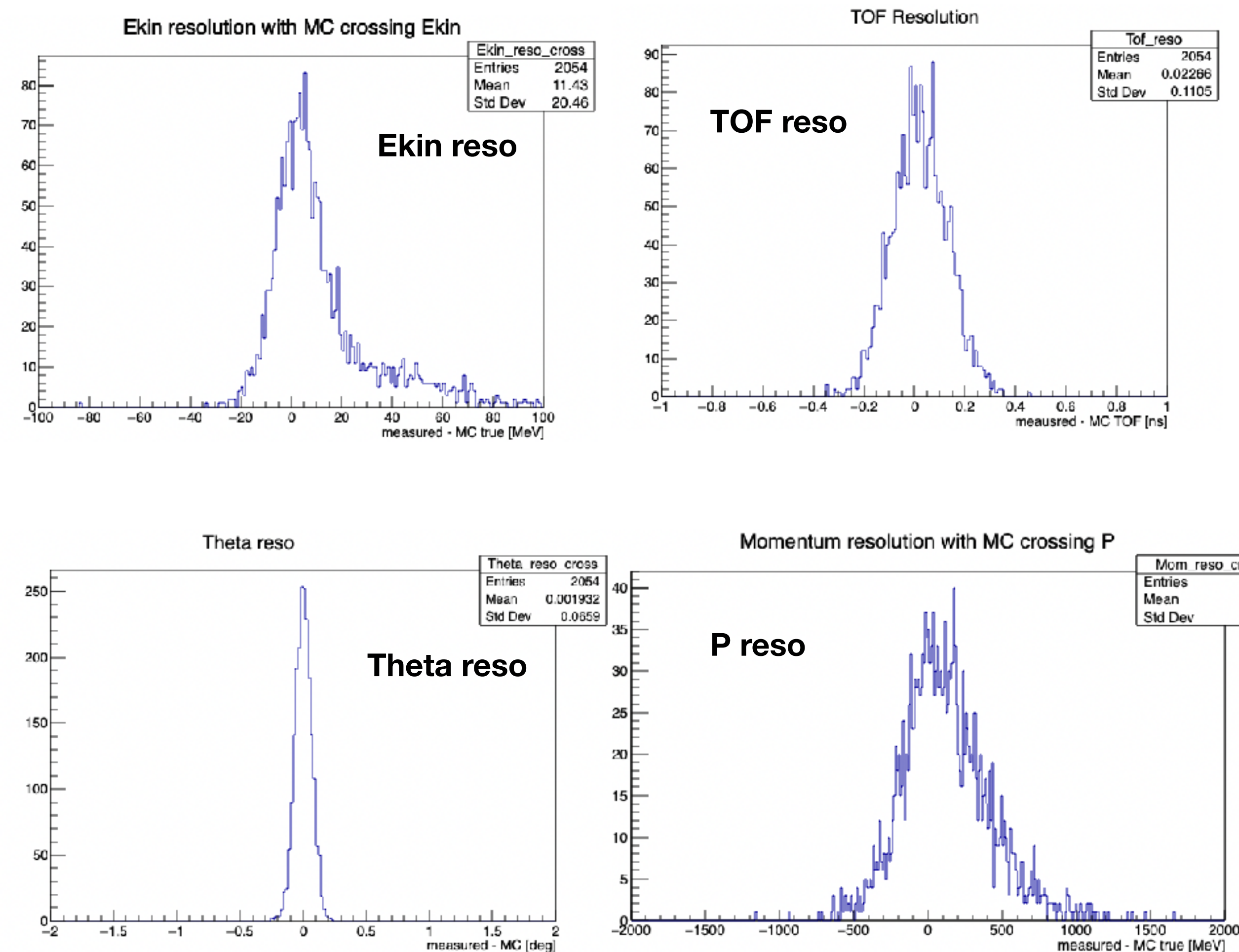
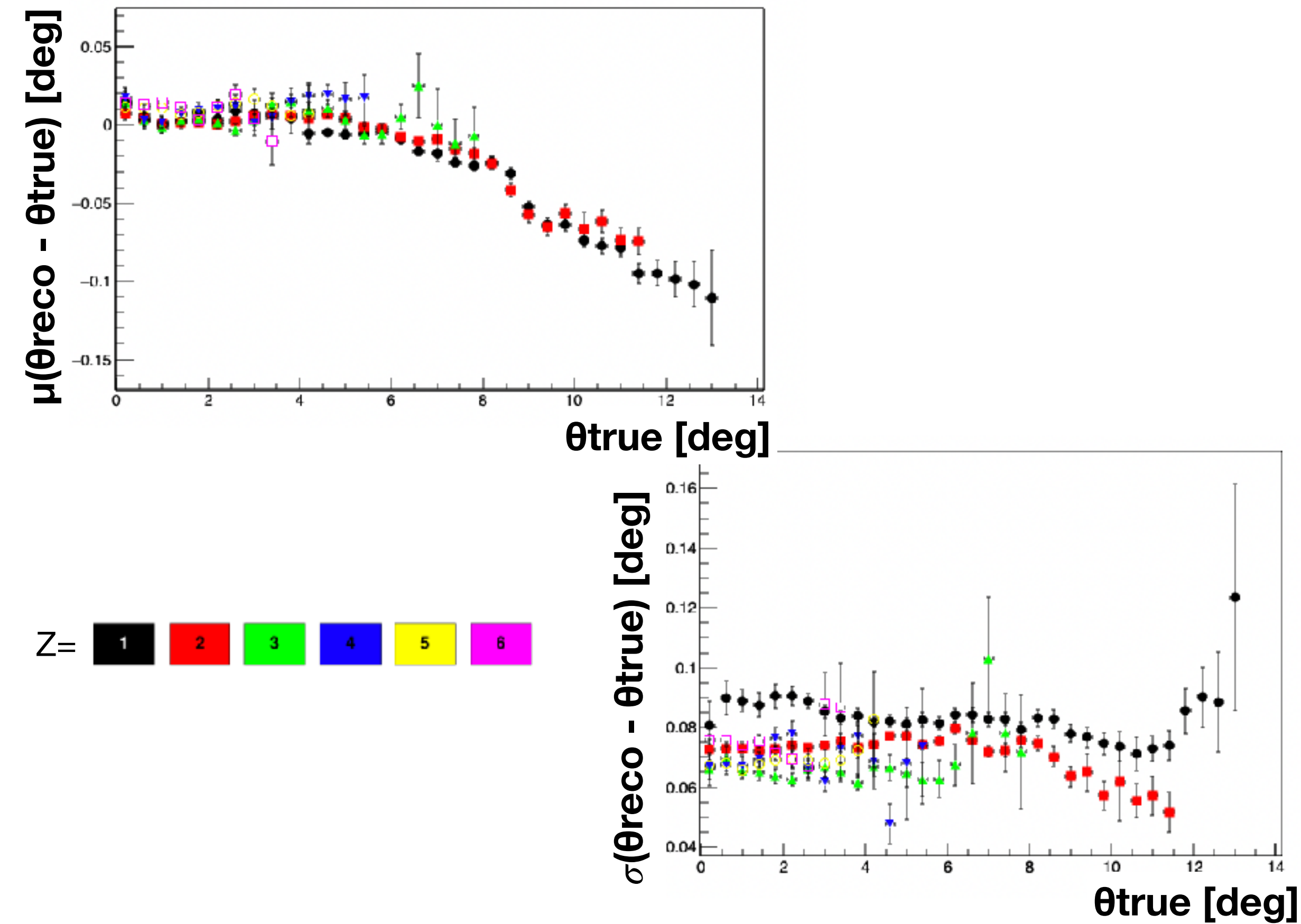


```

BaseReco.cxx      GlobalReco.hxx      GlobalRecoMC.hxx
BaseReco.hxx      GlobalRecoAna.cxx   GlobalRecoMassAna.cxx
CMakeLists.txt    GlobalRecoAna.hxx   GlobalRecoMassAna.hxx
GlobalReco.cxx    GlobalRecoMC.cxx    GlobalToeReco.cxx
MacYun:TAGfoot > pwd
/Users/yun/myFOOT/shoes/glbrecov1_shoe/libs/src/TAGfoot
  
```


Resolution studies

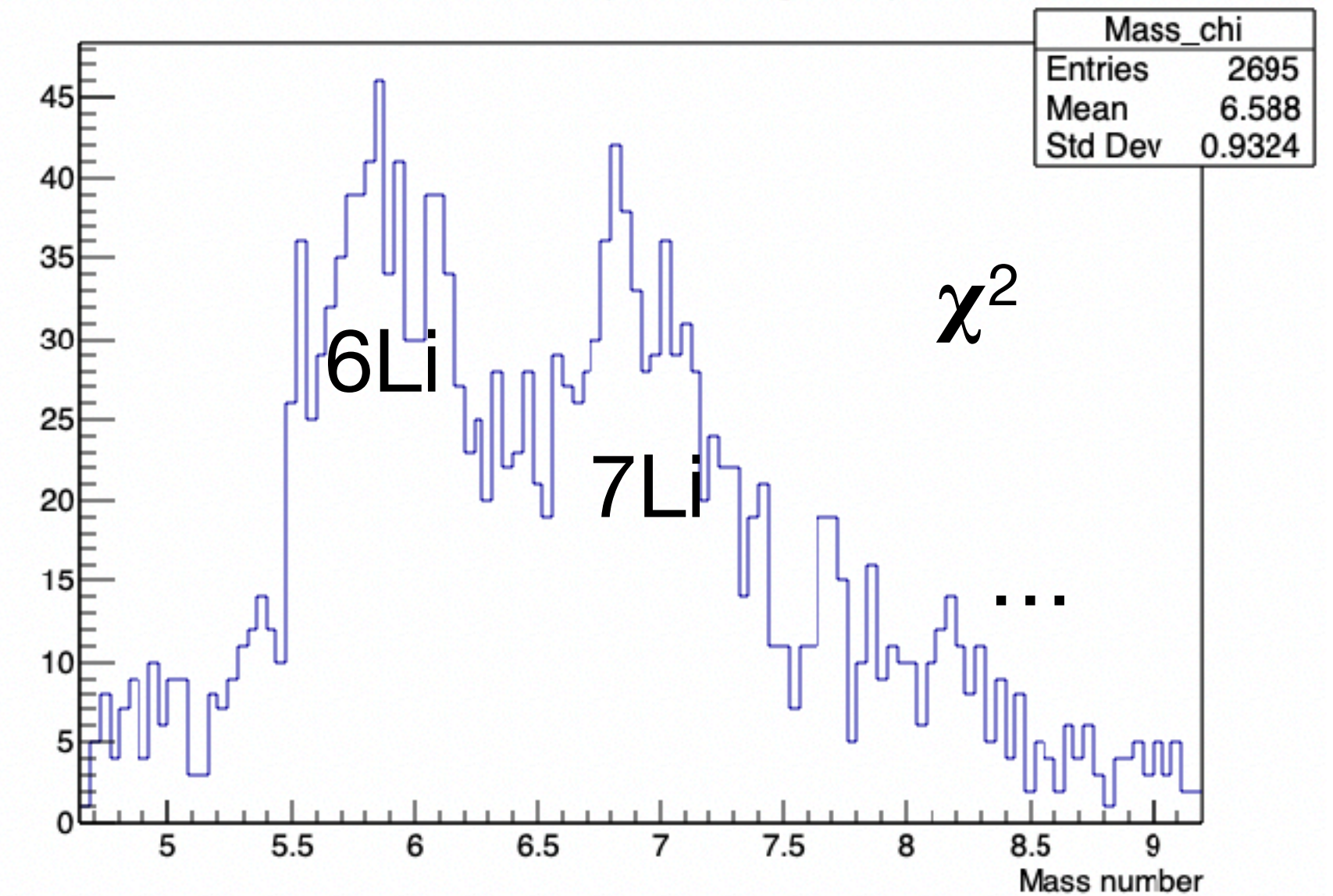
- Theta, Ekin, momentum and TOF resolution plots are calculated
- On the right the plots related to the **Z=3** reconstructed tracks are shown
- **Ekin** calculation needs to be checked



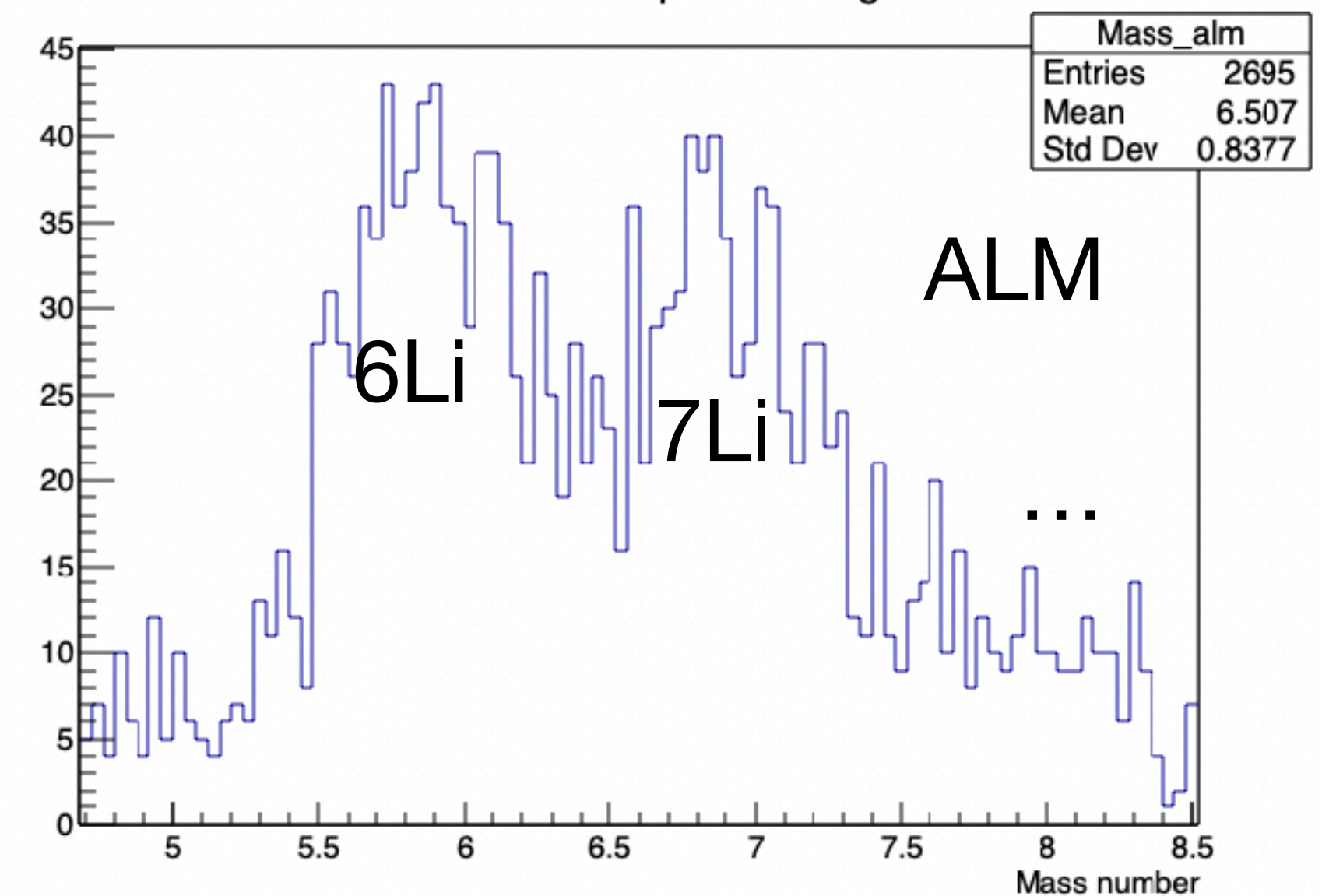
Mass reconstruction

- The mass reconstruction methods are developed in a **dedicated class**: GlobalRecoMassAna.cxx/hxx
- The input parameters are the foot TOF, P, Ekin measurements with the associated uncertainty
- The class provides an estimate of the mass with P-Ekin, TOF-P and TOF-Ekin measurements and It combines them with the Chi2 and Alm methods
- As output one can have 6 mass estimates with the associated error and the estimate of the TOF, P and Ekin from the chi2 and Alm algorithms
- At the moment the Ekin measurement from the Calo has to be checked with dedicated studies (see “calo studies” of this presentation)
- The input parameters (TOF, P, Ekin, Errors) need to be checked
- **The alm and chi2 minimisation parameters need to be studied and optimized**
- A check of the codes for errors, bugs etc. is more than welcome

Z=3 Reconstructed mass spectra using chi square method

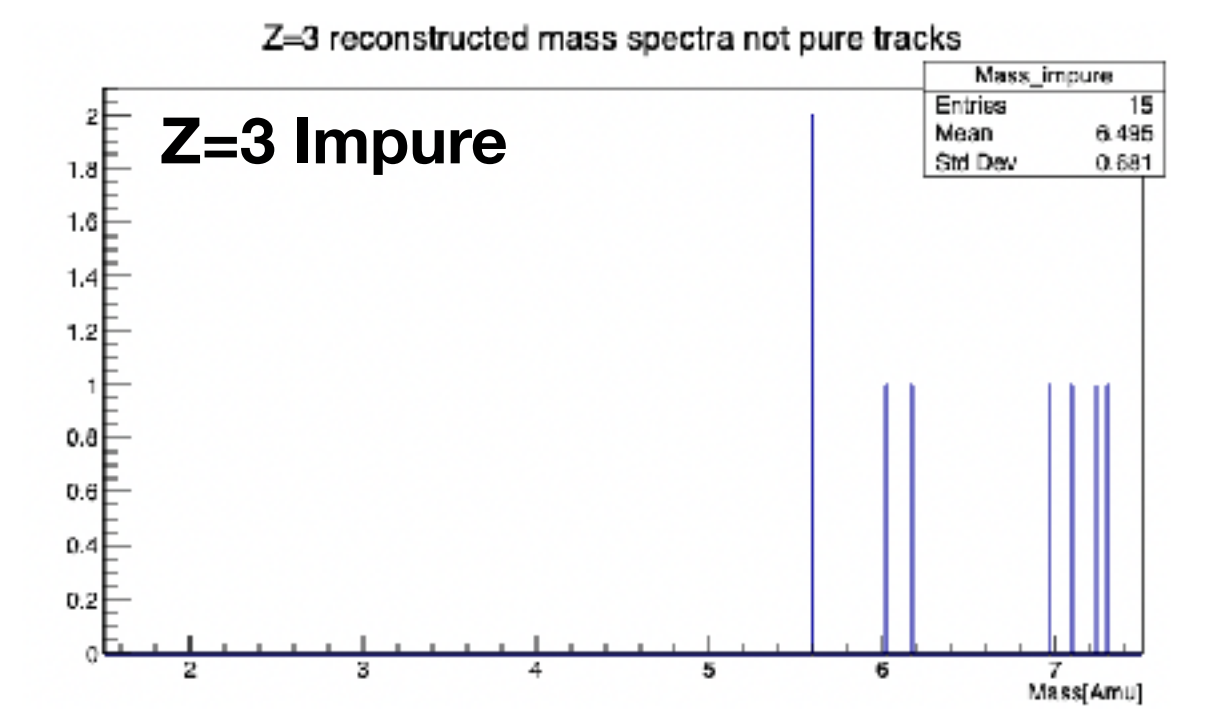
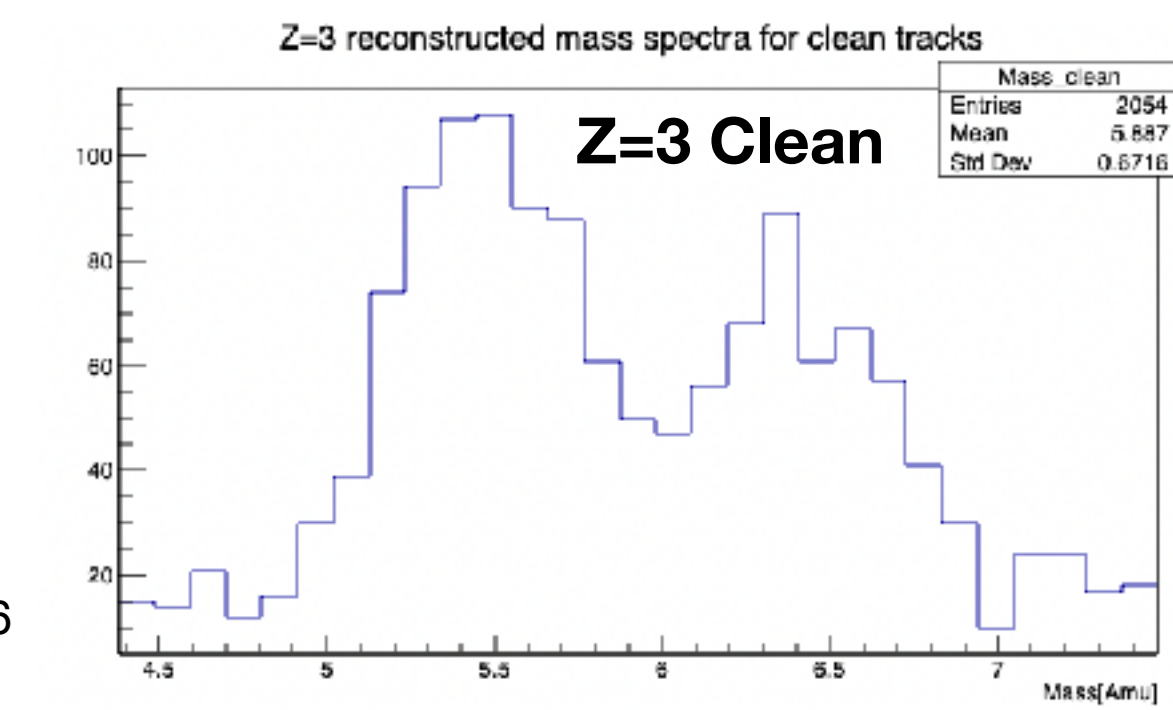
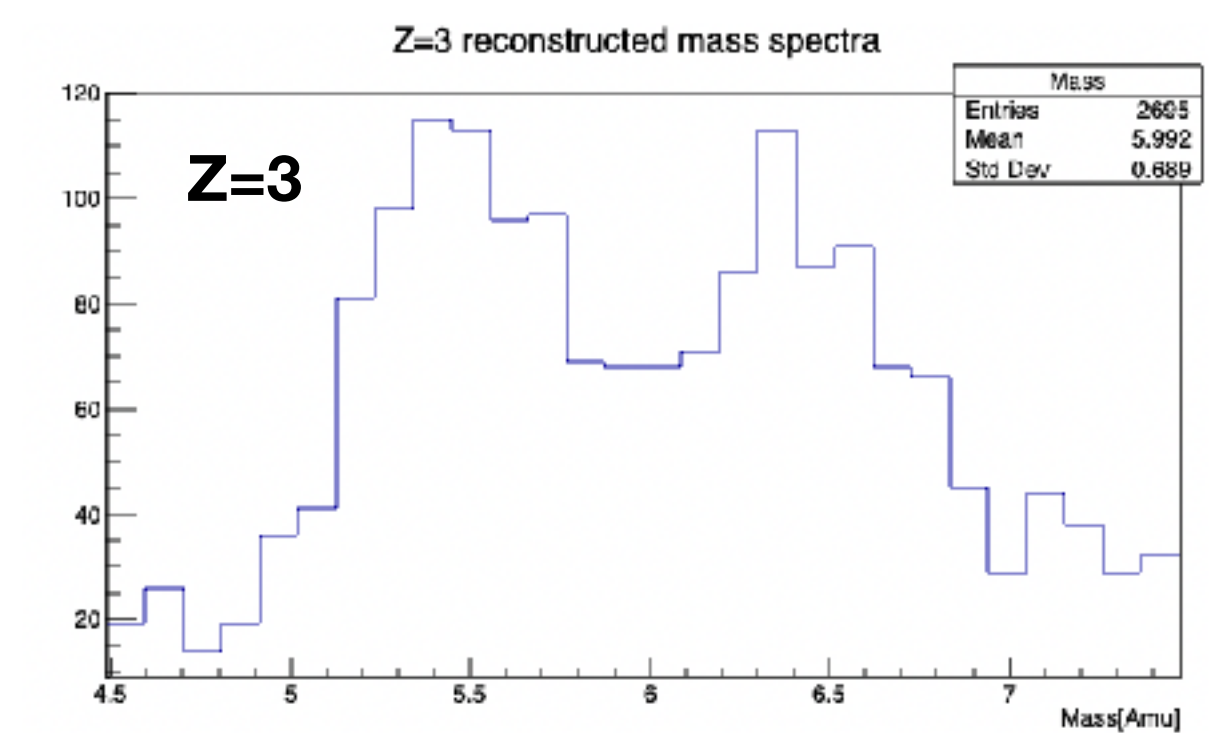
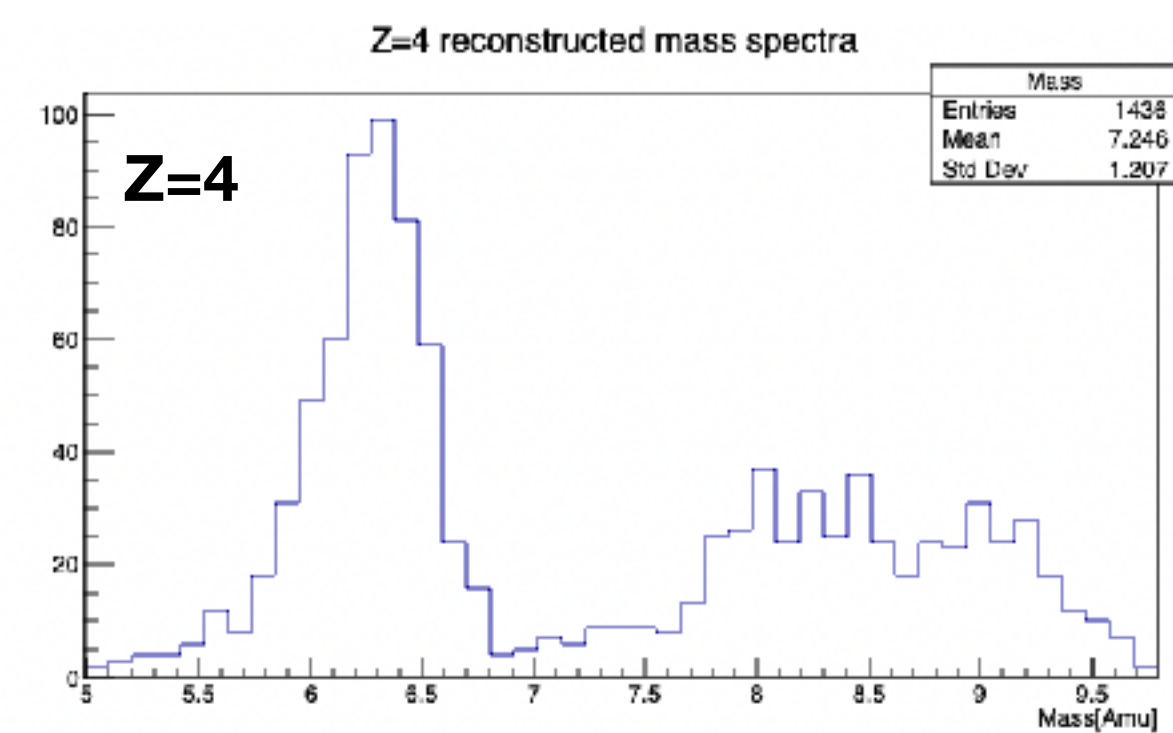
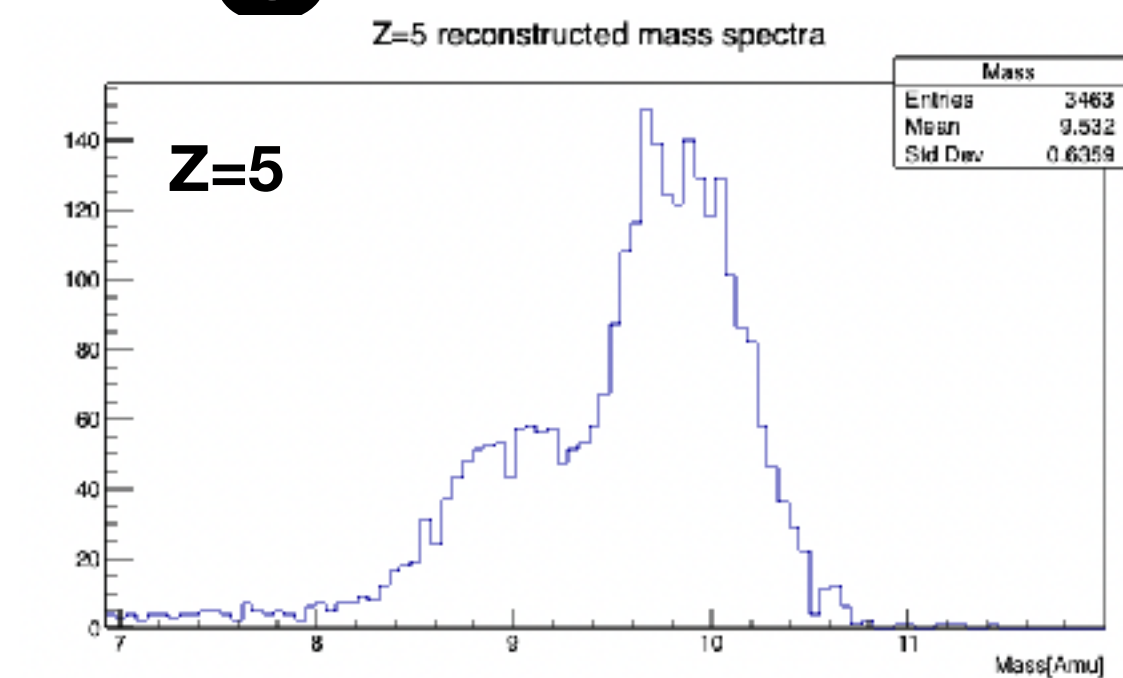
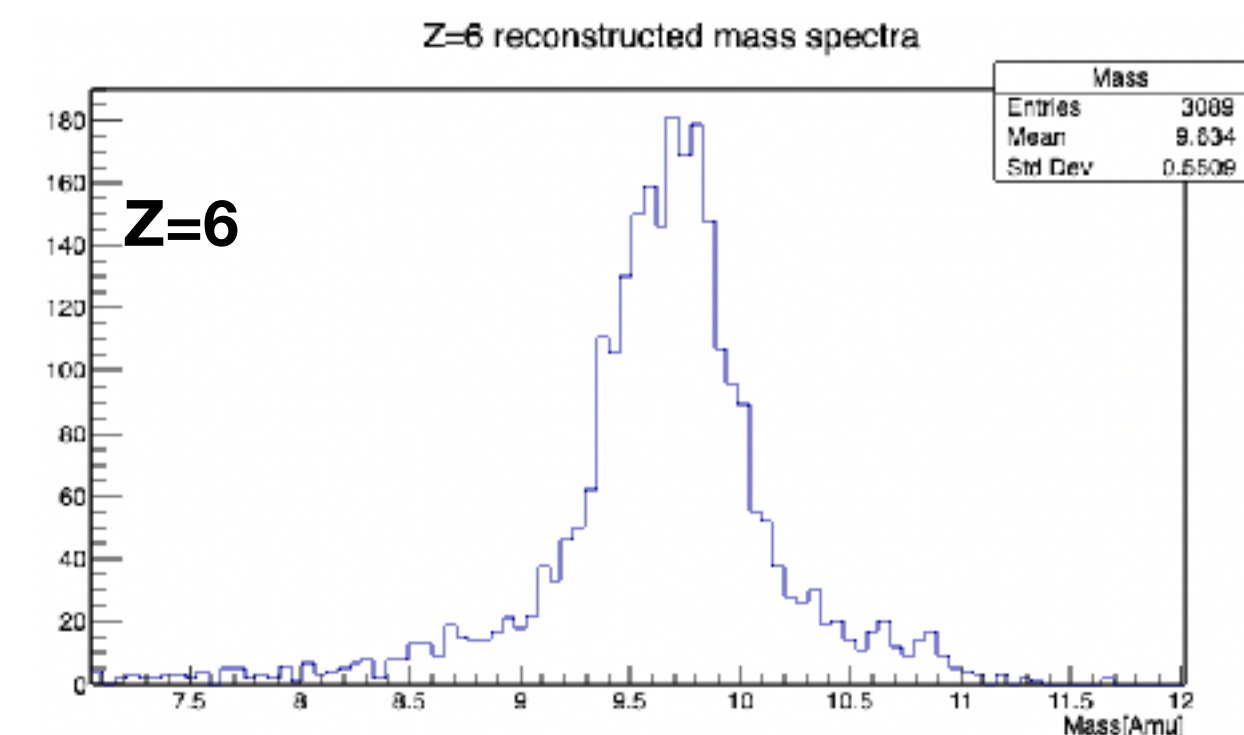


Z=3 Reconstructed mass spectra using alm method



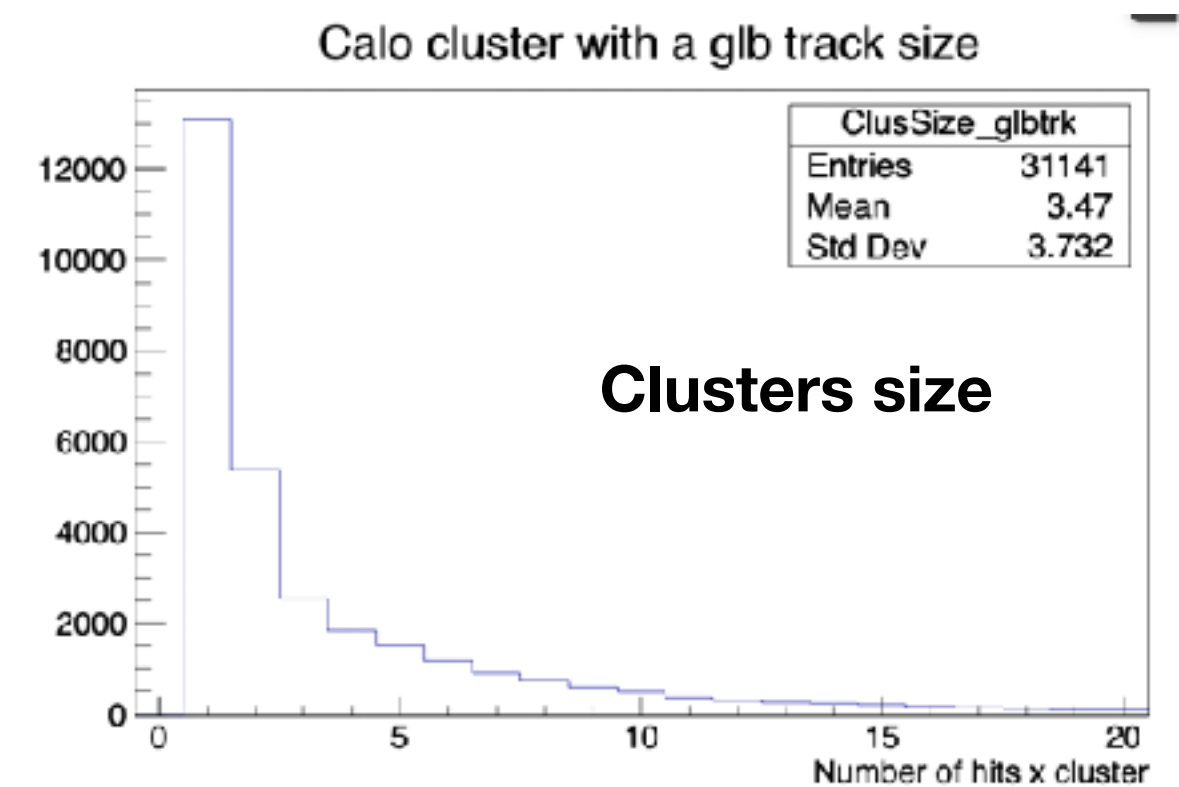
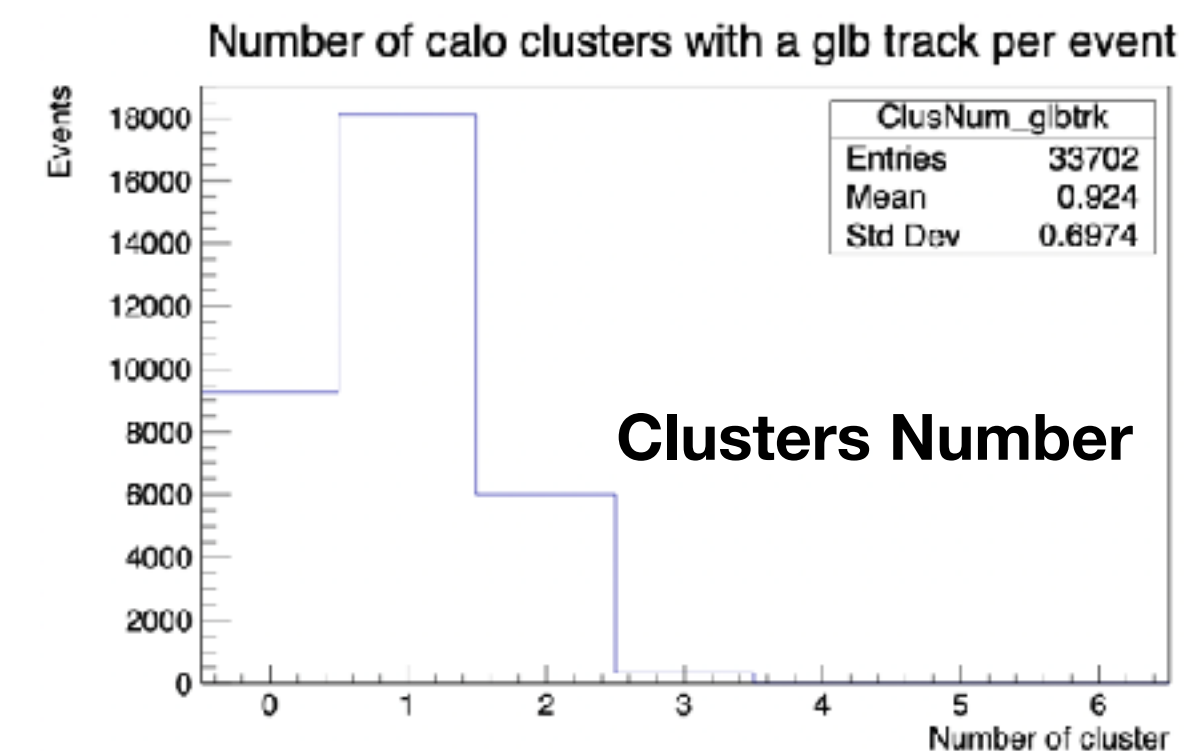
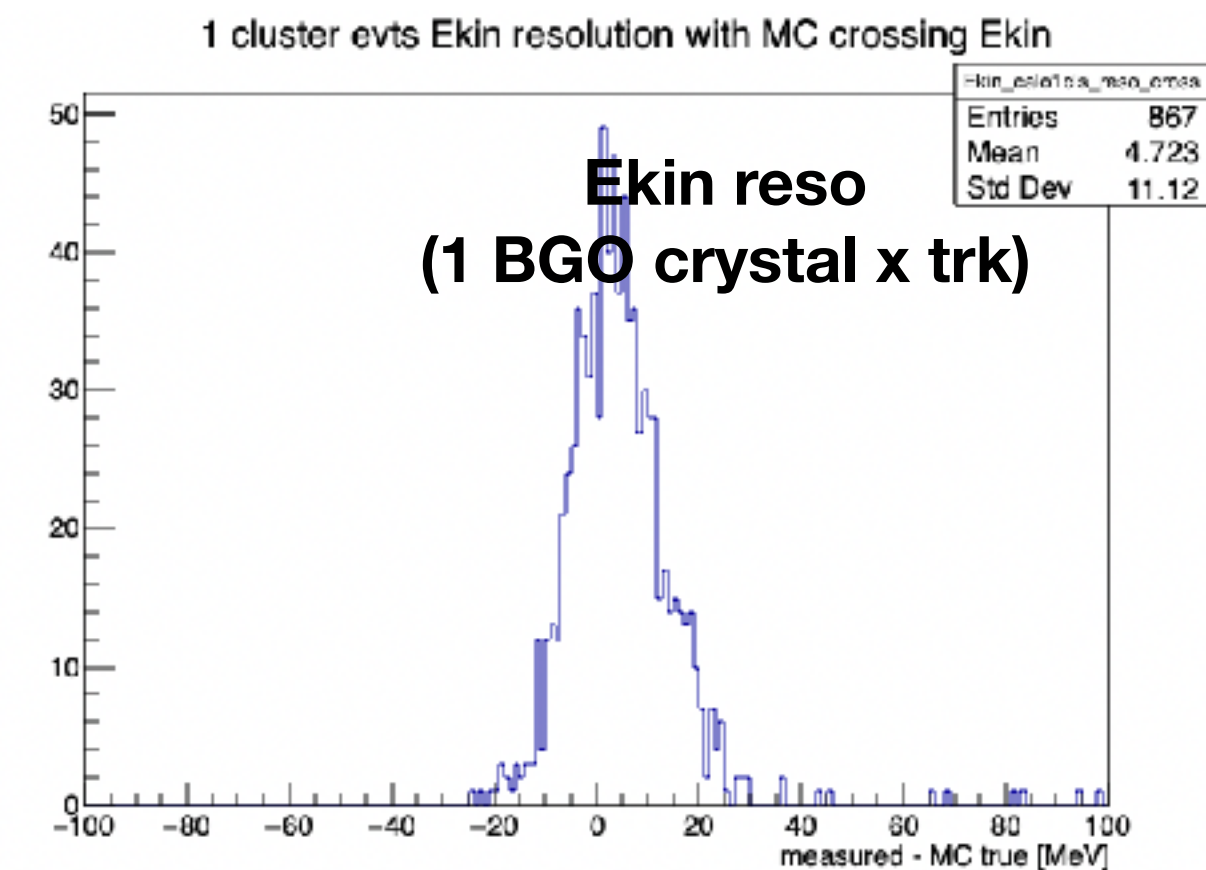
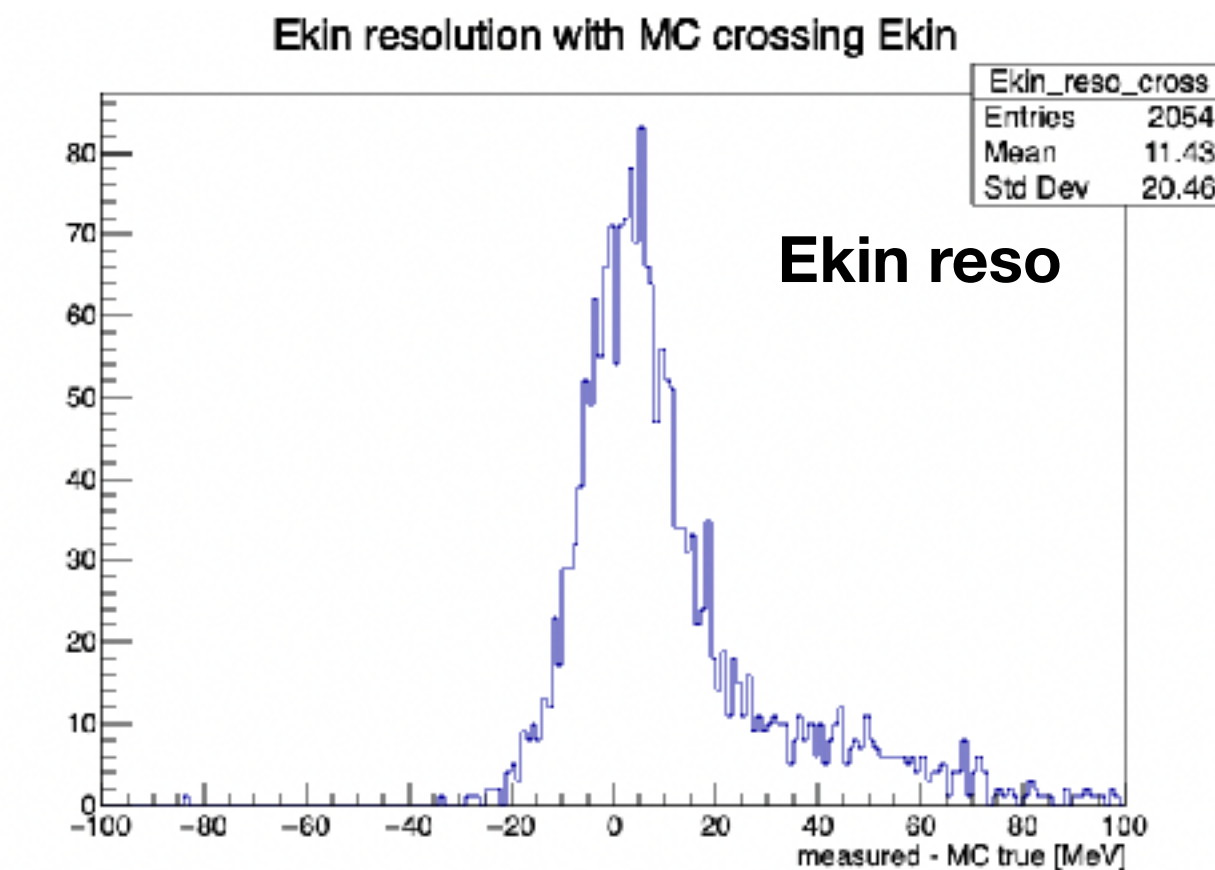
Mass spectra and combinatorial background

- The mass spectra for each Z are present
- Studies to evaluate the combinatorial background are ongoing
- Track purity seems to be very good
- The Z reconstruction precision is almost 100%, **need to add pileup and check the Z reconstruction actual performances**
- Need for statistics!



Calo studies

- At the moment the calorimeter Ekin energy measurement is higher than the MC true Ekin
- Considering only the tracks with 1 cluster, this effect is not present anymore
- The calo clustering is copy and paste from the VTX: it starts from a crystal with an over-threshold signal and it adds to the cluster all the adjacent crystals, iteratively
- Needs to verify if this clusterization algorithm is suitable for the calorimeter



To do list

- Ekin unfolding.
The theta unfolding is not necessary since the matrix is almost diagonal (with this simulation, without vtx pileup etc.)
- Efficiency definition and studies with a full MC simulation
- Full MC simulation with statistics
- Check TOE and Genfit output
- Check in the code for errors, bugs etc and improve the readability of the code

