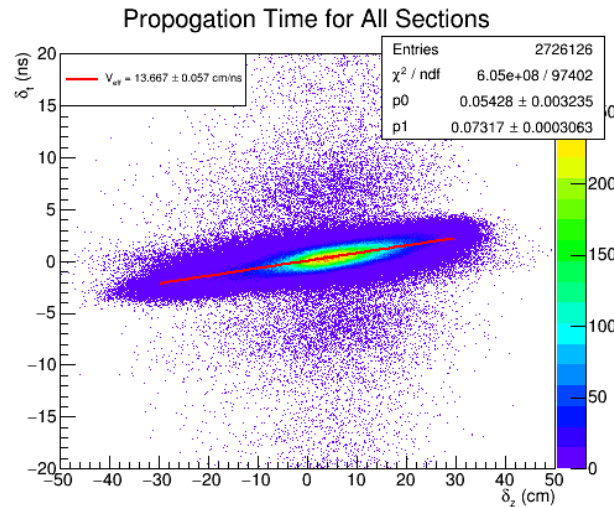
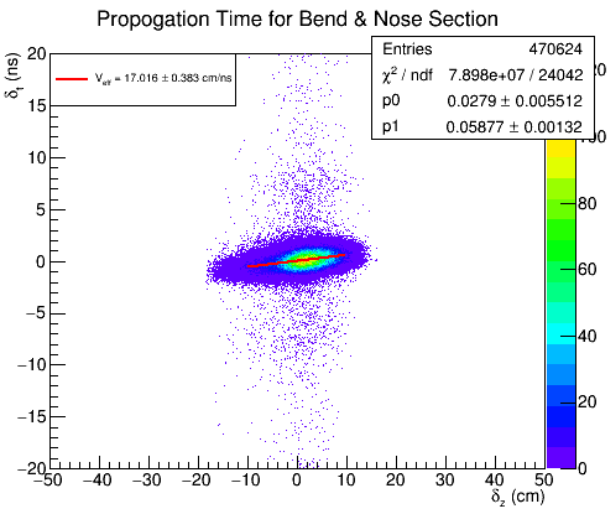
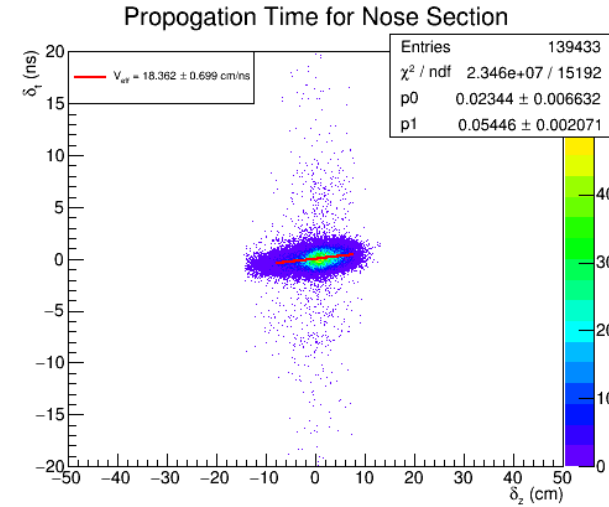
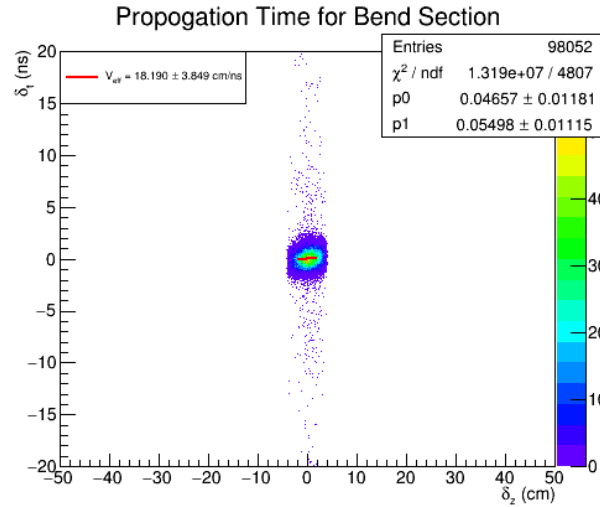
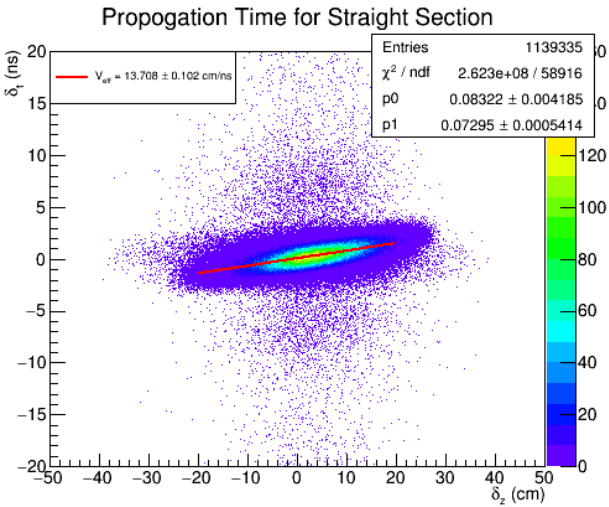


ST Propagation Time Correction Studies

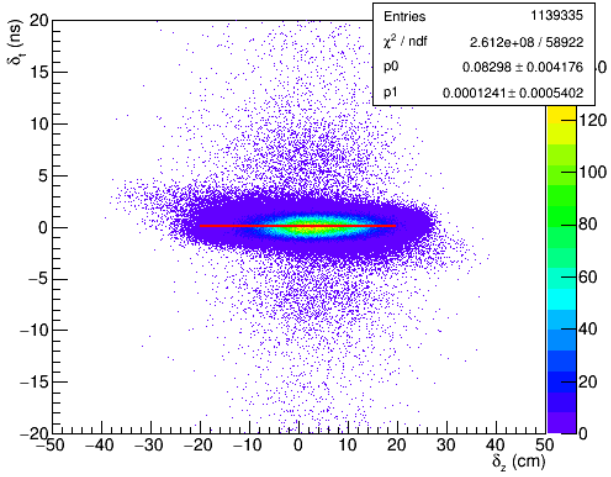
05/21/2015

ROOT Fitter Method (Straight Section)

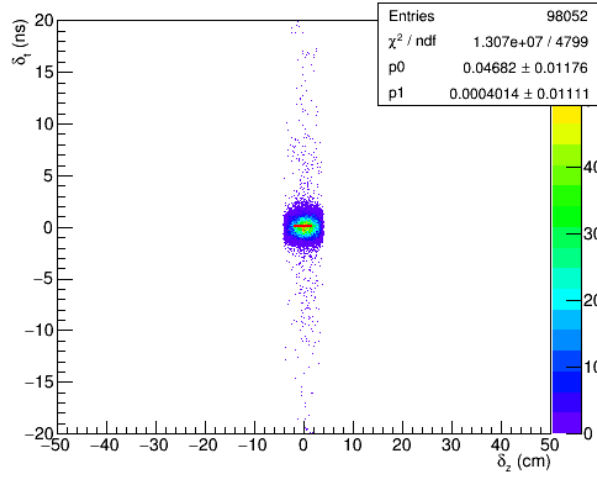


- For Straight Section:
 - $V_{\text{eff}} \approx 13.7 \text{ cm/ns}$

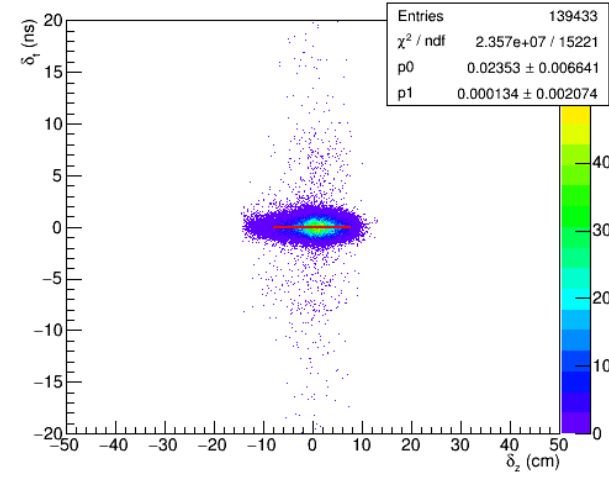
Propogation Time for Straight Section Corrected



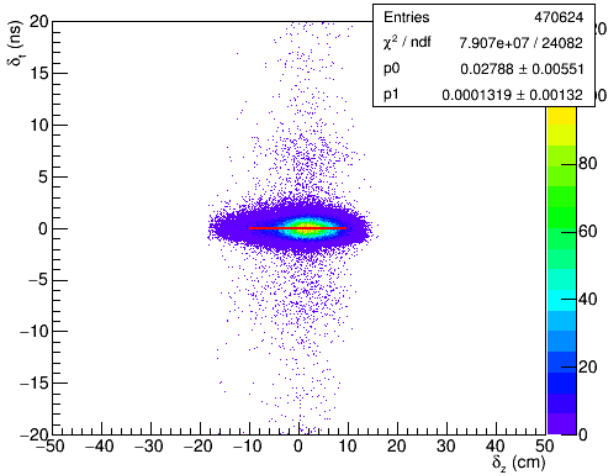
Propogation Time for Bend Section Corrected



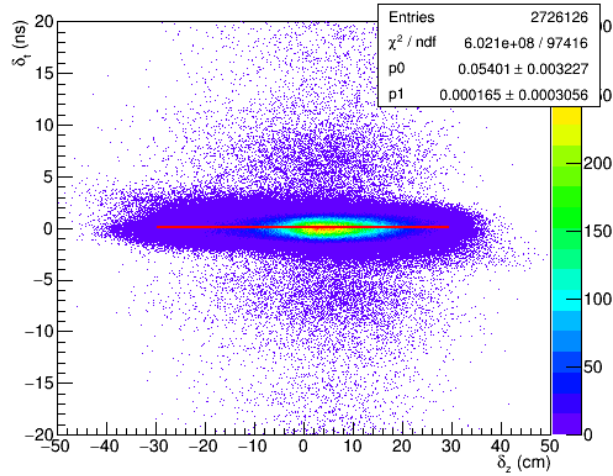
Propogation Time for Nose Section Corrected



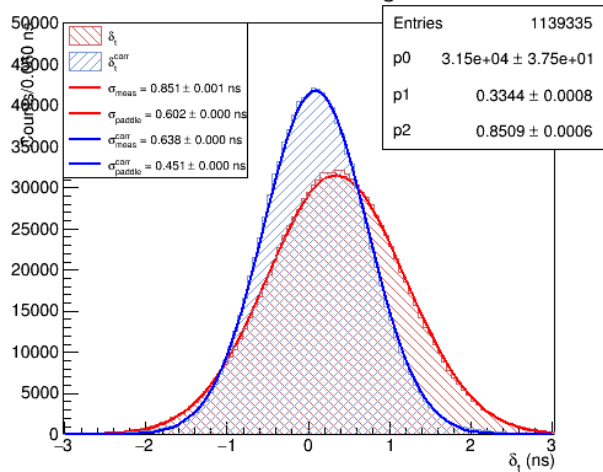
Propogation Time for Bend & Nose Section Corrected



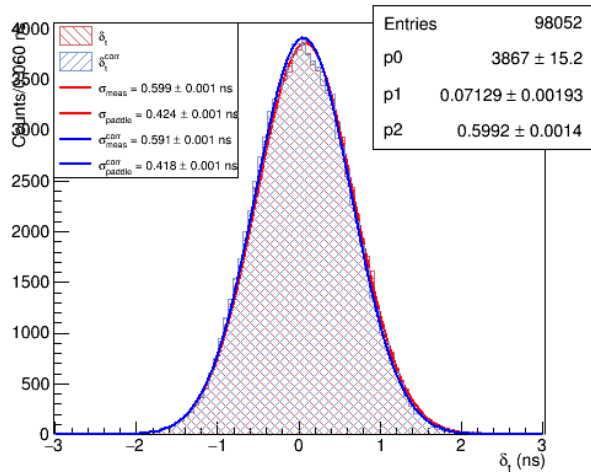
Propogation Time for All Sections Corrected



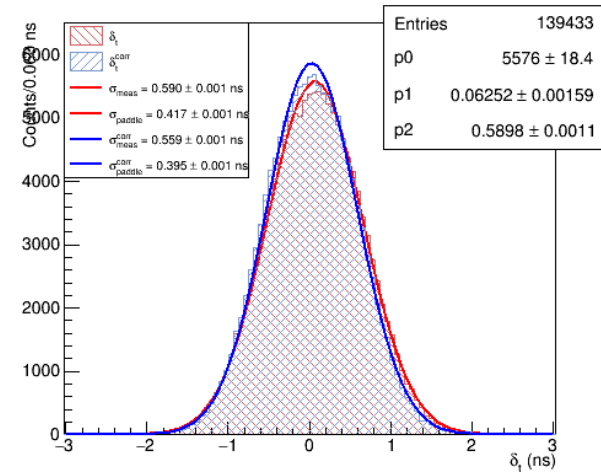
Time Resolution for Straight Section



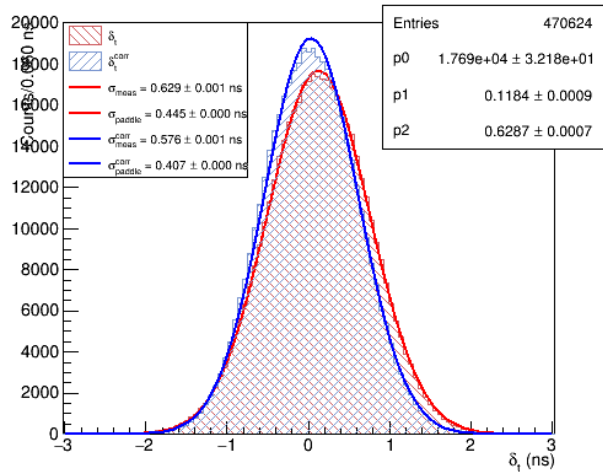
Time Resolution for Bend Section



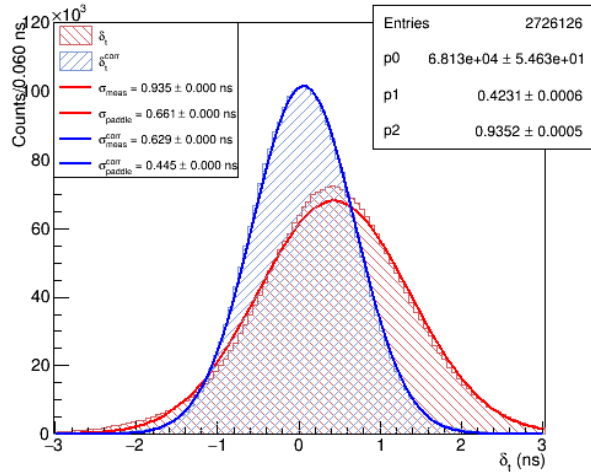
Time Resolution for Nose Section



Time Resolution for Bend & Nose Section

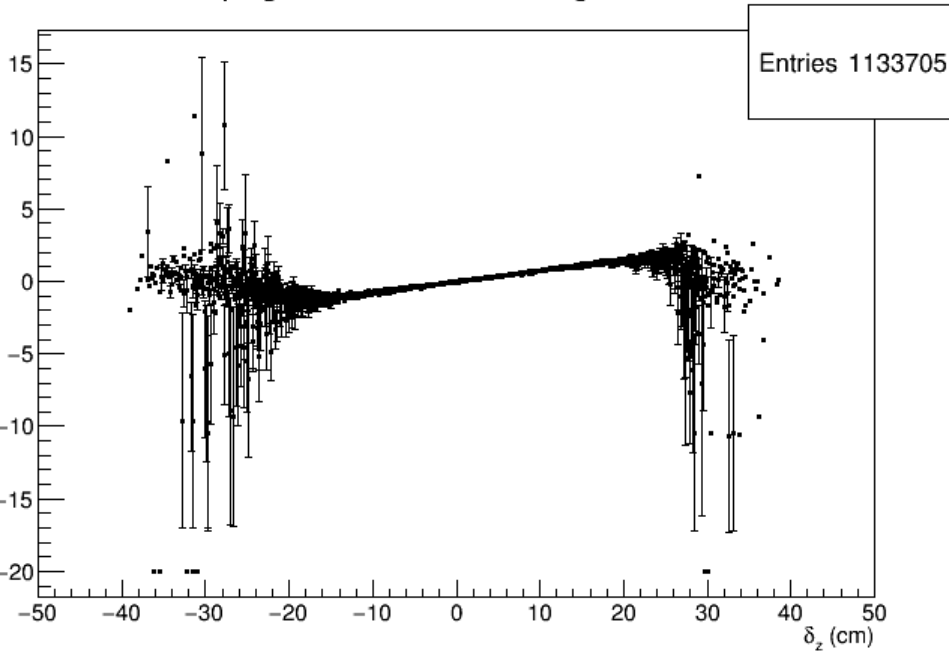


Time Resolution for All Sections

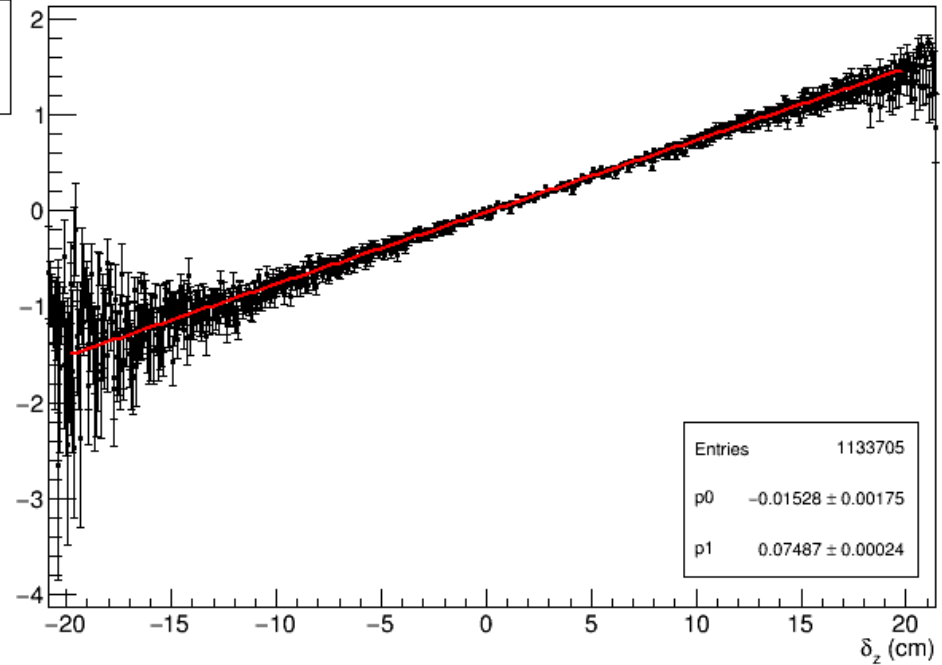


ProfileX Method (Straight Section)

Propogation Time for Straight Section



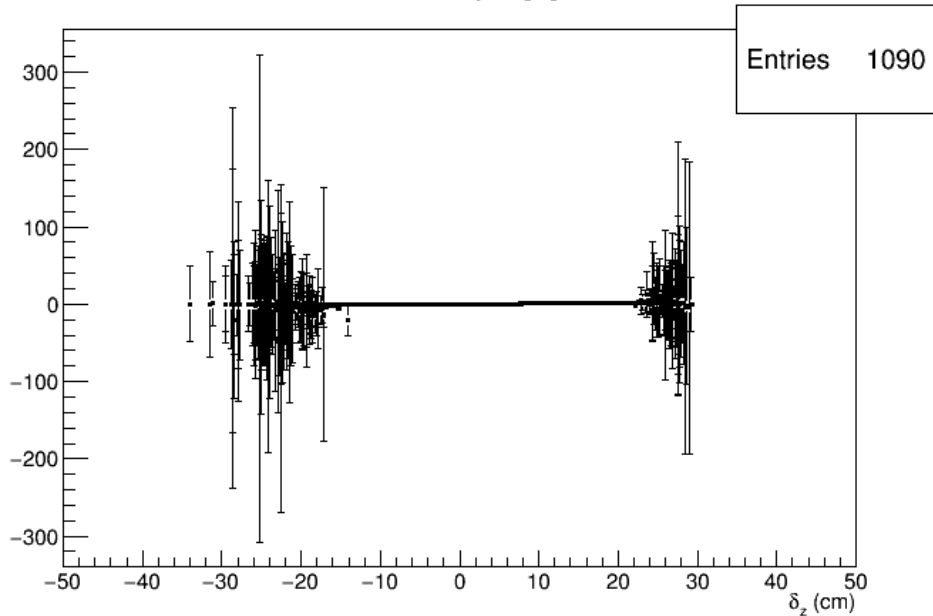
Propogation Time for Straight Section



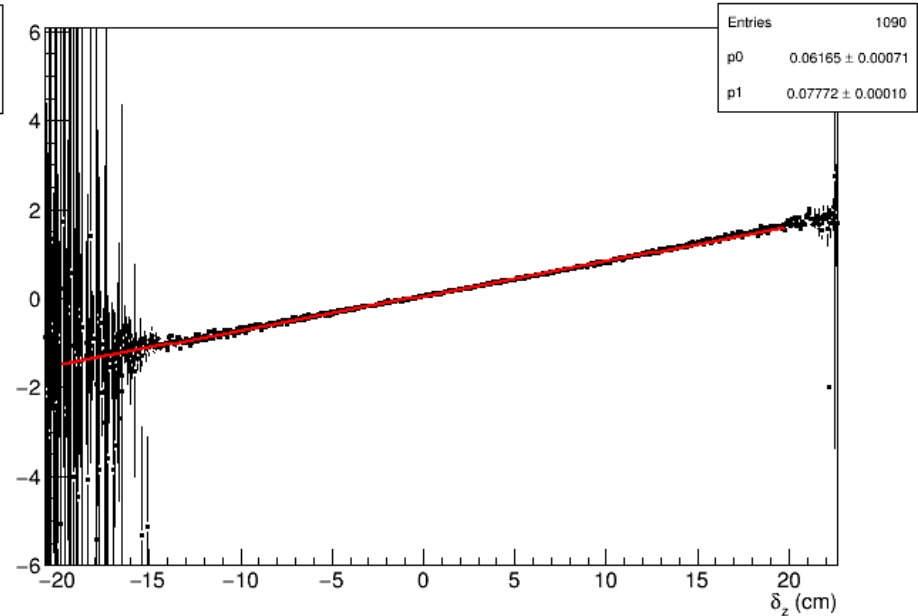
- Calculates the mean value of δt and its error for each bin ($500 \mu\text{m}$) along δz
 - The error calculated is the standard error on the mean
- $V_{\text{eff}} \approx 13.35 \text{ cm/ns}$

FitSlicesY (Straight Section)

Fitted value of par[1]=Mean



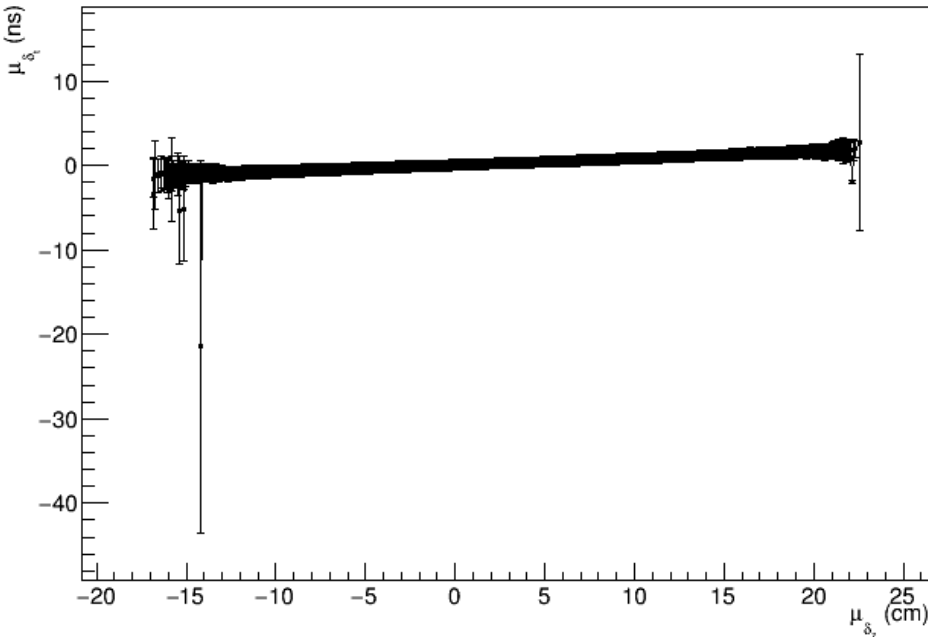
Fitted value of par[1]=Mean



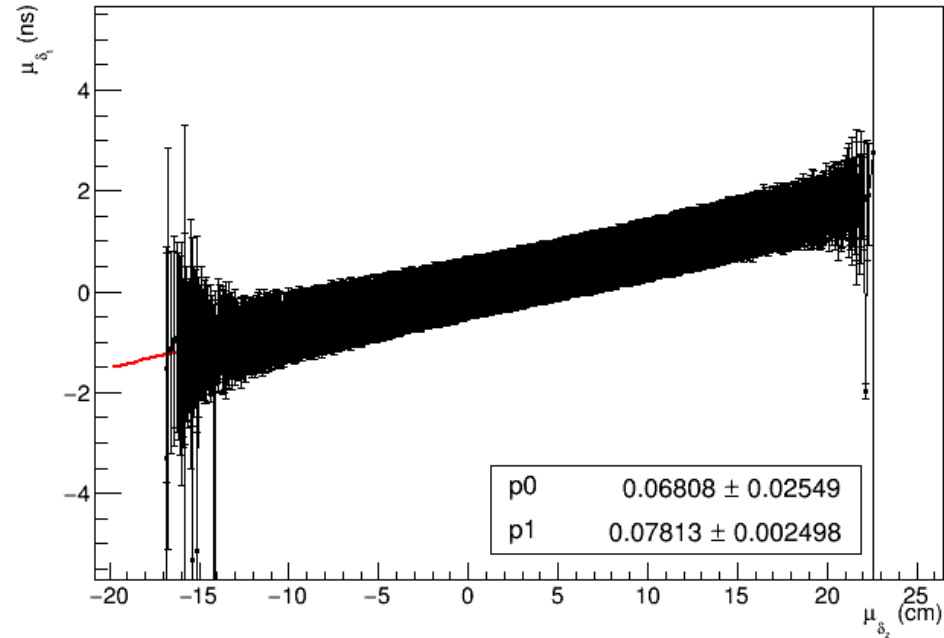
- Project slices along δt and fits them with a Gaussian for each bin ($500 \mu\text{m}$) along δz
 - The error calculated is the error on the mean as calculated by the fit
- $V_{\text{eff}} \approx 12.86 \text{ cm/ns}$

Custom Method (Straight Section)

Straight Section



Straight Section



- Project slices along δt and fit them with a Gaussian for each bin (500 μm) along δz
 - Require that the number of entries for each projection be greater than some predefined value (100 entries)
 - Other cuts are obviously possible
 - The error calculated is σ returned by the fit
- $V_{\text{eff}} \approx 12.79 \text{ cm/ns}$

Which Method Is Best?

- ROOT Fitter (Minuit & Fumili) is a black box
 - Outliers have same weight as “good” data points
- ProfileX underestimates the error
- FitSlicesY underestimates the error
- Custom method provides much more control
 - Cuts on σ , χ^2 / ndf , number of entries etc.
- What about binning?
 - Current binning scheme:
 - 500 μm in Z, 60 ps in TDC time
 - Analysis is extremely sensitive to binning