



University of
New Hampshire



Bump hunt from 2016 data

Rafayel Paremuzyan
University of New Hampshire

HPS collaboration meeting at JLab, May 29-31, 2019

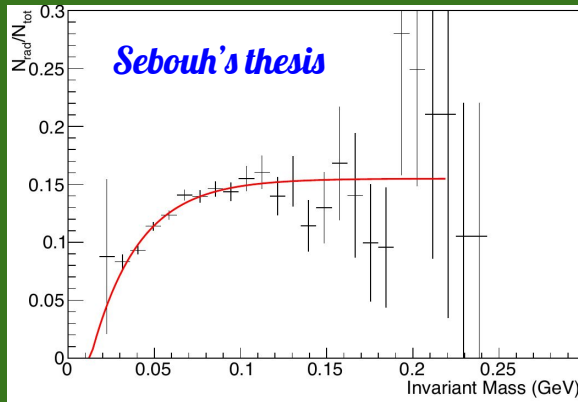
Status

Event selection

- Most of it is done in pass1 data.
- Needs to be re-obtained w/ pass4 data.
- cl. Δt , d_0 , χ^2/NDF , trk-cl time diff, ESum.

Cuts are obtained by maximizing figure of merit: e.g. $S/\text{Sqrt}(Bgr)$

Radiative fraction



Unlike to 2015, it is not flat

Bump hunting

Omar has released the code, although some features needs to be added

Mass resolution

This was parametrized by Sebouh. Codes are available.

Though the Moeller mass was obtained from Carbon data which gives smaller sigma 1.7 MeV vs 2.2 MeV

Systematic uncertainties

- frad
- Mass resolution
- fitting

Has not started yet.



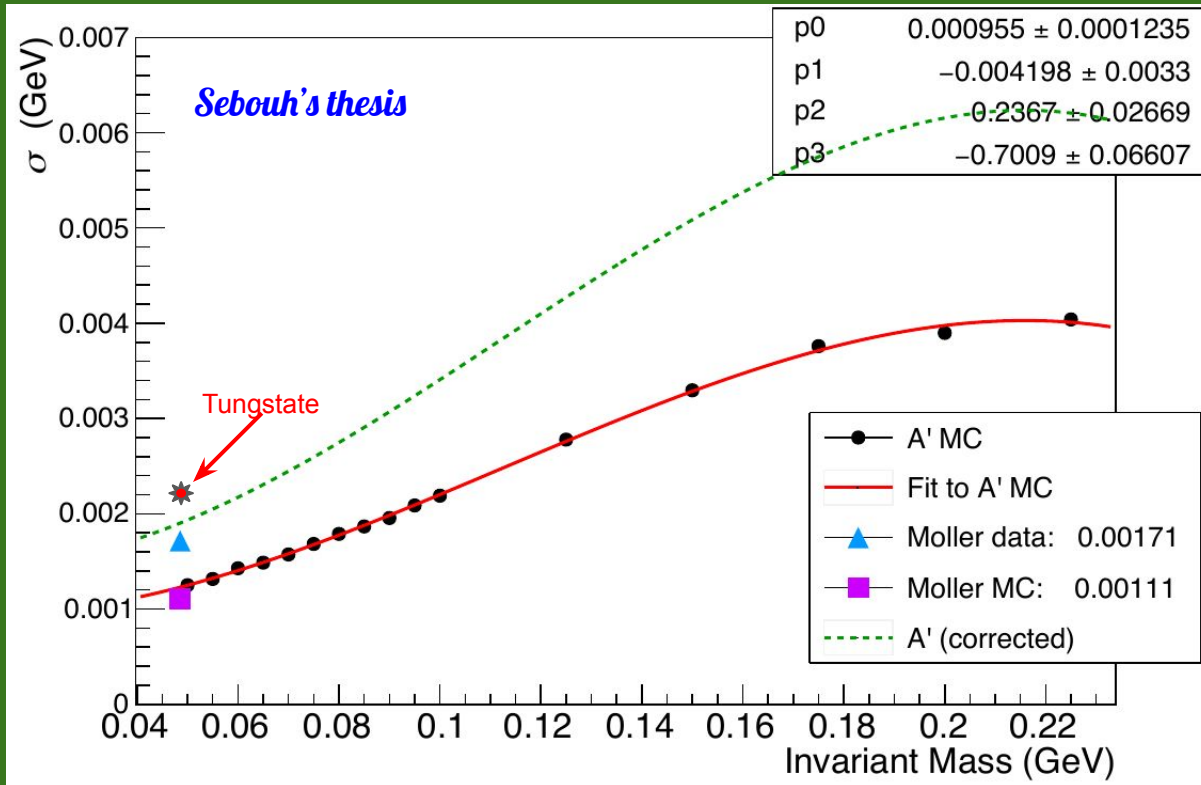
2016 Resonance Search Task list

Created by Rafayel Paremuzyan, last modified by Omar Moreno on Feb 10, 2019

Task	Subtask	Comment	Document
MC and Data agreement	<input type="checkbox"/> Check the tracking efficiency (Matt, Omar) <input type="checkbox"/> Check single hit efficiency? (Matt, Omar) Mat has progress, see his talk <input checked="" type="checkbox"/> MG5 cross section converges? (Bradley) <input type="checkbox"/> Any significant discrepancy between MC and data selection cut? (Matt)	Unlike to 2015 data, we see that the Normalized data is about 30% higher than the normalized MC	
Event Selection	<input type="checkbox"/> Optimize event selection cuts (Rafo) Done for pass1, needs to be revalidated with pass4 <input type="checkbox"/> Optimize energy/momentum sum cut (Rafo? Omar?) Done for pass1, needs to be revalidated with pass4 <input type="checkbox"/> Generate final e+e- invariant mass spectrum (Rafo)		
Determine mass resolution with the pass4 data	<input type="checkbox"/> Generate A'-beam MC at several different masses along with Moller-beam MC (Bradley). Done <input type="checkbox"/> Check data-MC agreement for Moller data (Matt? Omar?) <input type="checkbox"/> Develop a cutflow to isolate Moller peak and fit using a Crystal Ball function to extract the mass resolution (Rafo? Omar?) <input type="checkbox"/> Isolate A' invariant mass peaks and fit each using a Crystal Ball function to extract mass resolution (Rafo) <input type="checkbox"/> Fit A' mass resolution as a function of A' mass to obtain mass parametrization. (Rafo) <input type="checkbox"/> Determine mass scale correction (Kyle)	Is done for paas1, need to run codes over new MC data	
Bump Hunting	<input type="checkbox"/> Run the BumpHunter (Kyle) <input type="checkbox"/> Use Crystal-Ball instead of the Gauss for the signal shape (Omar) <input type="checkbox"/> Add CLs limit calculation <input type="checkbox"/> Import mass resolution parametrization (Kyle) <input type="checkbox"/> Optimize fitting function and window size. Requires knowledge of the mass resolution (Kyle) <input type="checkbox"/> Incorporate mass resolution and scale systematic. <input type="checkbox"/> Add "Pulls to exclusion limits" conversion in the BumpHunter?		
Systematics	<input type="checkbox"/> fRad (Matt) <input type="checkbox"/> Mass resolution (Moeller mass fit, different target positions) (Rafo) <input type="checkbox"/> fits (Kyle)		

Backup slides

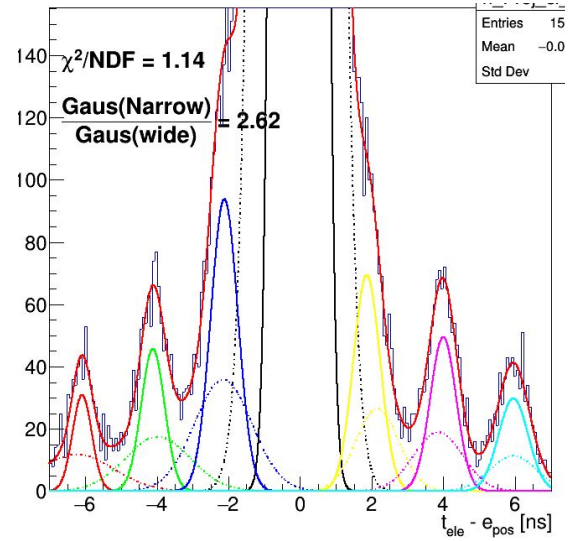
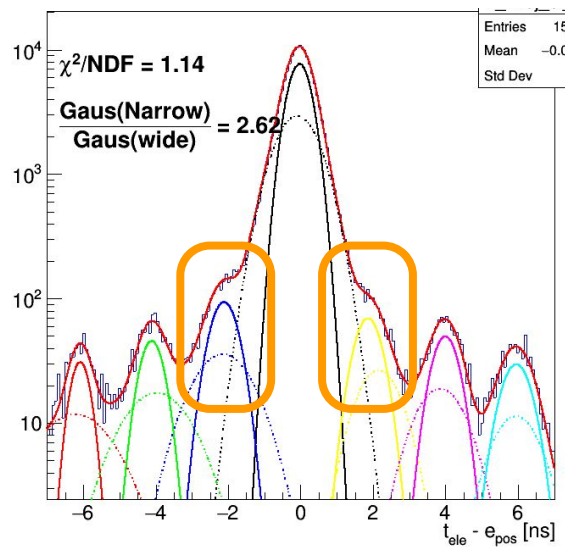
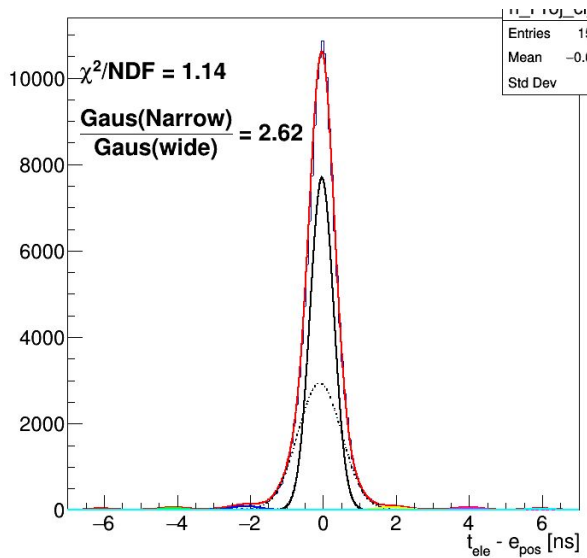
Moeller mass resolution is obtained from Carbon data.
Mass parametrization needs to be scaled wrt Tungstate mass resolution



$$\text{Peak} = A * (\text{Gaus}(x, \mu^1, \sigma^1) + d * \text{Gaus}(x, \mu^2, \sigma^2))$$

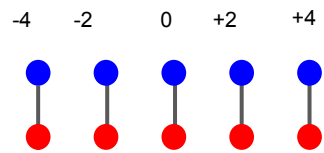
The fit function is a sum of 7 Peak functions: $F = \sum_{i=1}^7 \text{Peak}(i)$

'd' is unique for all peaks

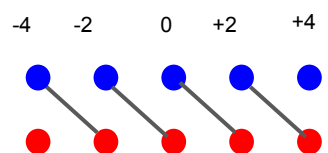


Shoulders on “-2 ns” and “+2 ns” are not symmetric, Not sure why

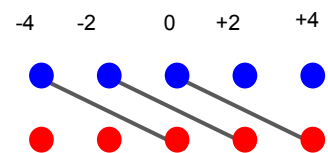
Will be checked with pass2



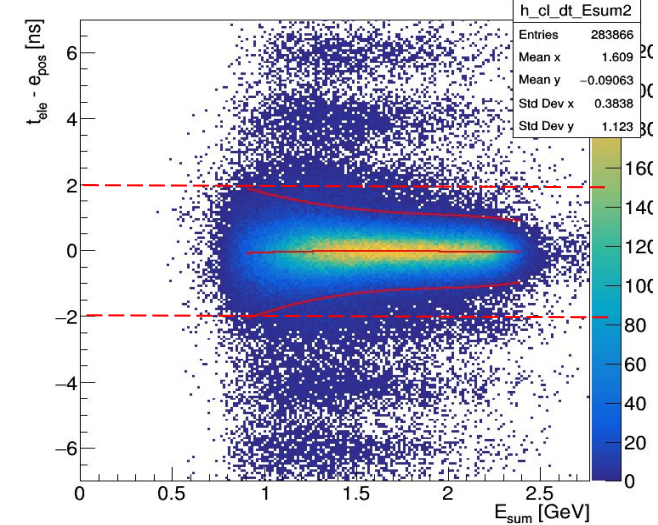
$\Delta t = 0\text{ns}$: 5 samples



$\Delta t = 2\text{ns}$: 4 samples

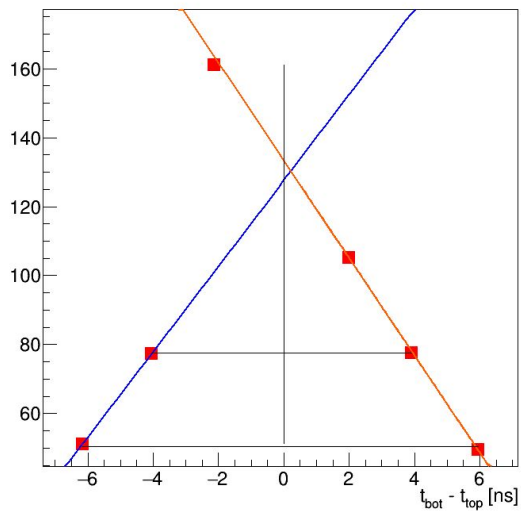
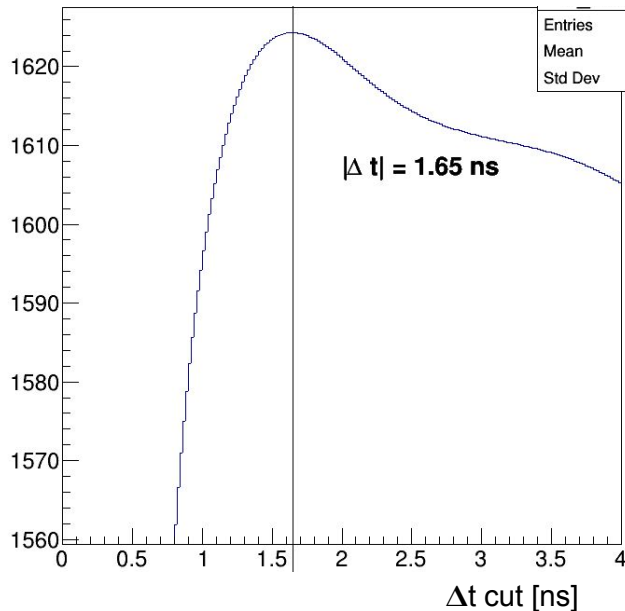


$\Delta t = 4\text{ns}$: 3 samples



There is an asymmetry here

Events at $\Delta t = -2\text{ns}$ are more than at $\Delta t = +2\text{ns}$



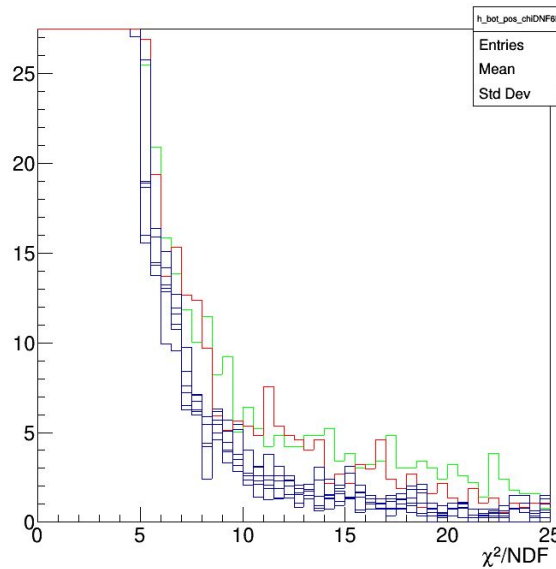
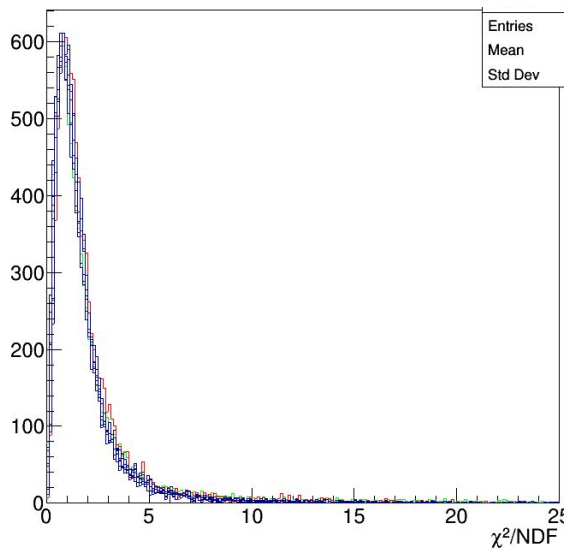
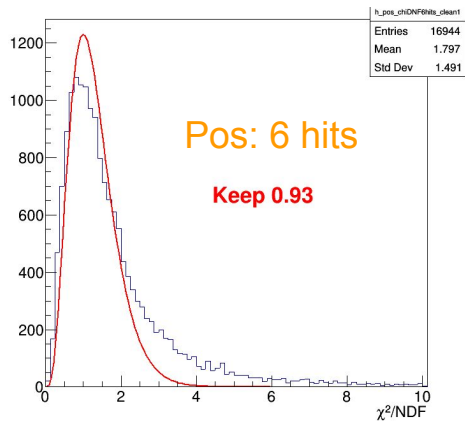
Δt cut is optimized by maximizing “Core/sqrt(total)” function as a function of cut value

Track chi2/NDF

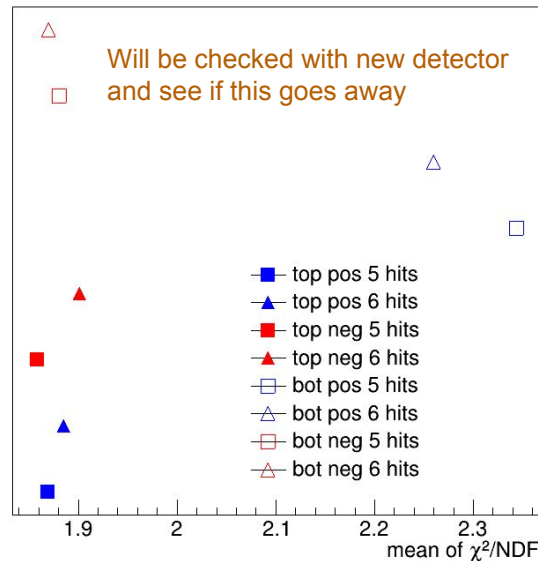
The chi2/NDF distribution is not described by the chi2/NDF PDF, but there are reasons for that: uncertainty in the geometry, not right material model etc.

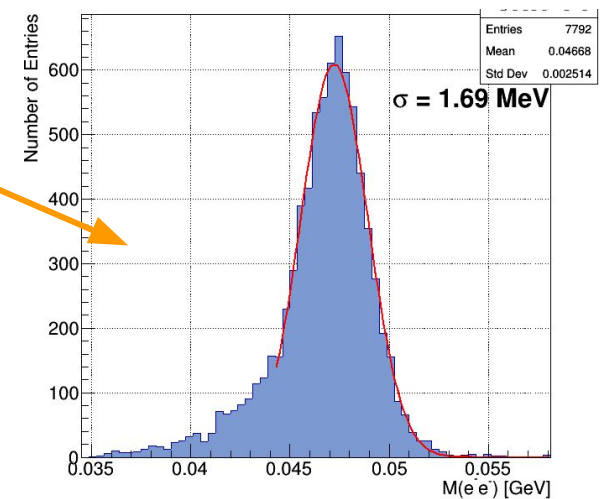
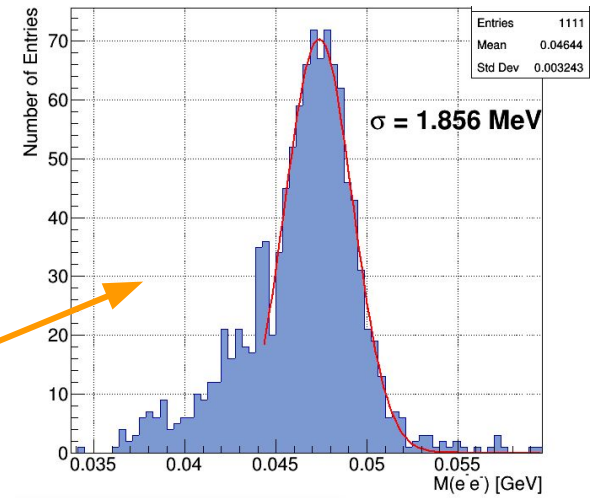
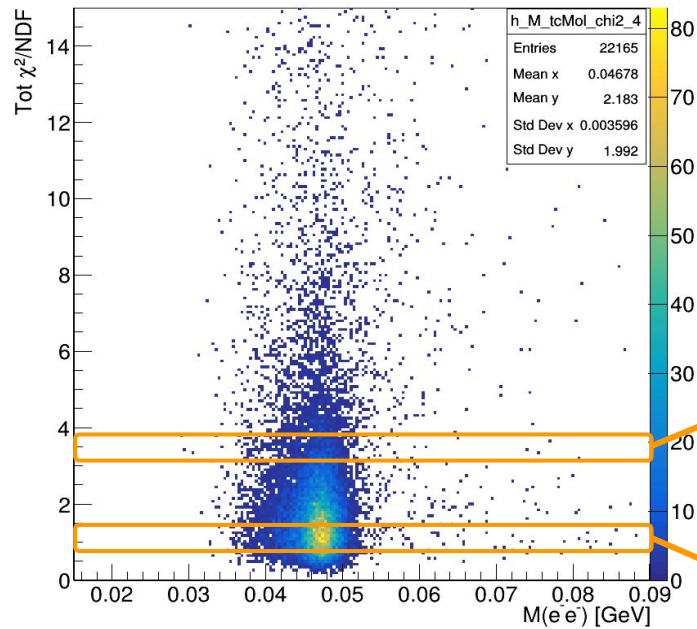
Subdividing into top/bot/neg/pos/5hits/6hits, distributions keep their shape, which is a good sign, only positrons at the bottom show slightly more tail

Select cleanest events: only 1 track in in detector half, and no extra hit is present in SVT



Mean of chi2/ndf



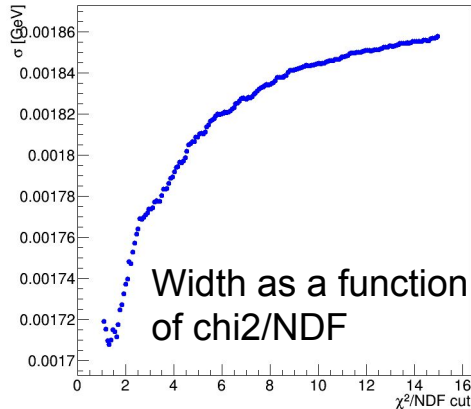
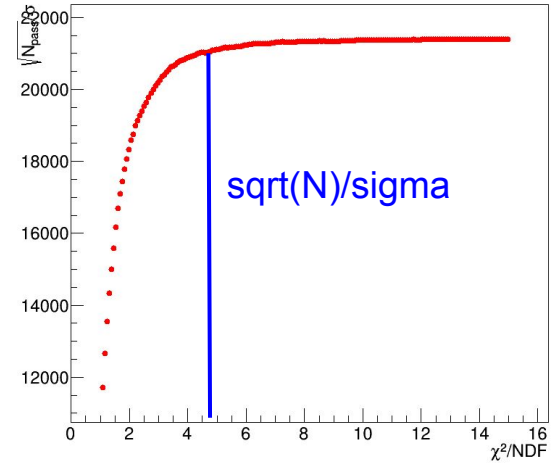
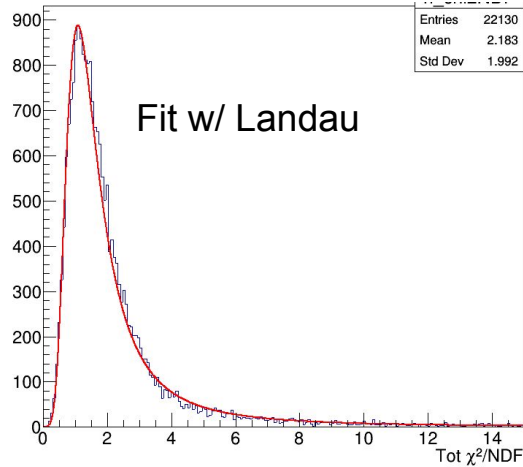
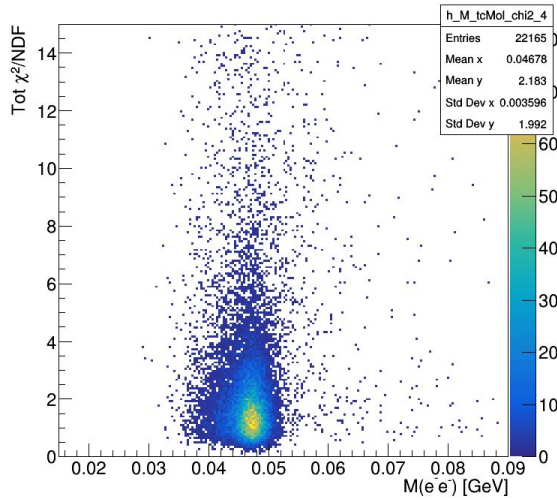


Instead of χ^2/NDF of a single electron track a total (combined) χ^2/NDF is used.

As it is expected mass depends on χ^2/NDF cut.

Optimize cut on χ^2/NDF to get the maximum signal sensitivity

Carbon data

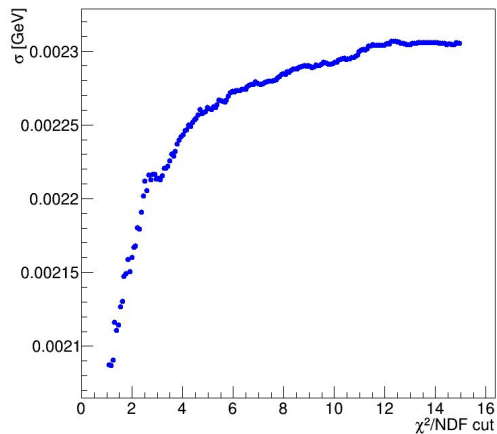
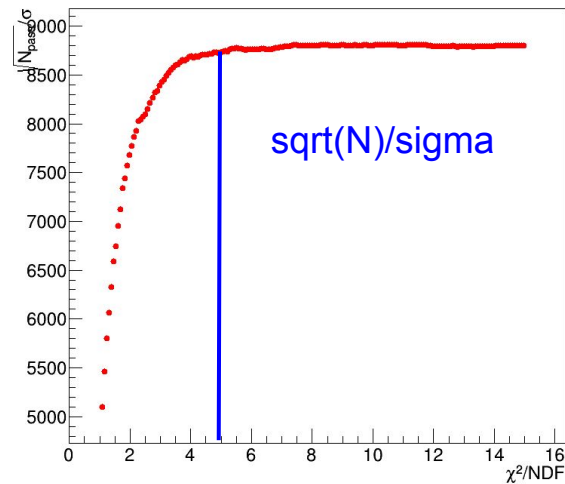
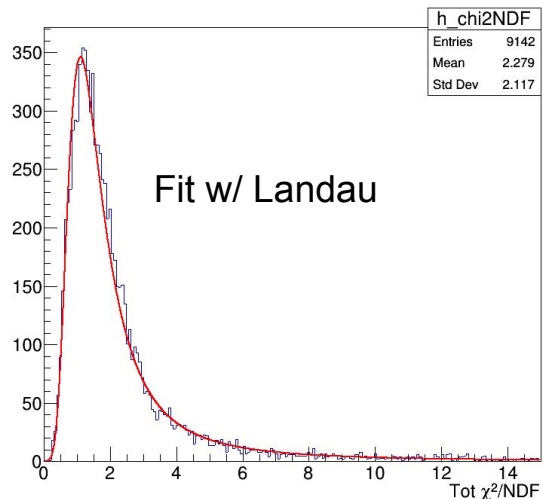
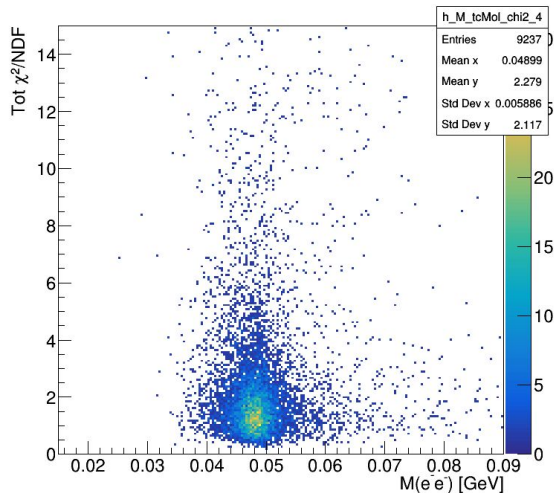


The goal: maximize \sqrt{N}/σ

No apparent maximum, but keeping it tighter without losing significant signal sensitivity is safer.

Proposal is to choose $\chi^2/\text{NDF} < 5$.

Tungsten data

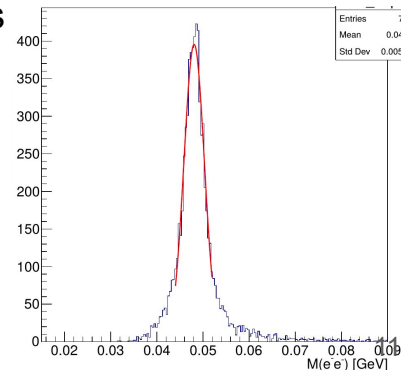


Similarities: significance doesn't converge in both cases

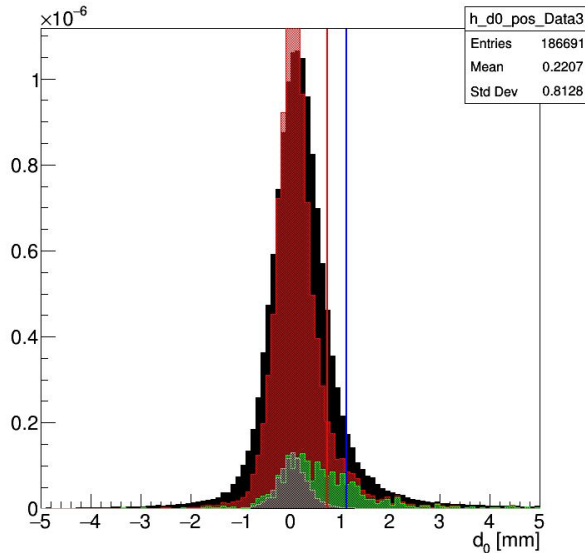
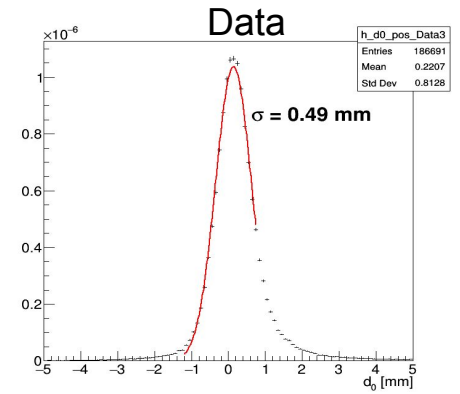
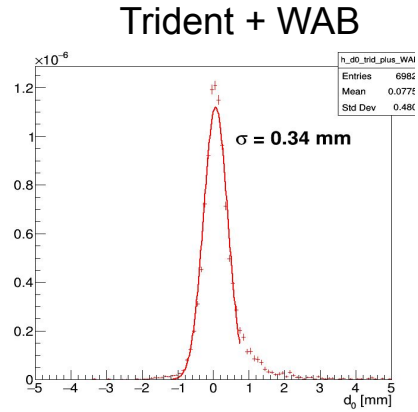
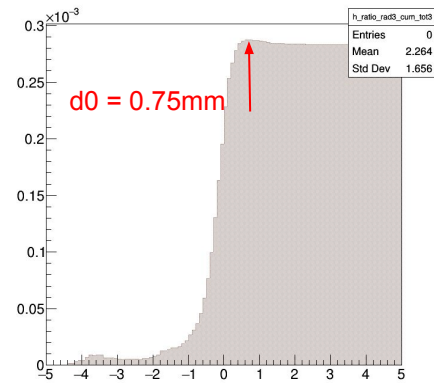
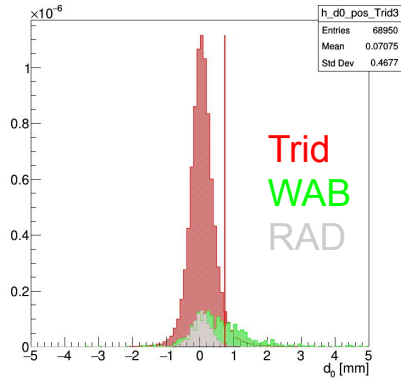
Differences:

- sigma is quite a bit larger
- The tails are on the opposite side of the peak

Need more digging into to understand sigma differences between C12 and W targets



d0 cut (suppressing cWABs)



Using only MC, the d0 cut is optimized as $N_{\text{rad}}/\sqrt{N_{\text{trid}} + N_{\text{WAB}}}$

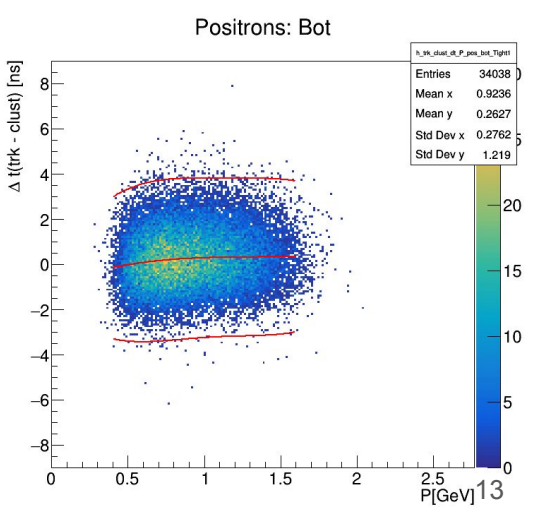
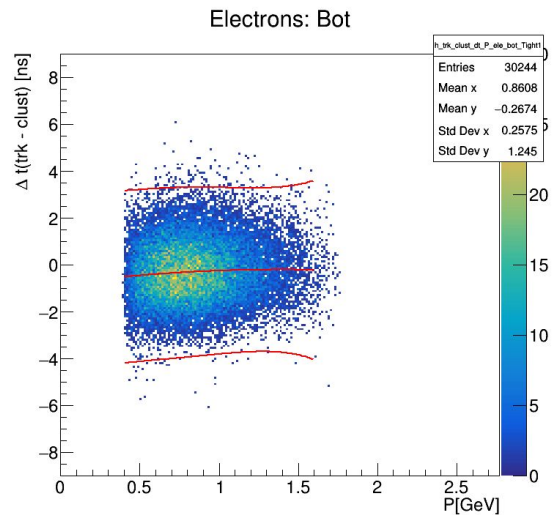
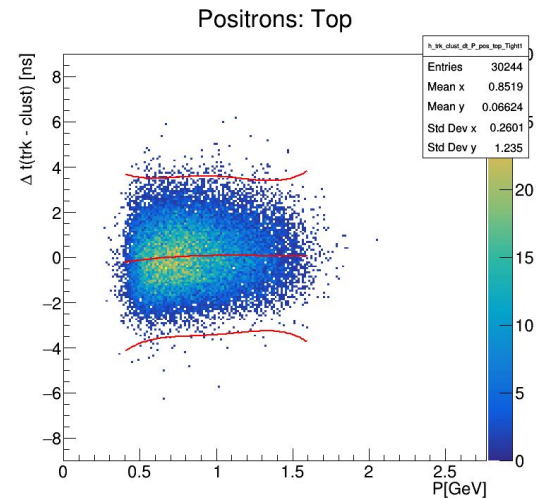
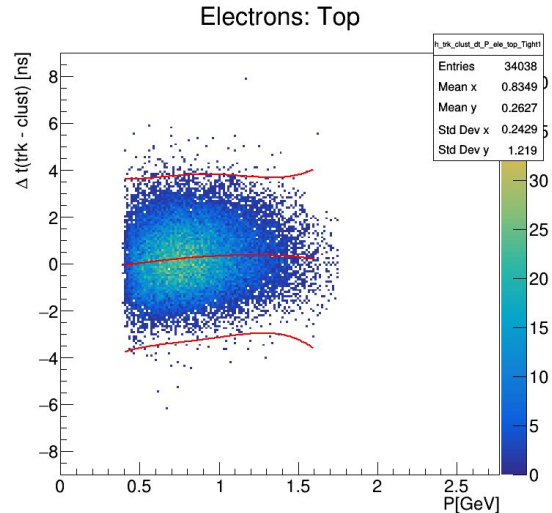
$$d0_cut(\text{data}) = \mu(\text{data}) + (d0_cut - \mu(\text{MC}))\sigma(\text{Data})/\sigma(\text{MC})$$

$$d0 < 1.18$$

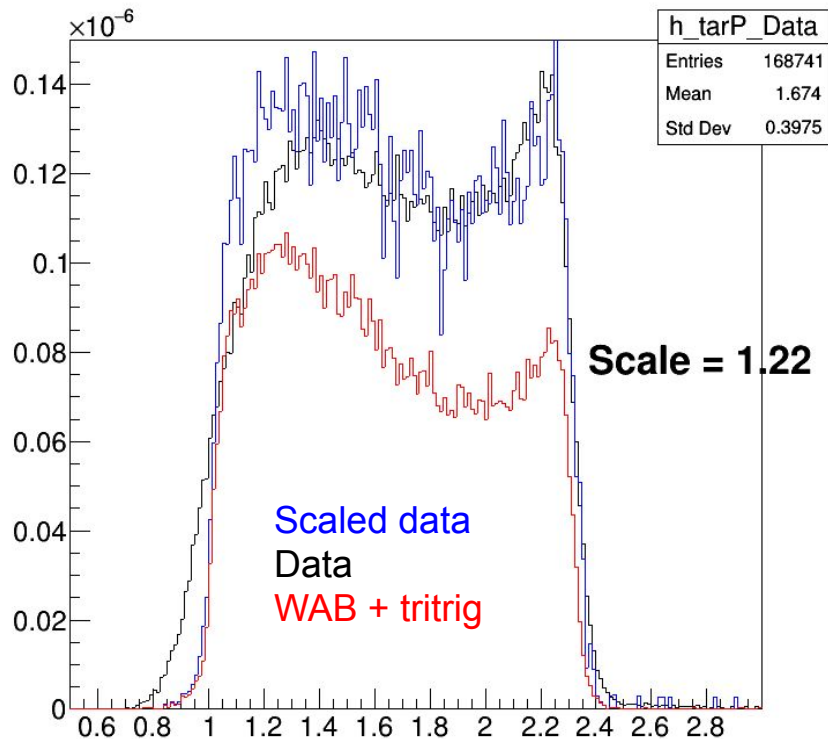
Track cluster time difference

Mean +/- 3sigma cuts can be applied
With tight cuts on other variables,
essentially no bgr is present under the
Gaussian.

As a systematic can be to put fixed cuts



Data MC comparison



Data and MC “shapes” agree to each other, however MC is lower the data by about 20%
Data is higher than the MC.

No issue is found in the data luminosity calculation.

Will be checked with the new cooking and new MC with the new detector.