

On the way to reach  
high resolution spectra  
with FINUDA

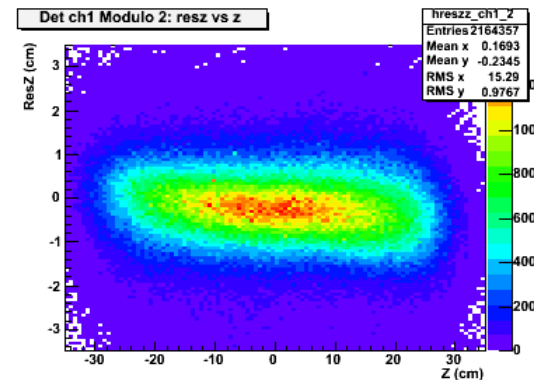
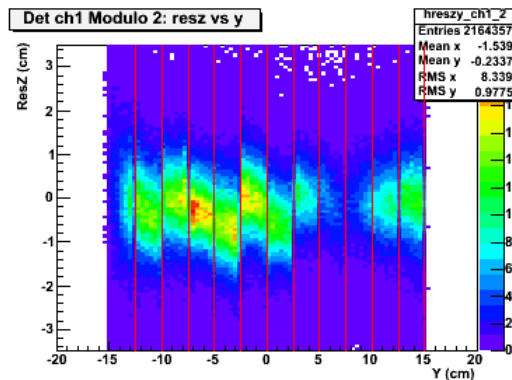
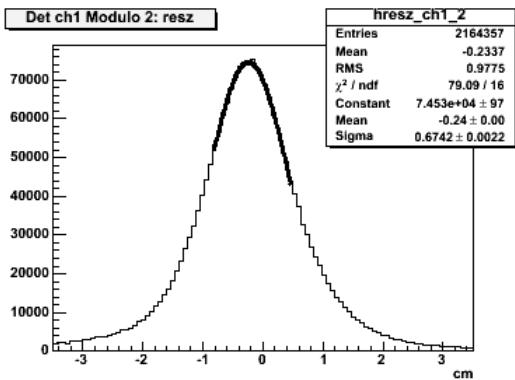
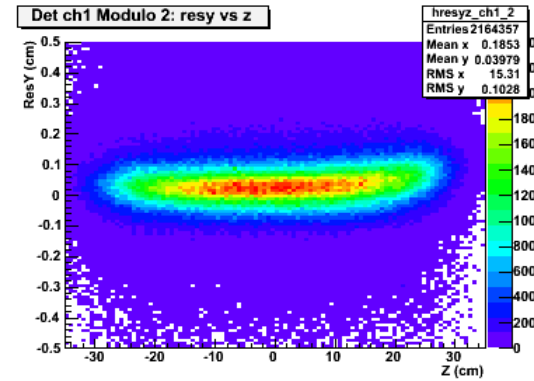
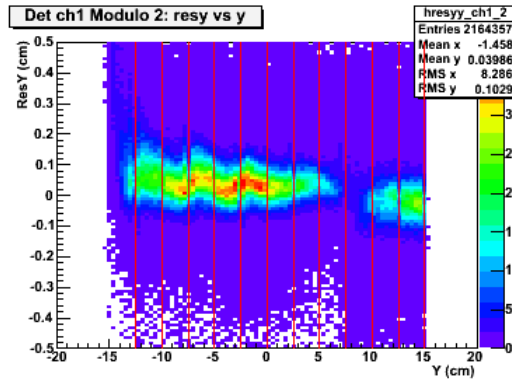
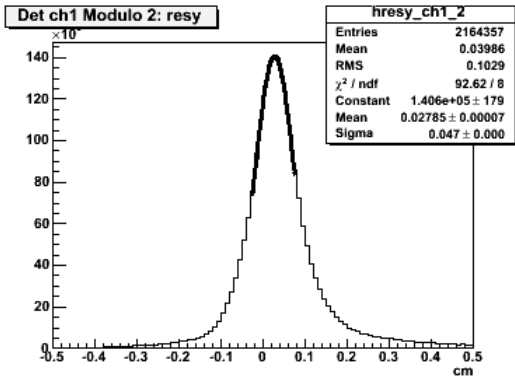
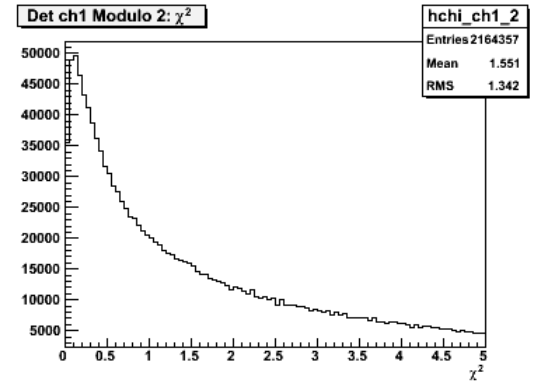
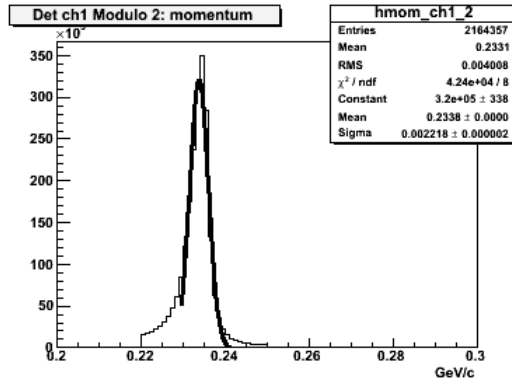
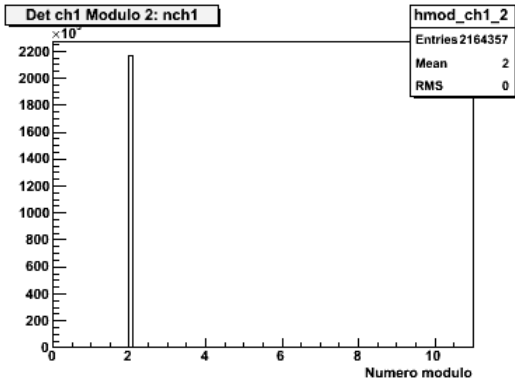
S. Piano

Finuda TRIESTE

# Calibration and Alignment

- VDET aligned (looking at V60504)
- DCH1 and DCH2:
  - Different behavior of negative and positive tracks
  - Residual distributions depend on anodic wire
    - $Z$  few centimeters
    - $\Phi$  few millimeters
- STRAW:
  - Waiting for reliable 2003-2004 Cosmic Rays Calibration Table on FINUDA Data Base (the basis of the global alignment...)

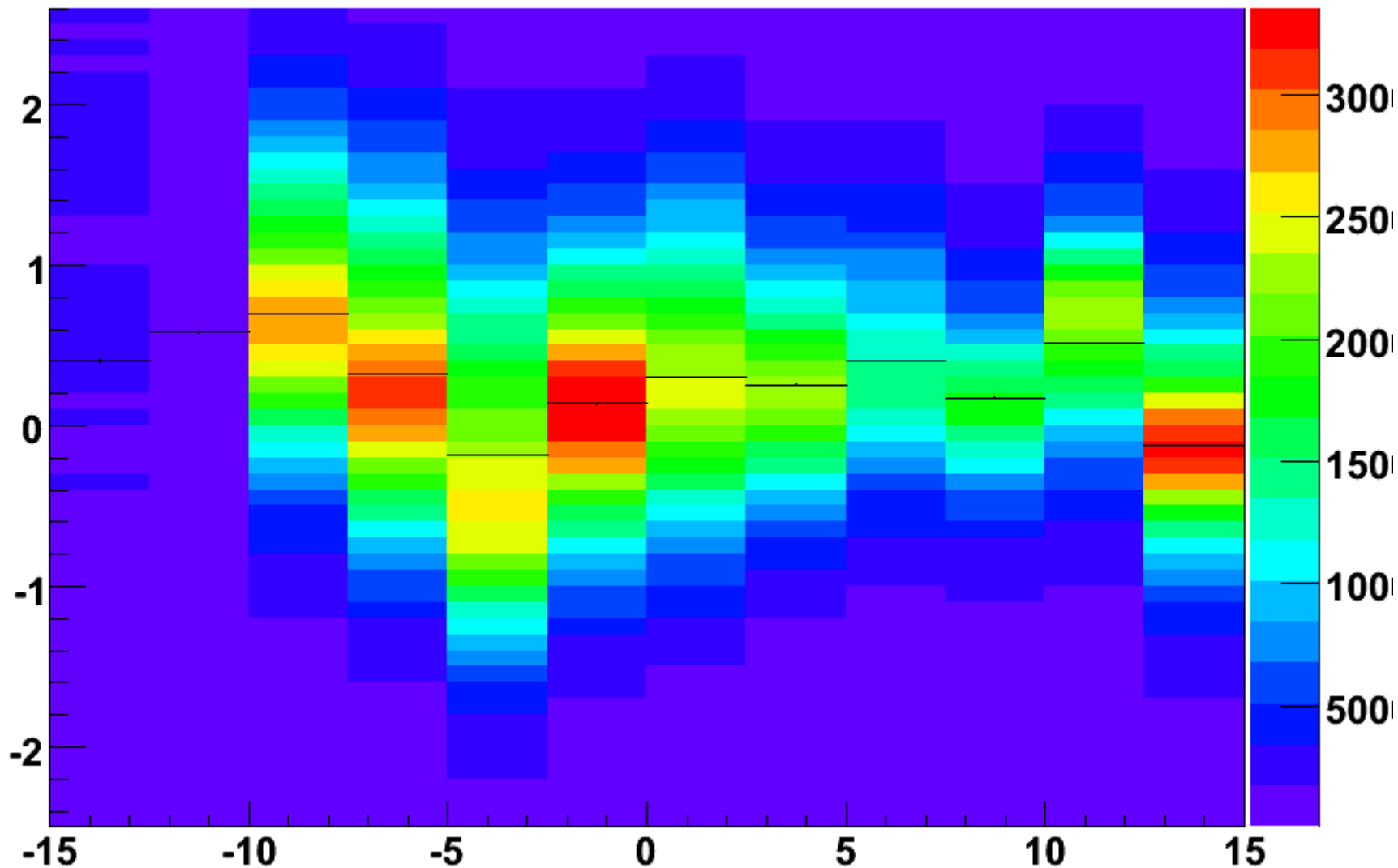
# Drift Chamber Residual Distributions (Germano)



# Z Side Residuals vs Anodic Wire

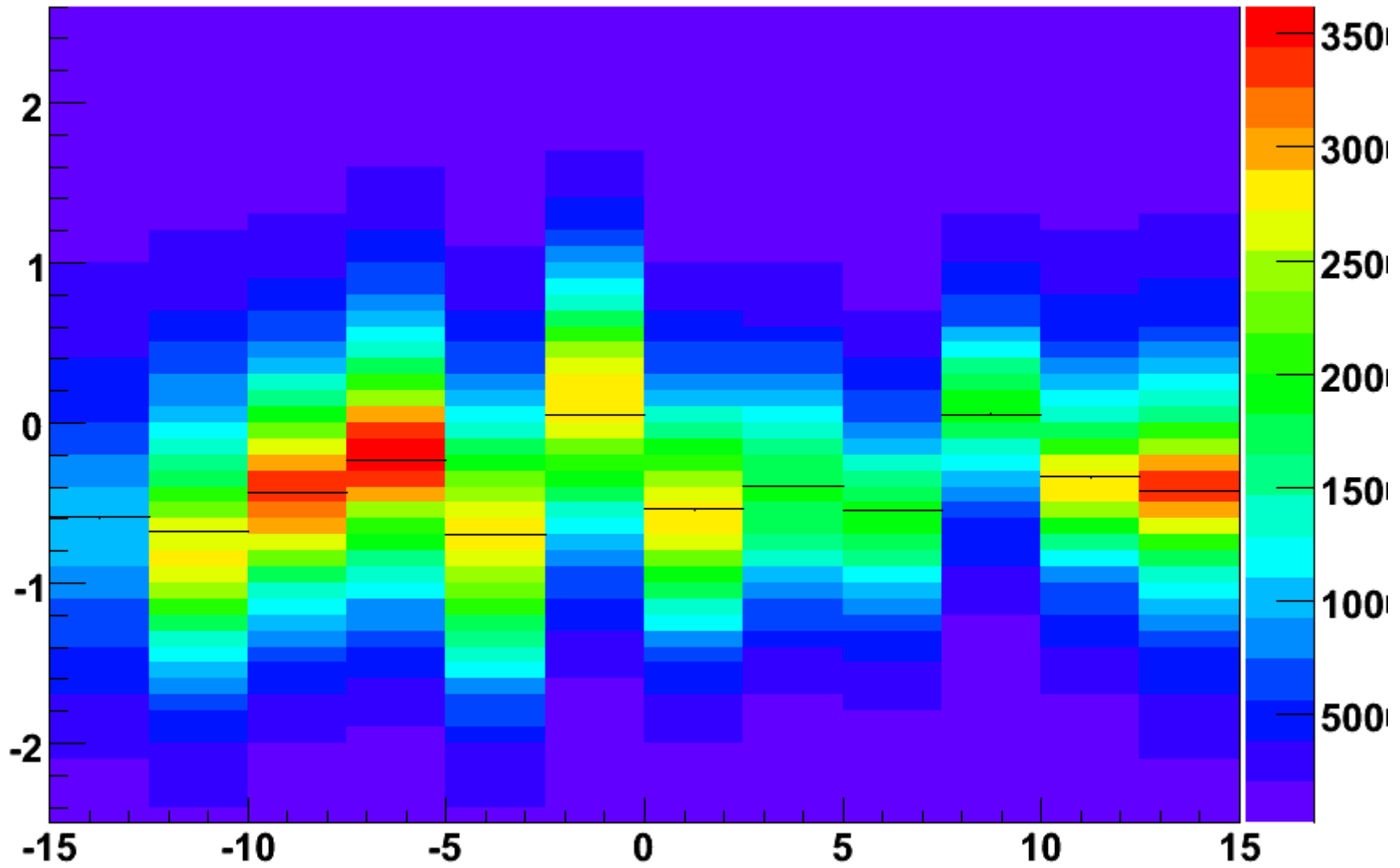
reszvsy\_ch1\_2

Entries 4437478



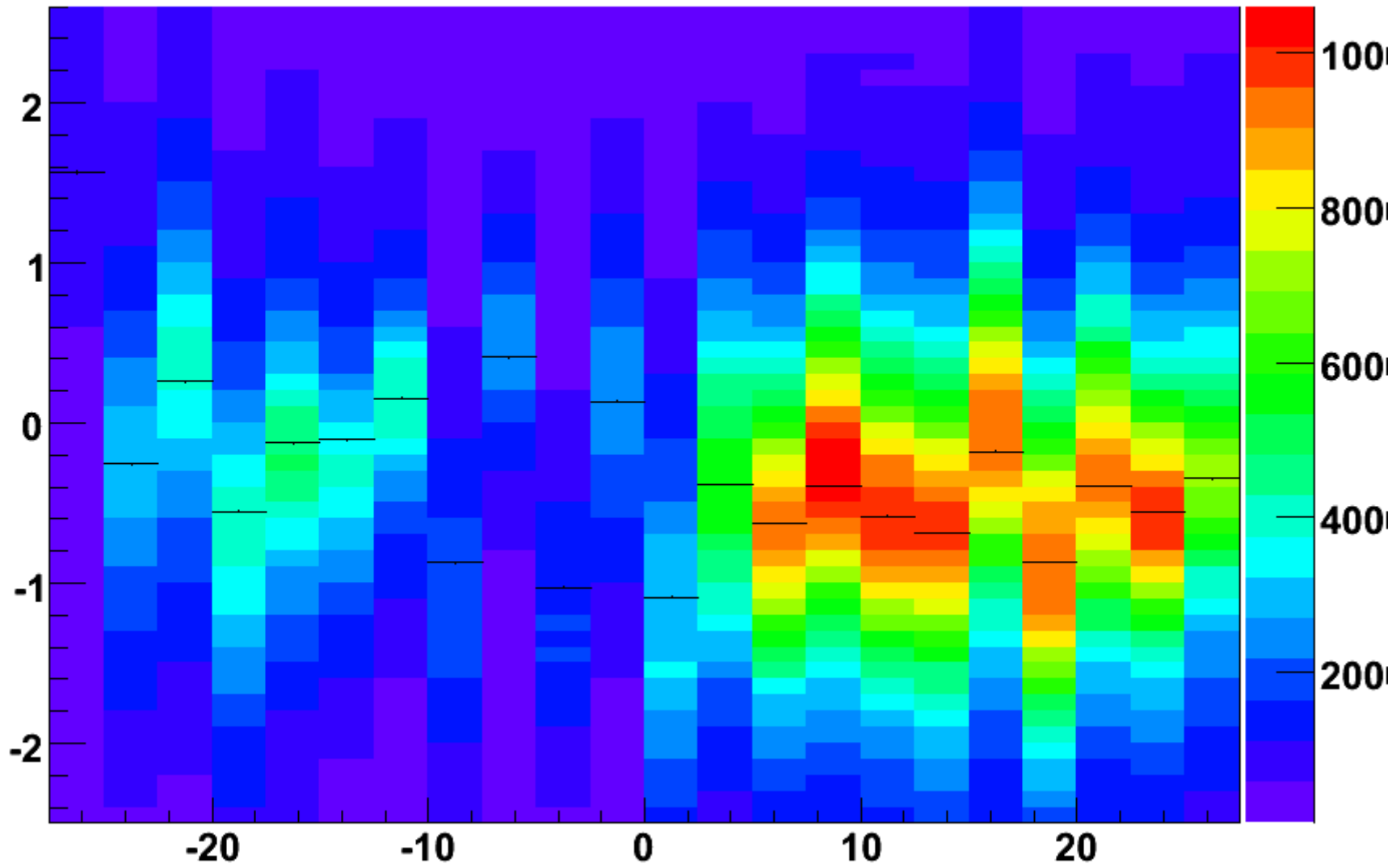
reszvsy\_ch1\_4

Entries 4564434



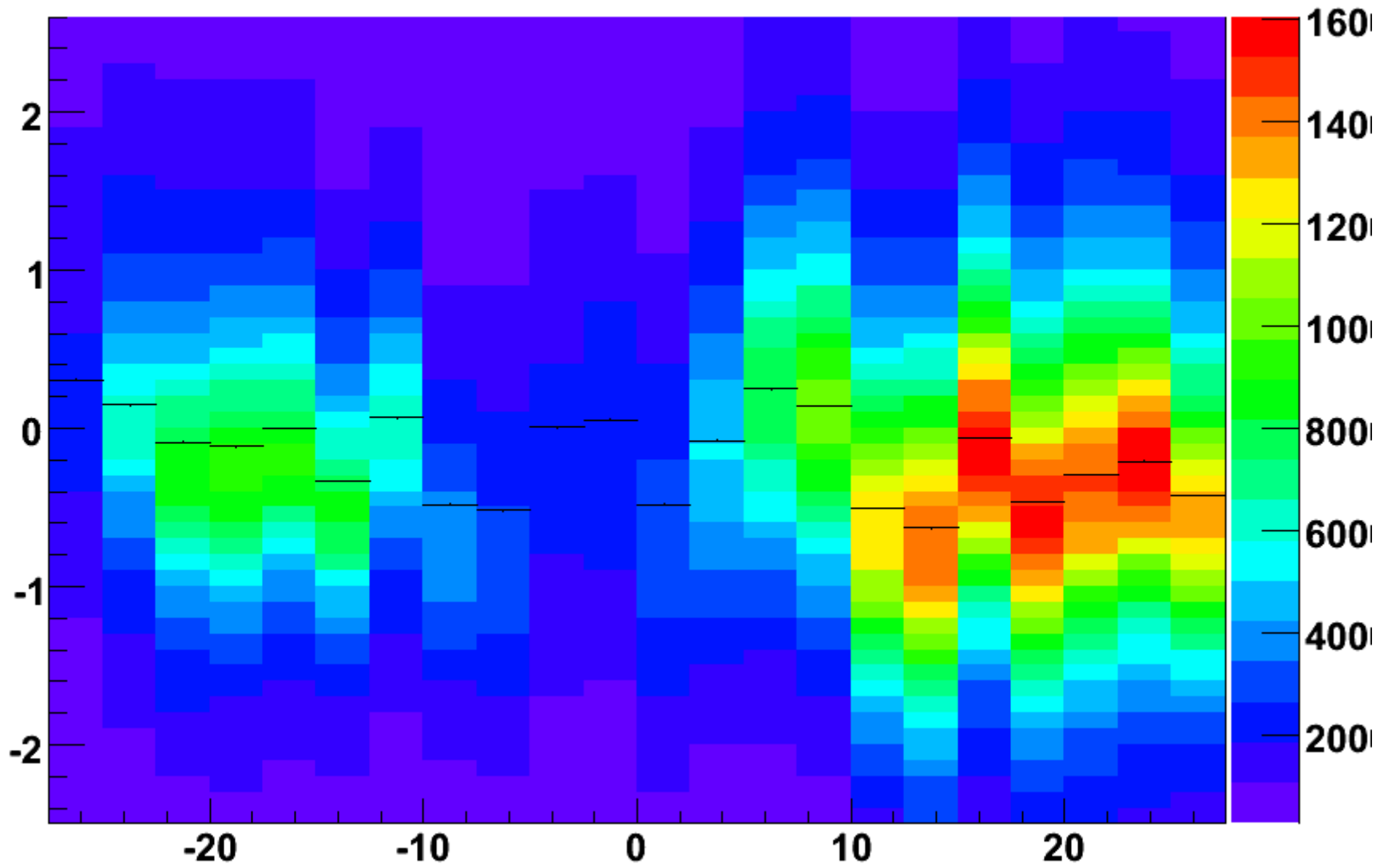
reszvsy\_ch2\_0

Entries 3099563



reszvsy\_ch2\_2

Entries 4725890



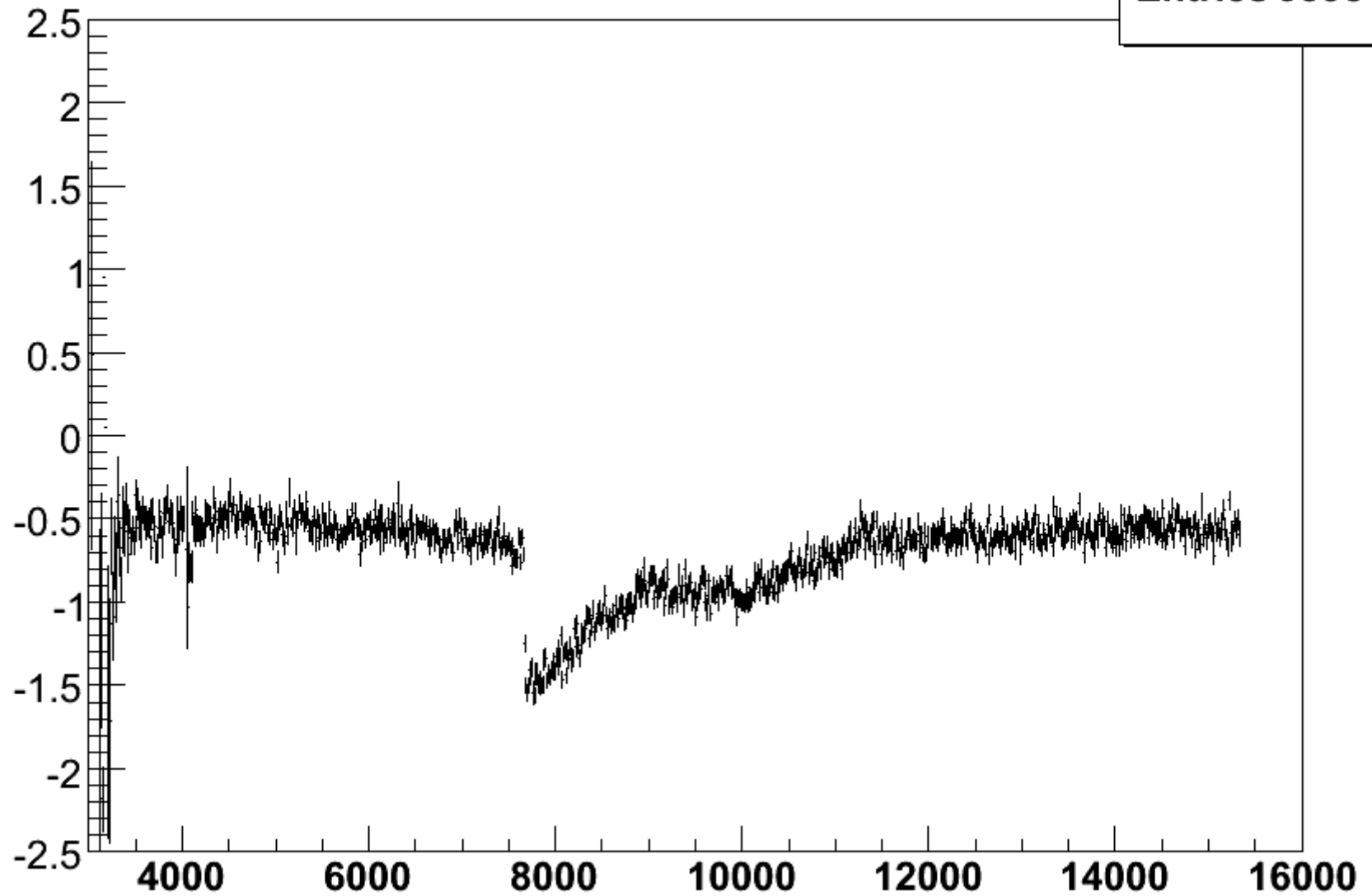
# Calibration and Alignment

- VDET aligned (looking at V60504)
- DCH1 and DCH2:
  - Different behavior of negative and positive tracks
  - Residual distributions depend on anodic wire and on **TIME !!!**
    - Z few centimeters
    - $\Phi$  few millimeters
- STRAW:
  - Waiting for reliable 2003-2004 Cosmic Rays Calibration Table on FINUDA Data Base (the basis of the global alignment...)



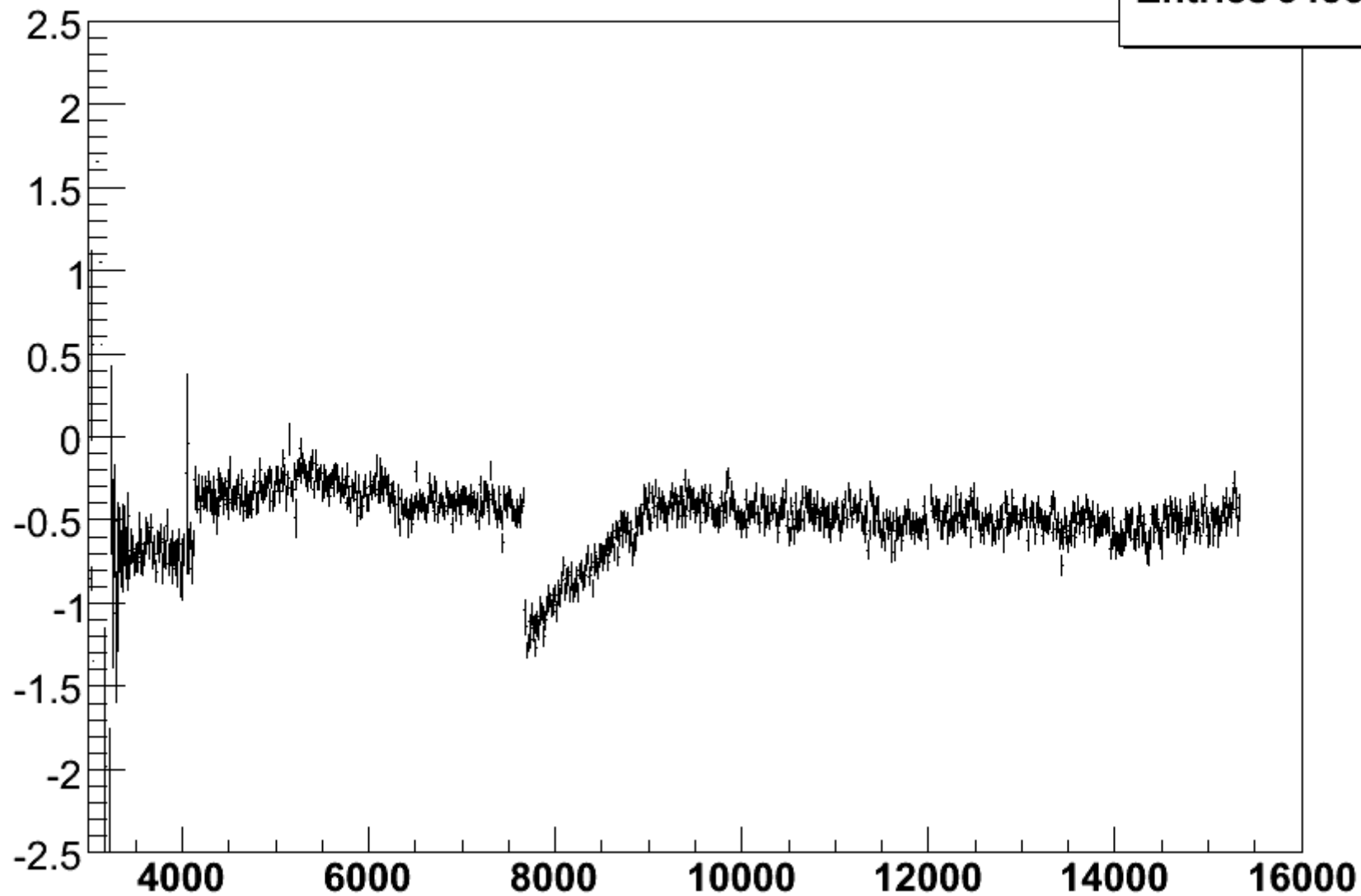
runvsreszvsy\_ch1\_0\_w\_03

Entries 359548



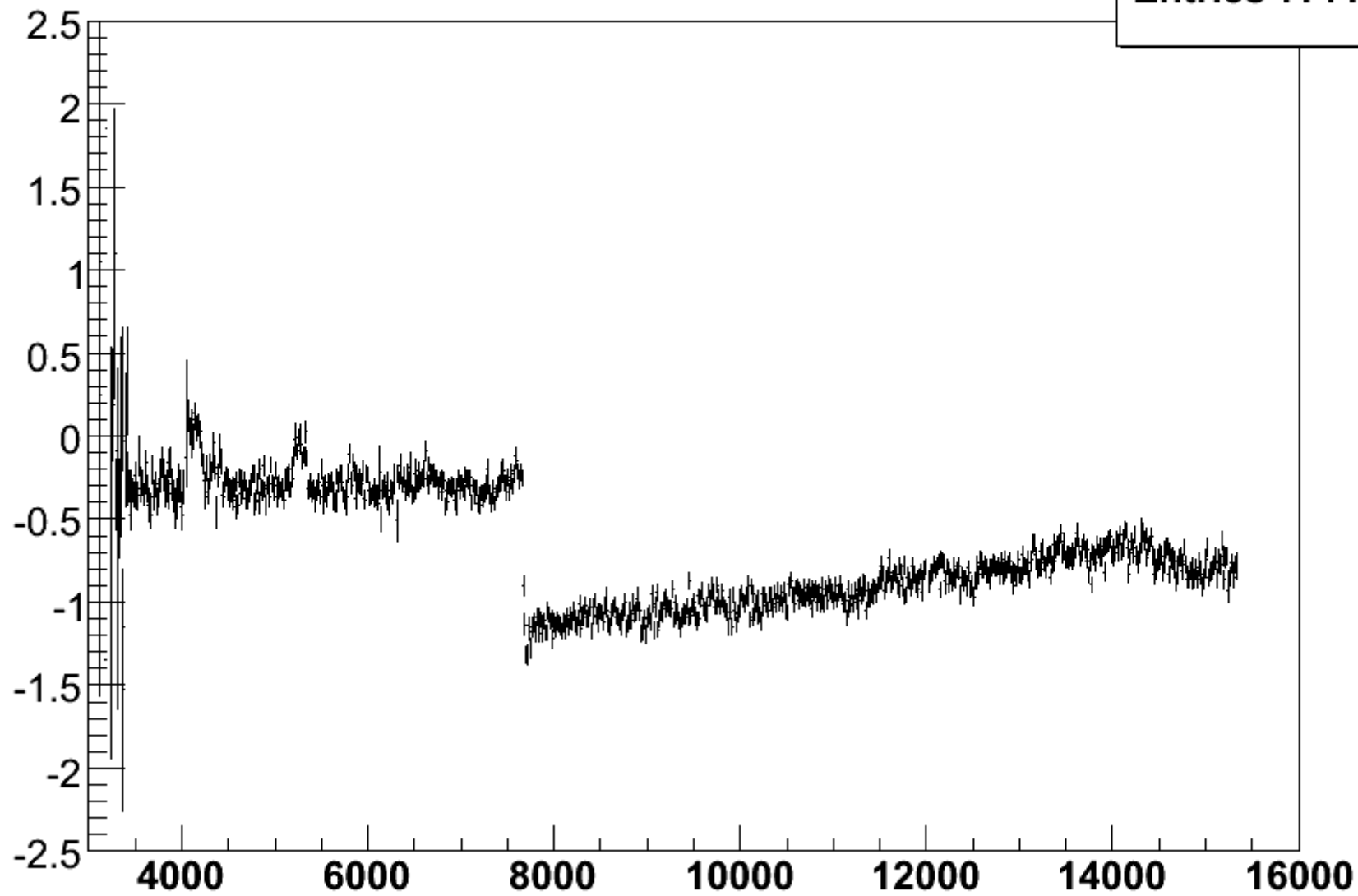
runvsreszvsy\_ch1\_0\_w\_05

Entries 345600



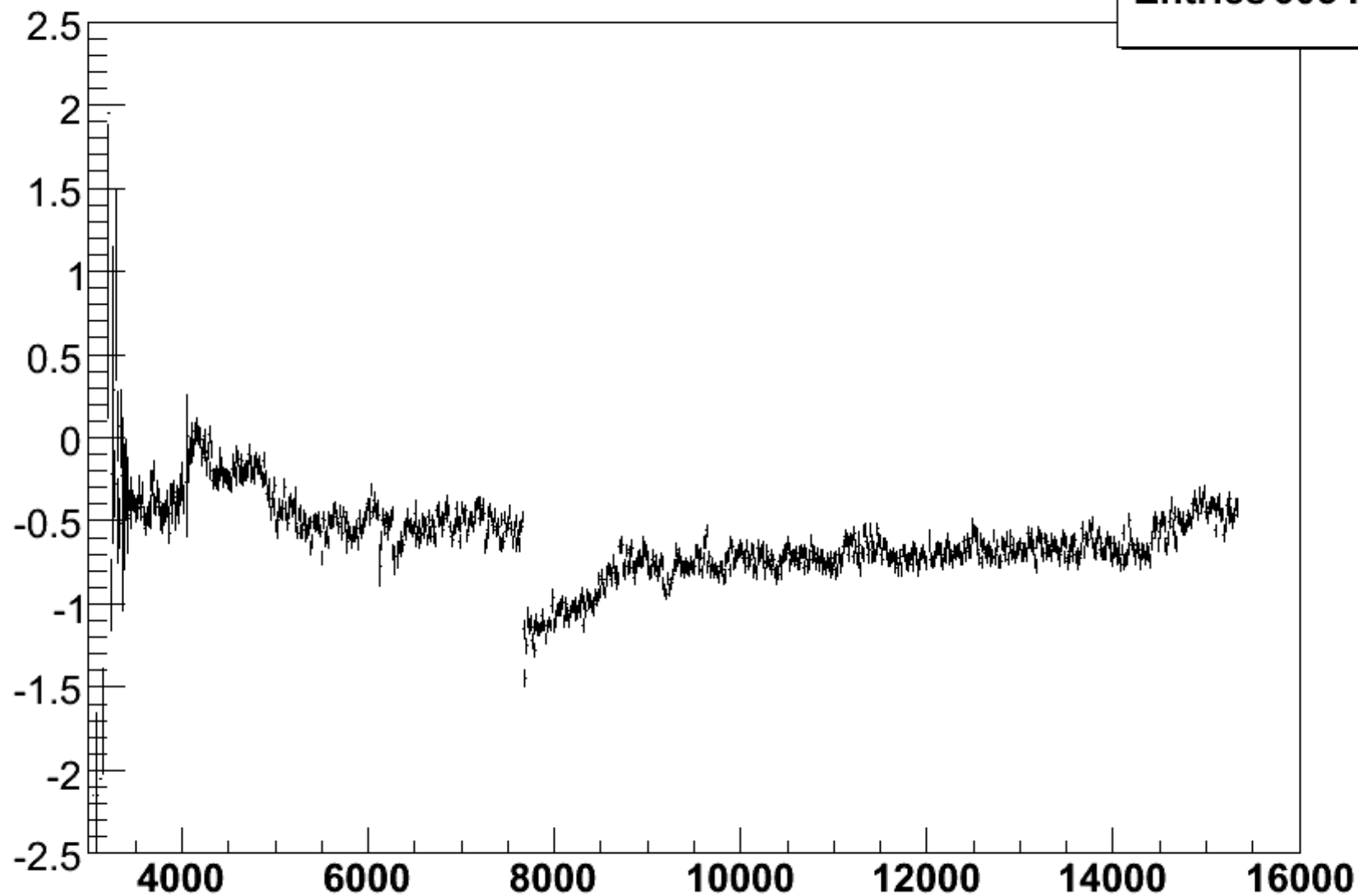
runvsreszvsy\_ch1\_0\_w\_10

Entries 414415



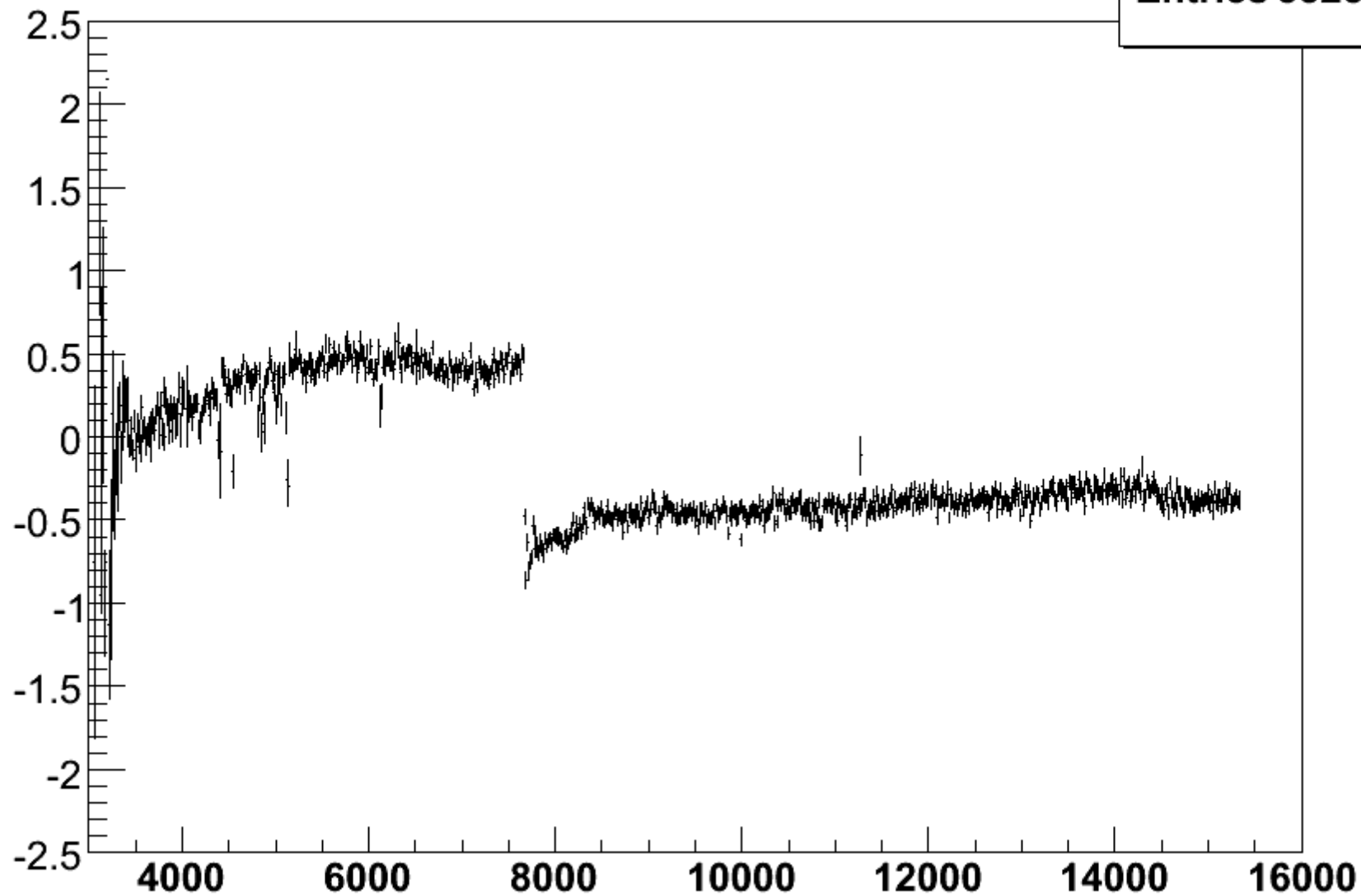
runvsreszvsy\_ch1\_1\_w\_04

Entries 603465



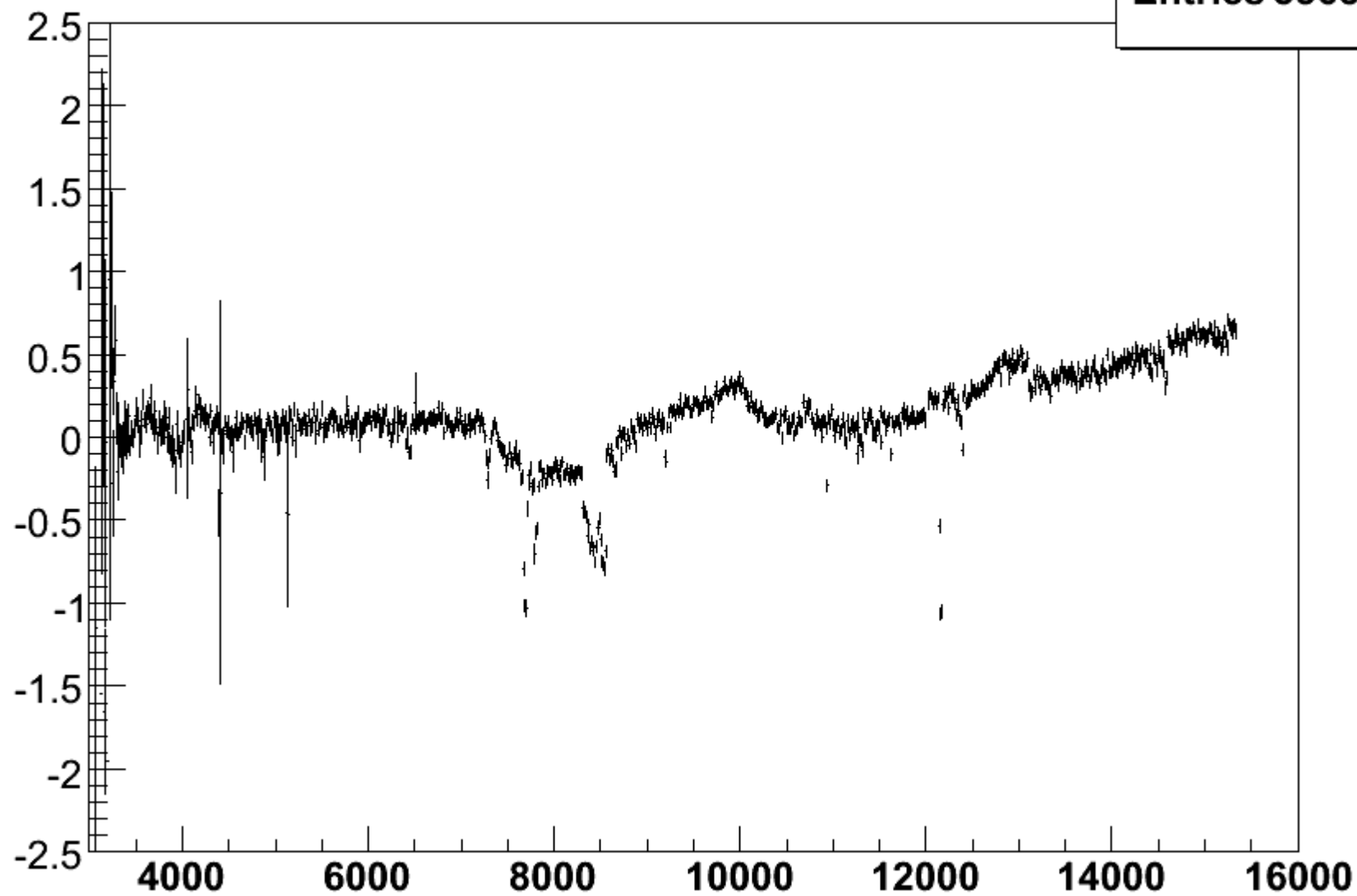
runvsreszvsy\_ch1\_2\_w\_04

Entries 552562



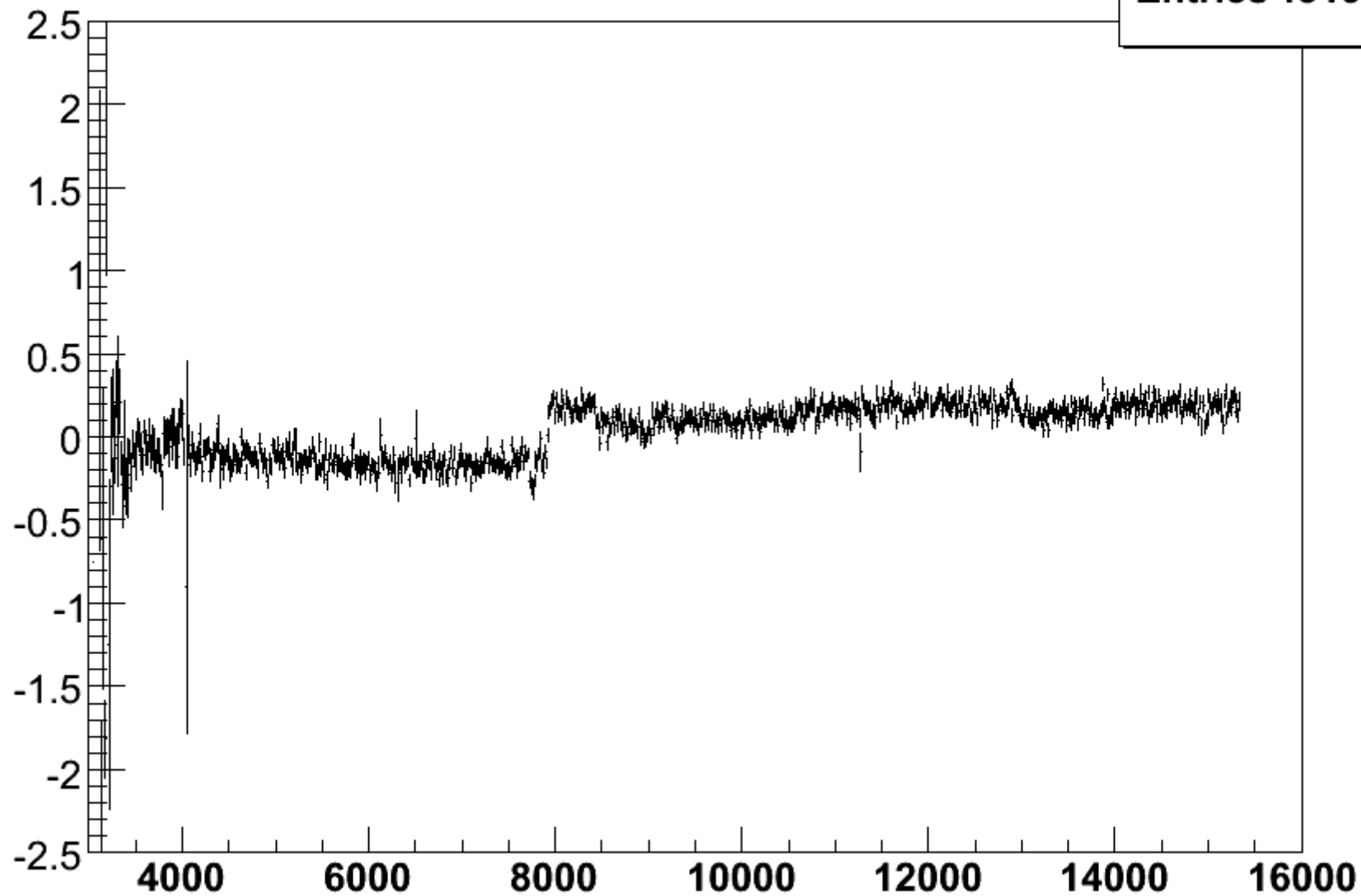
runvsreszvsy\_ch1\_2\_w\_05

Entries 556832



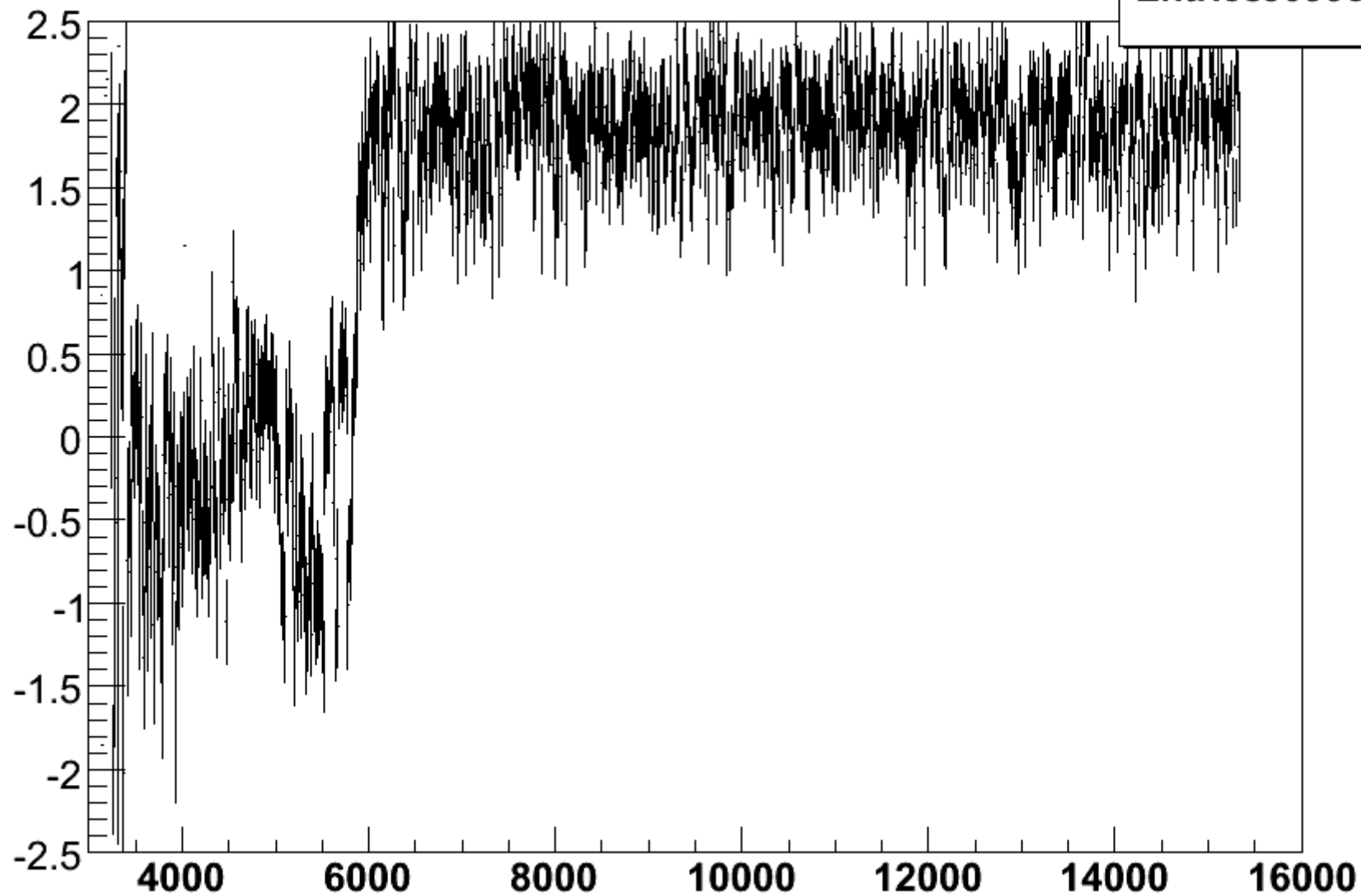
runvsreszvsy\_ch1\_3\_w\_06

Entries 431934



runvsreszvsy\_ch2\_0\_w\_00

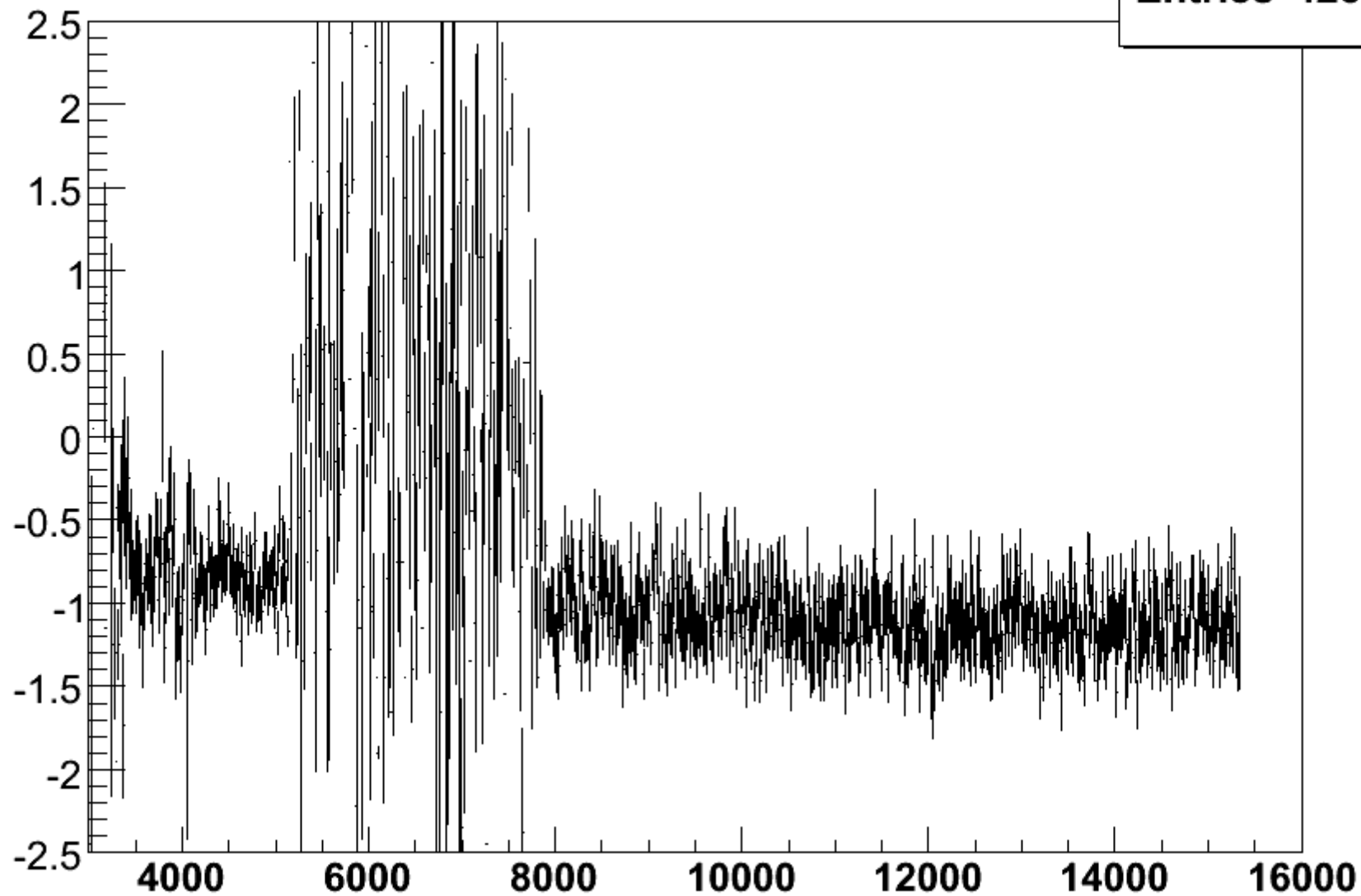
Entries3099563





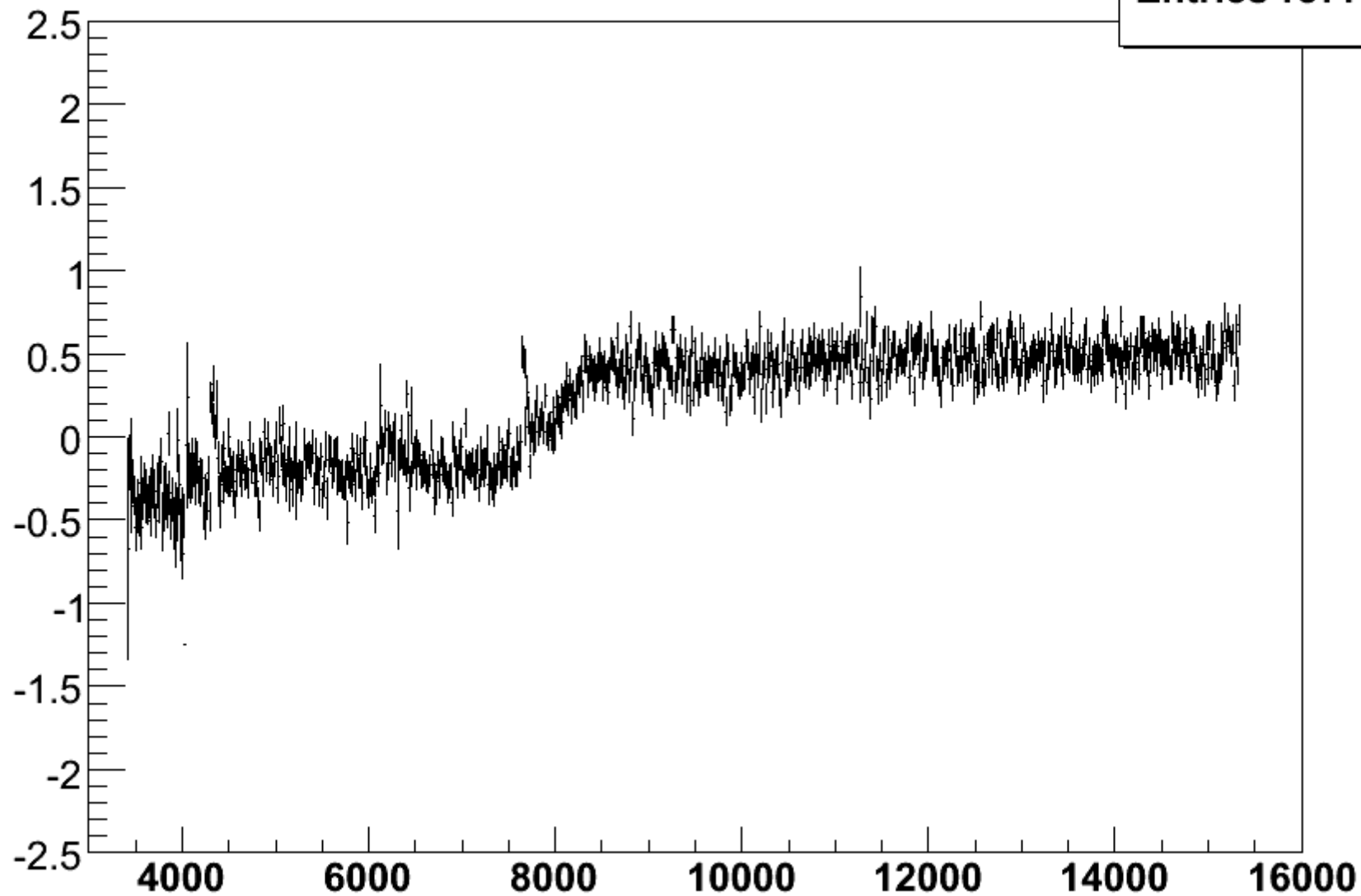
runvsreszvsy\_ch2\_0\_w\_09

Entries 42312



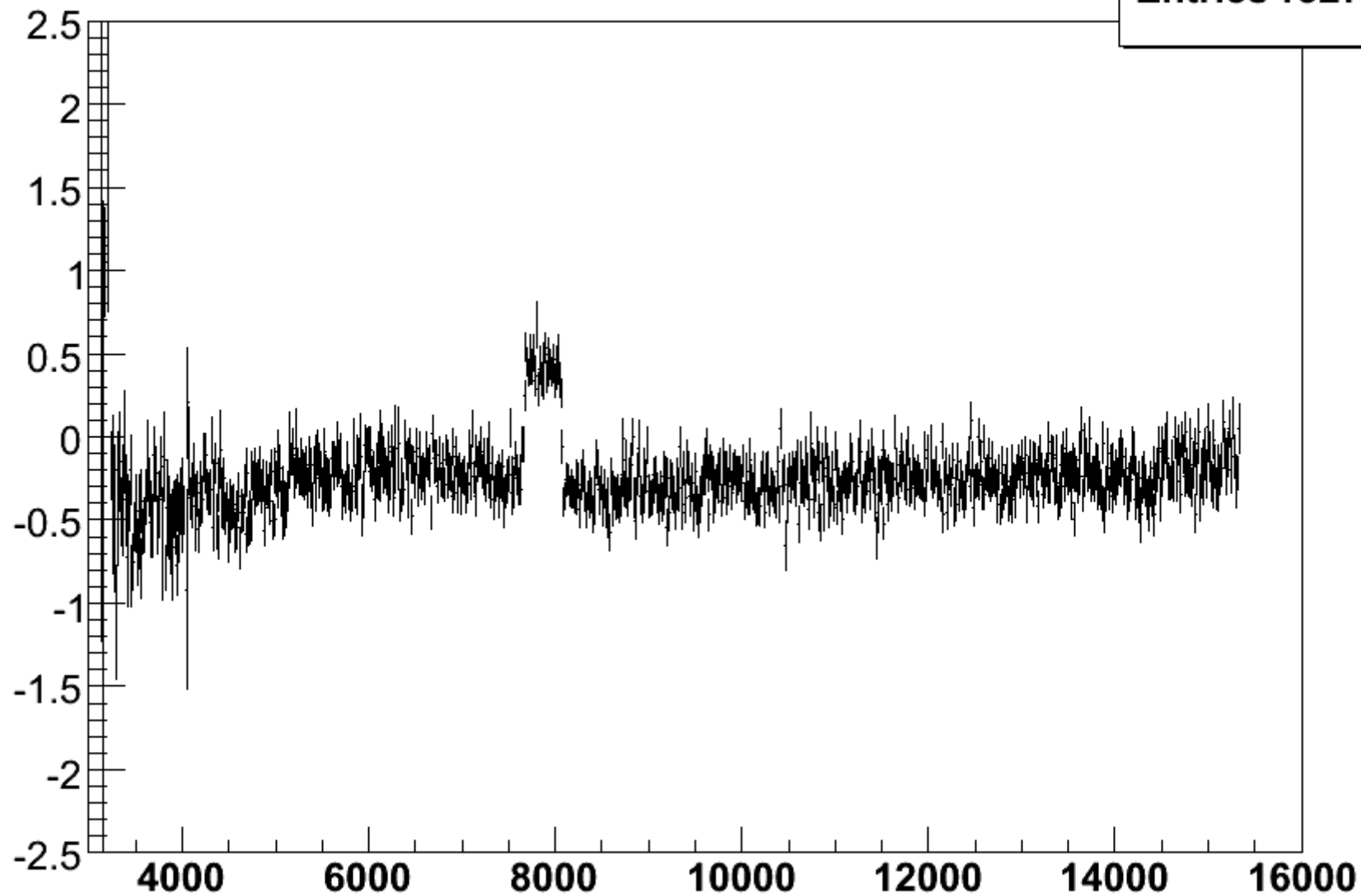
runvsreszvsy\_ch2\_1\_w\_05

Entries 187755



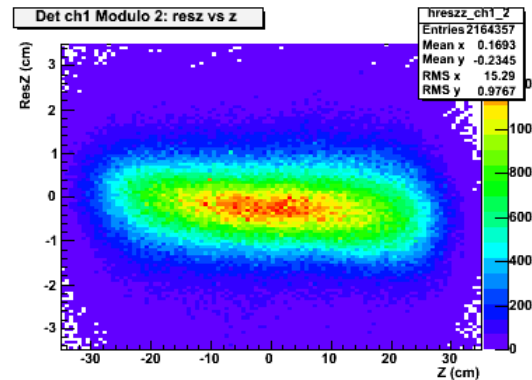
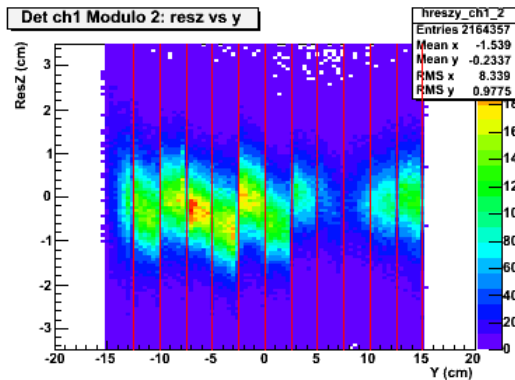
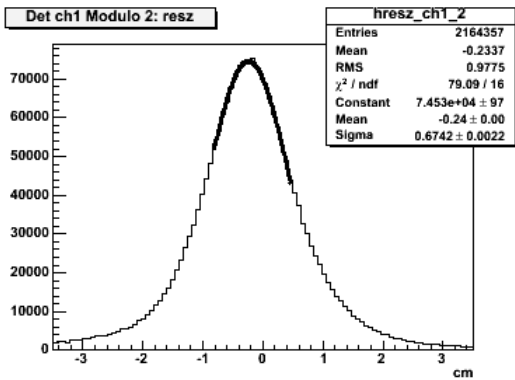
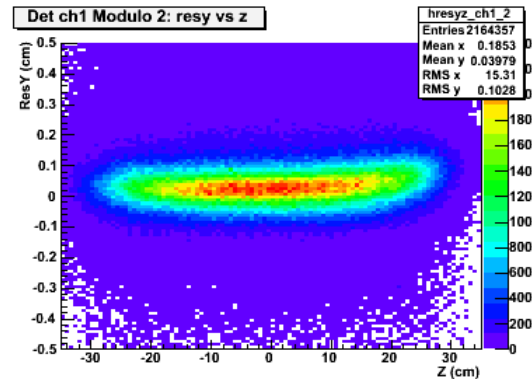
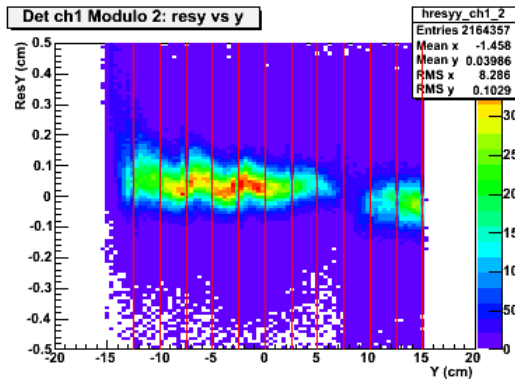
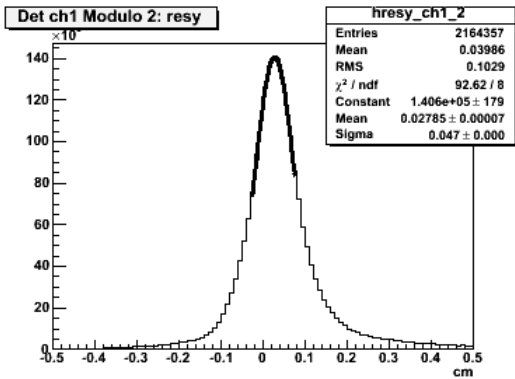
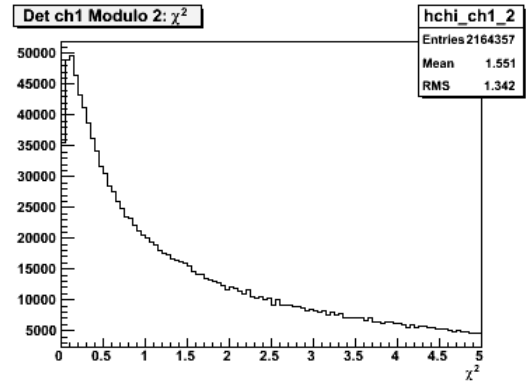
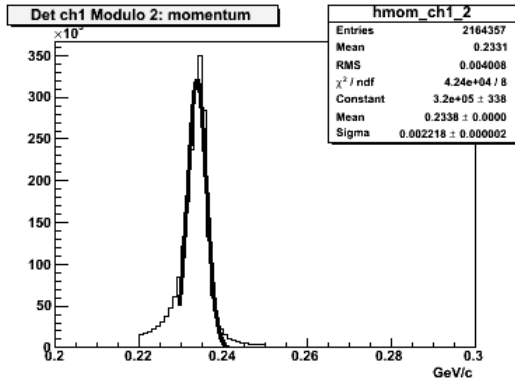
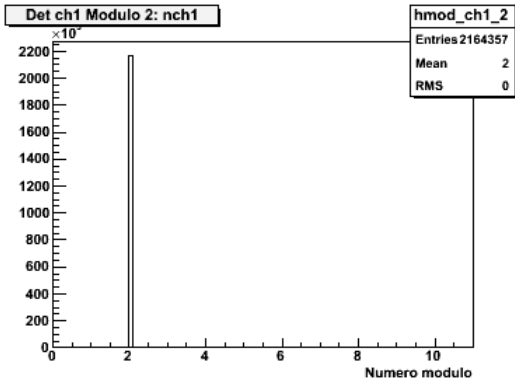
runvsreszvsy\_ch2\_4\_w\_05

Entries 132765



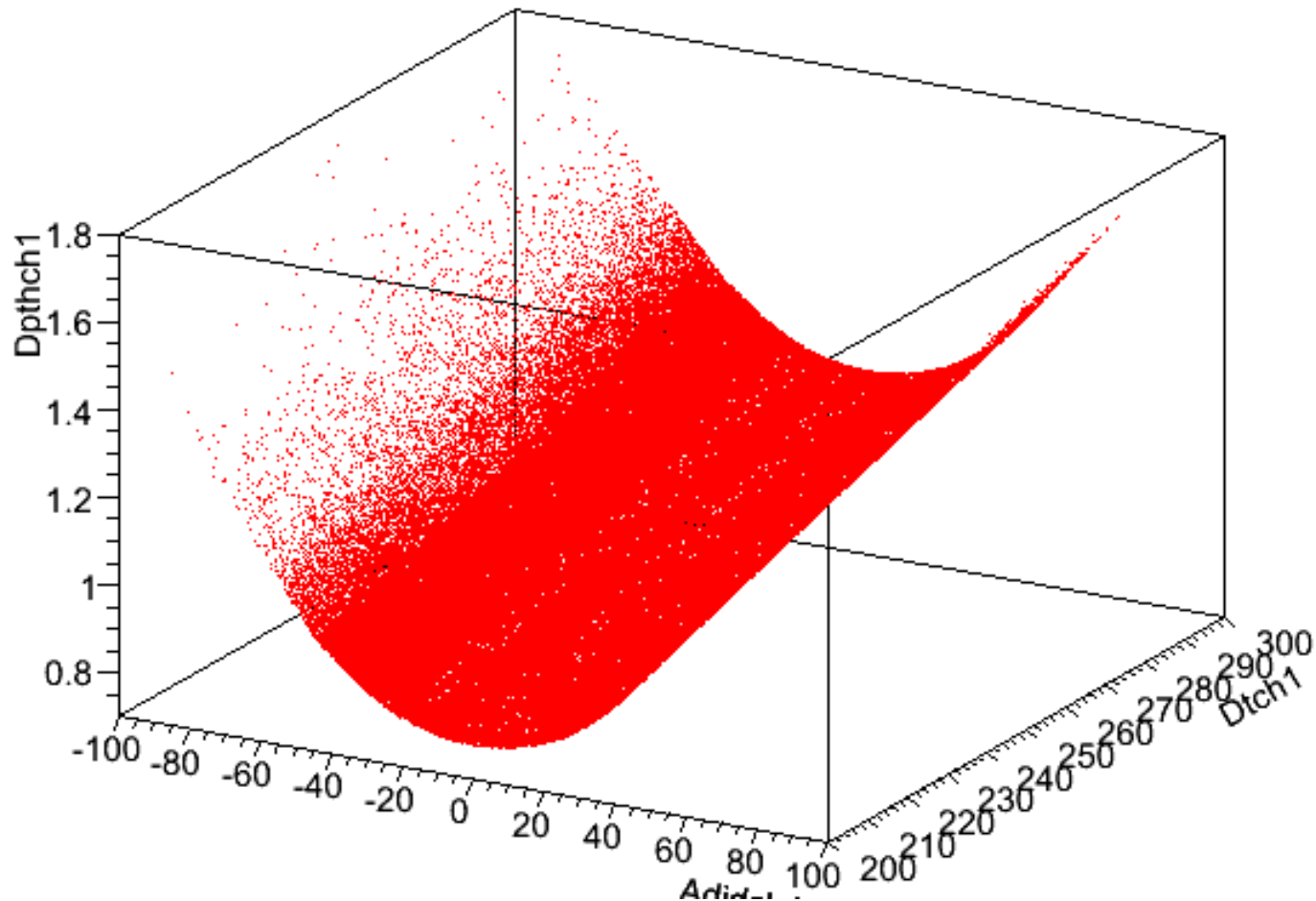
# Drift Chamber Z Side ... the ADC SIDE of the moon

- Residual mean values of few centimeters
- Strong dependence on Time
- Existing ADC calibrations are not good enough to get acceptable resolution on long and short tracks
- Please, check the ADC calibrations run-by-run



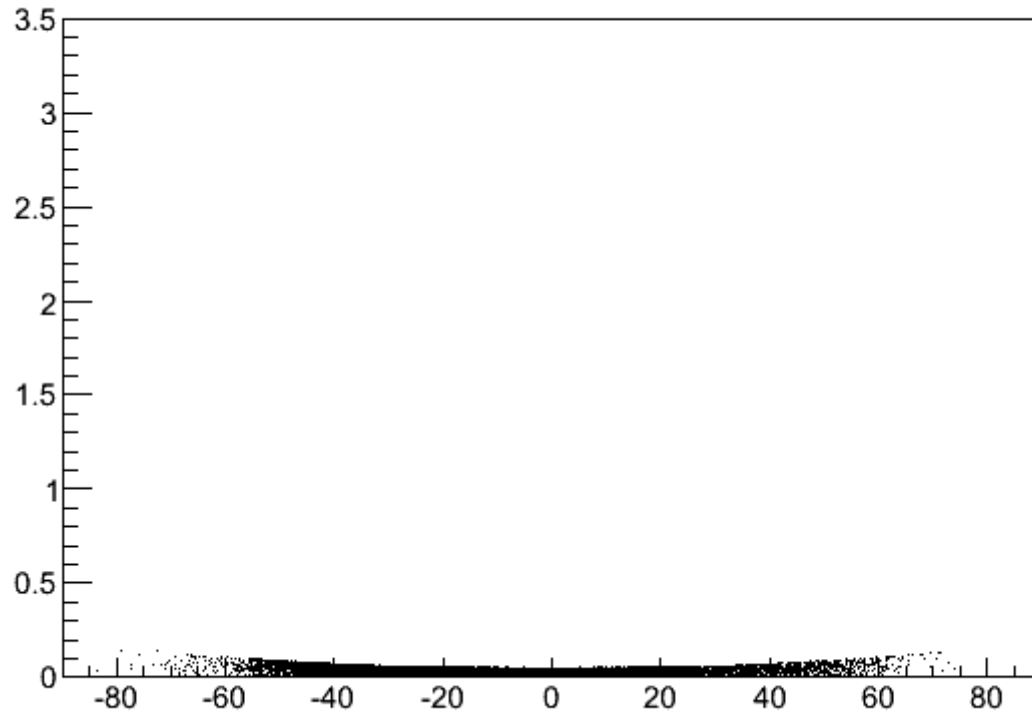
# $\Phi$ Side: Garfield Correction vs Drift Time and vs Attach Angle

Dpthch1:Dch1:Adjch1 [Mom > 0.030 && Mom < 0.800 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Dtch1 > 200 && Dtch1 < 300]



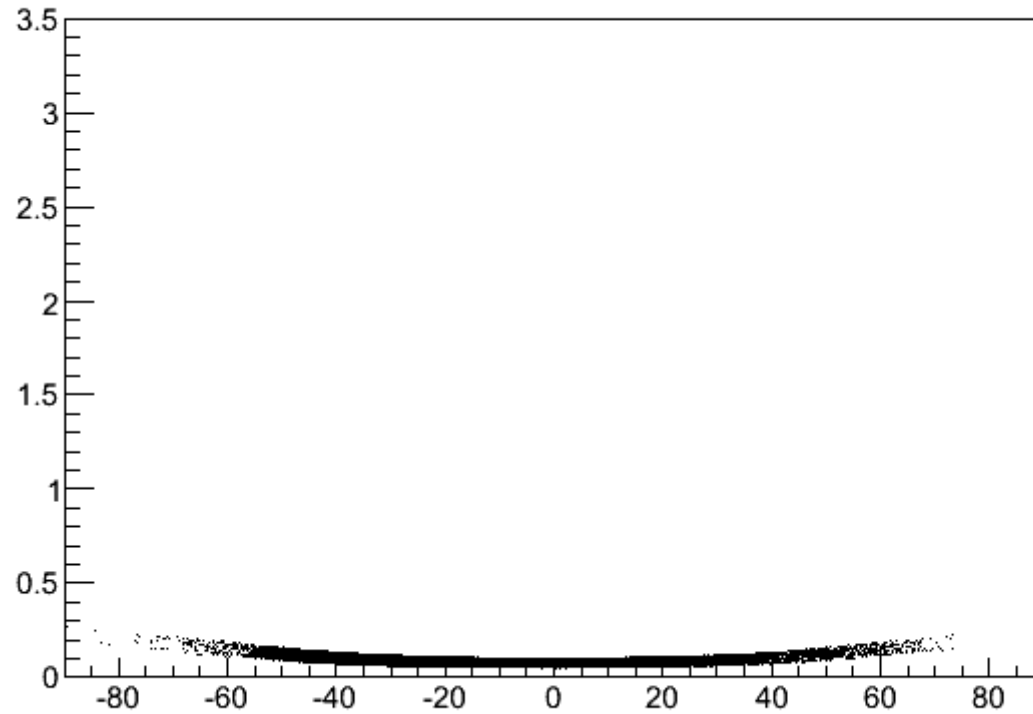
# Garfield Correction vs Attach Angle with $0 \text{ ns} < \text{Drift Time} < 10 \text{ ns}$

Dpthch1:Adjch1 [Mom > 0.030 && Mom < 0.600 && CH2 < 100 && Nch1 > 0 && Nch1 < 9 && Dch1 > 0 && Dtzch1 < 10]



# Garfield Correction vs Attach Angle with $10 \text{ ns} < \text{Drift Time} < 20 \text{ ns}$

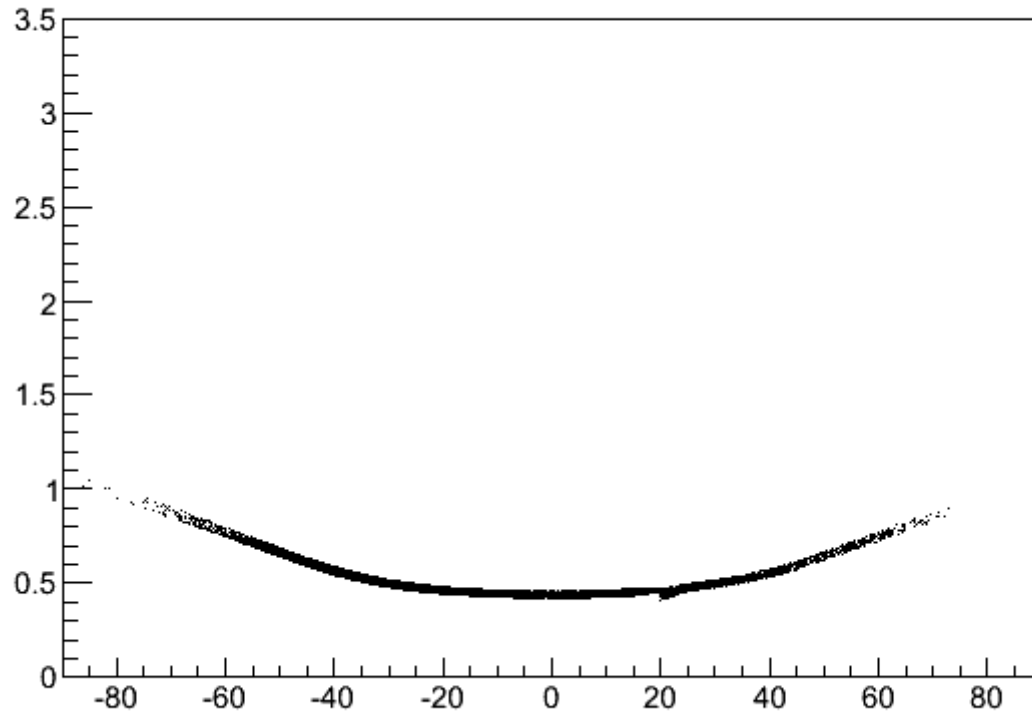
Dp[hch1:Adj]ch1 [Mom > 0.030 && Mom < 0.600 && CH2 < 100 && Nch1 > 0 && Nch1 < 9 && Dch1 > 10 && Dch1 < 20]





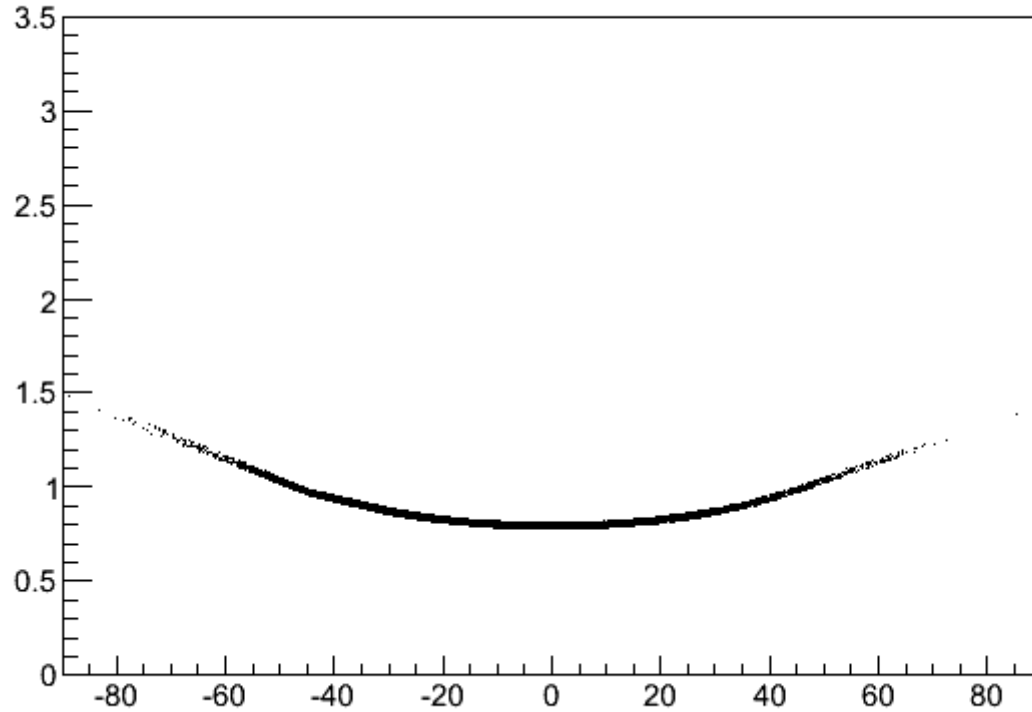
# Garfield Correction vs Attach Angle with $100 \text{ ns} < \text{Drift Time} < 110 \text{ ns}$

Dpthch1:Adjch1 {Mom > 0.030 && Mom < 0.600 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Drch1 > 100 && Drch1 < 110}



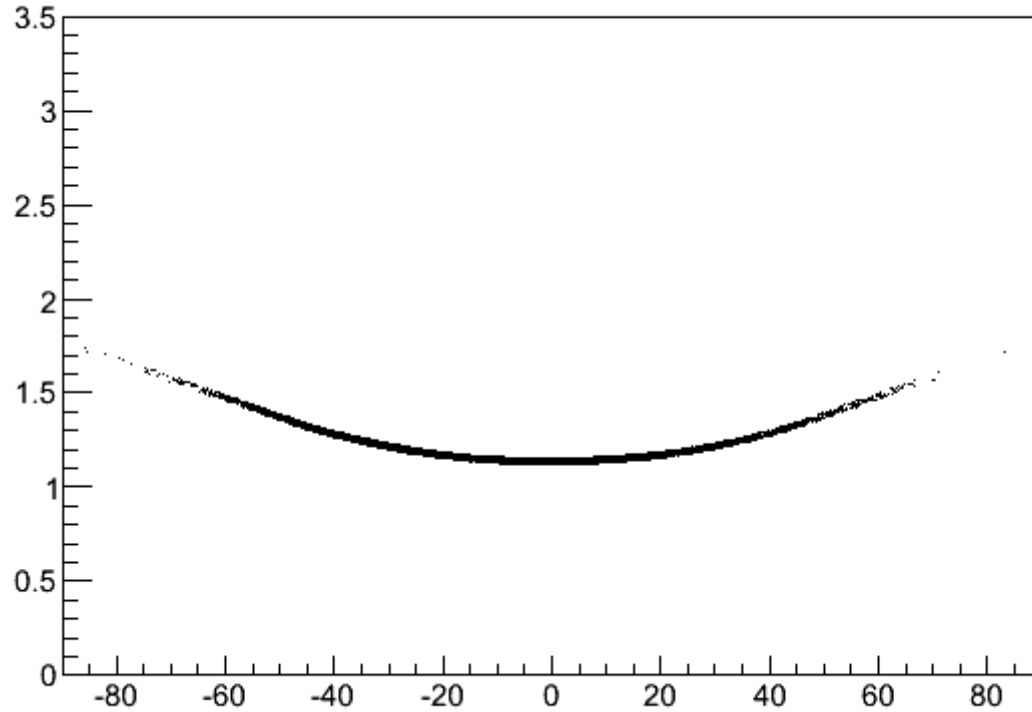
# Garfield Correction vs Attach Angle with $200 \text{ ns} < \text{Drift Time} < 210 \text{ ns}$

Dpthch1:Adjch1 [Mom > 0.030 && Mom < 0.600 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Dch1 > 200 && Dch1 < 210]



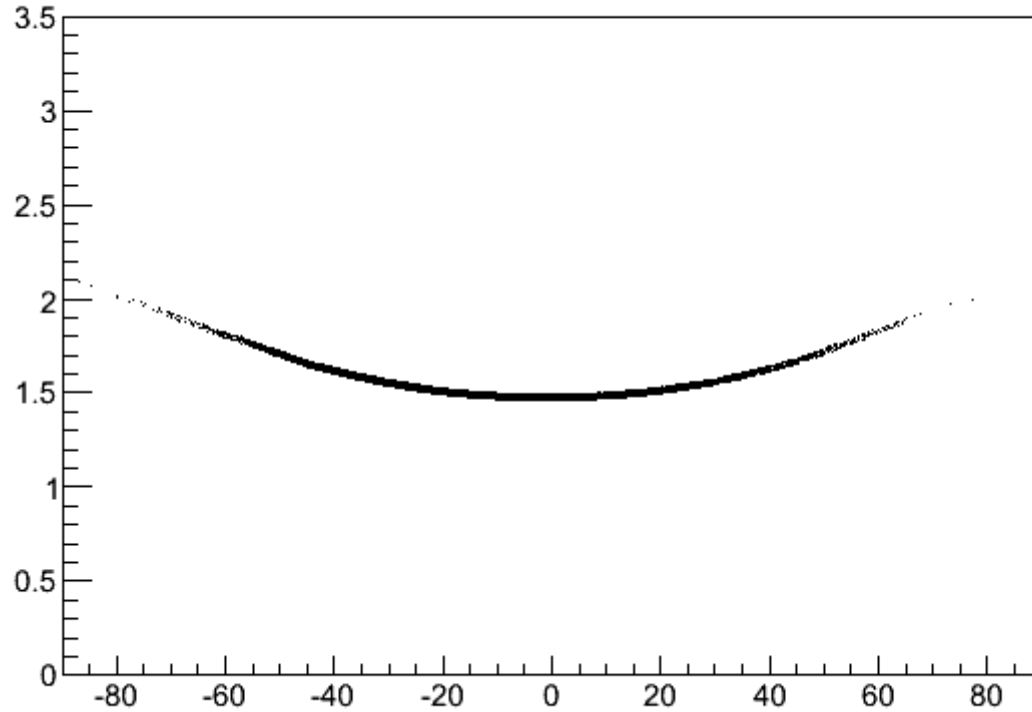
# Garfield Correction vs Attach Angle with $300 \text{ ns} < \text{Drift Time} < 310 \text{ ns}$

Dpthch1:Adjch1 [Mom > 0.030 && Mom < 0.600 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Dch1 > 300 && Dch1 < 310]



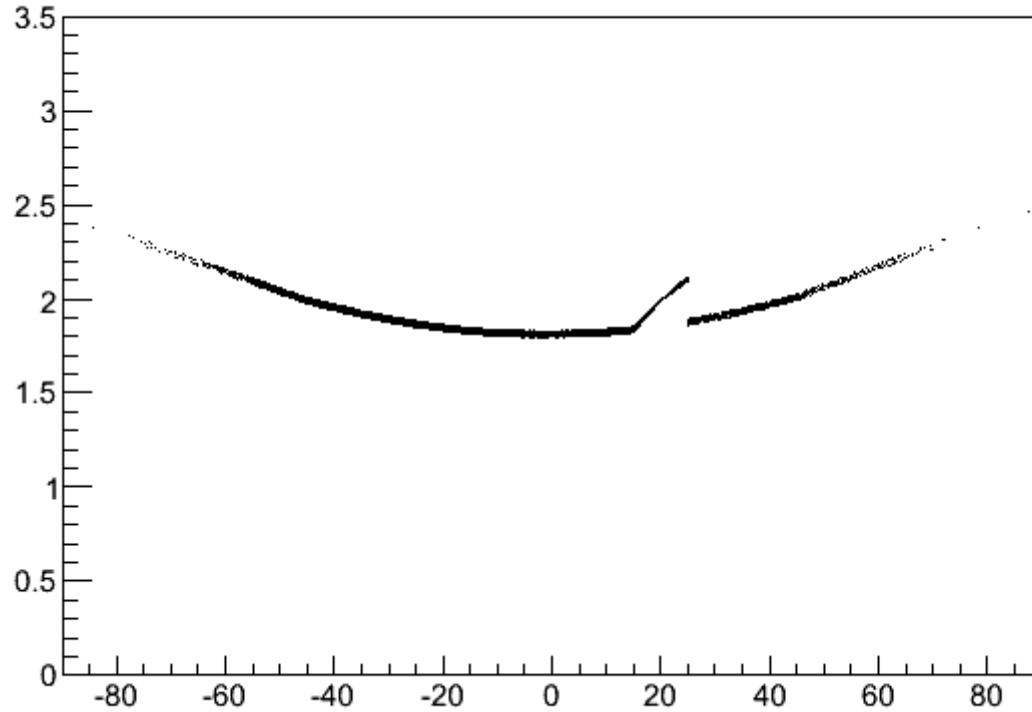
# Garfield Correction vs Attach Angle with $400 \text{ ns} < \text{Drift Time} < 410 \text{ ns}$

Dpthch1:Adjch1 [Mom > 0.030 && Mom < 0.600 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Dch1 > 400 && Dch1 < 410]



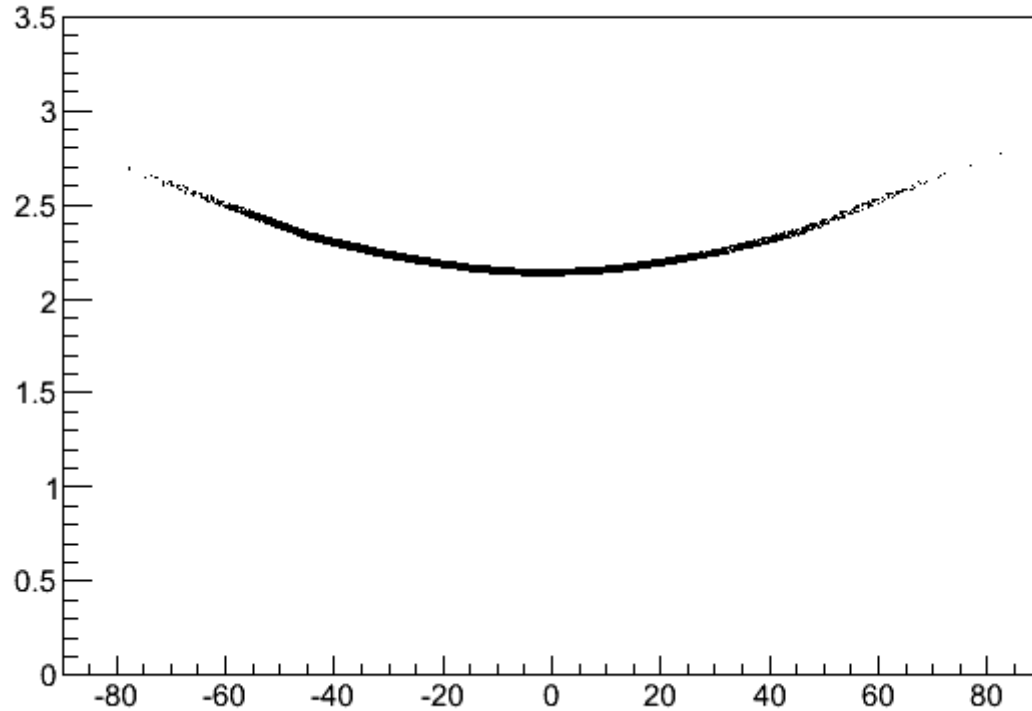
# Garfield Correction vs Attach Angle with $500 \text{ ns} < \text{Drift Time} < 510 \text{ ns}$

Dpthch1:Adjch1 [Mom > 0.030 && Mom < 0.600 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Dch1 > 500 && Dch1 < 510]



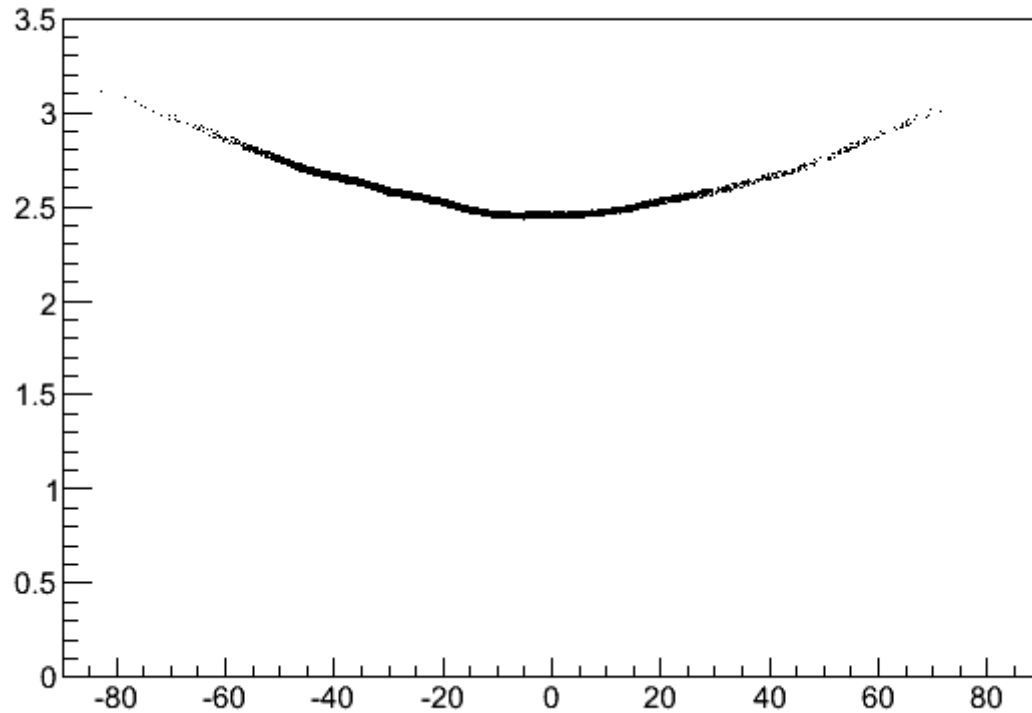
# Garfield Correction vs Attach Angle with $600 \text{ ns} < \text{Drift Time} < 610 \text{ ns}$

Dpthch1:Adjch1 [Mom > 0.030 && Mom < 0.600 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Dch1 > 600 && Dch1 < 610]

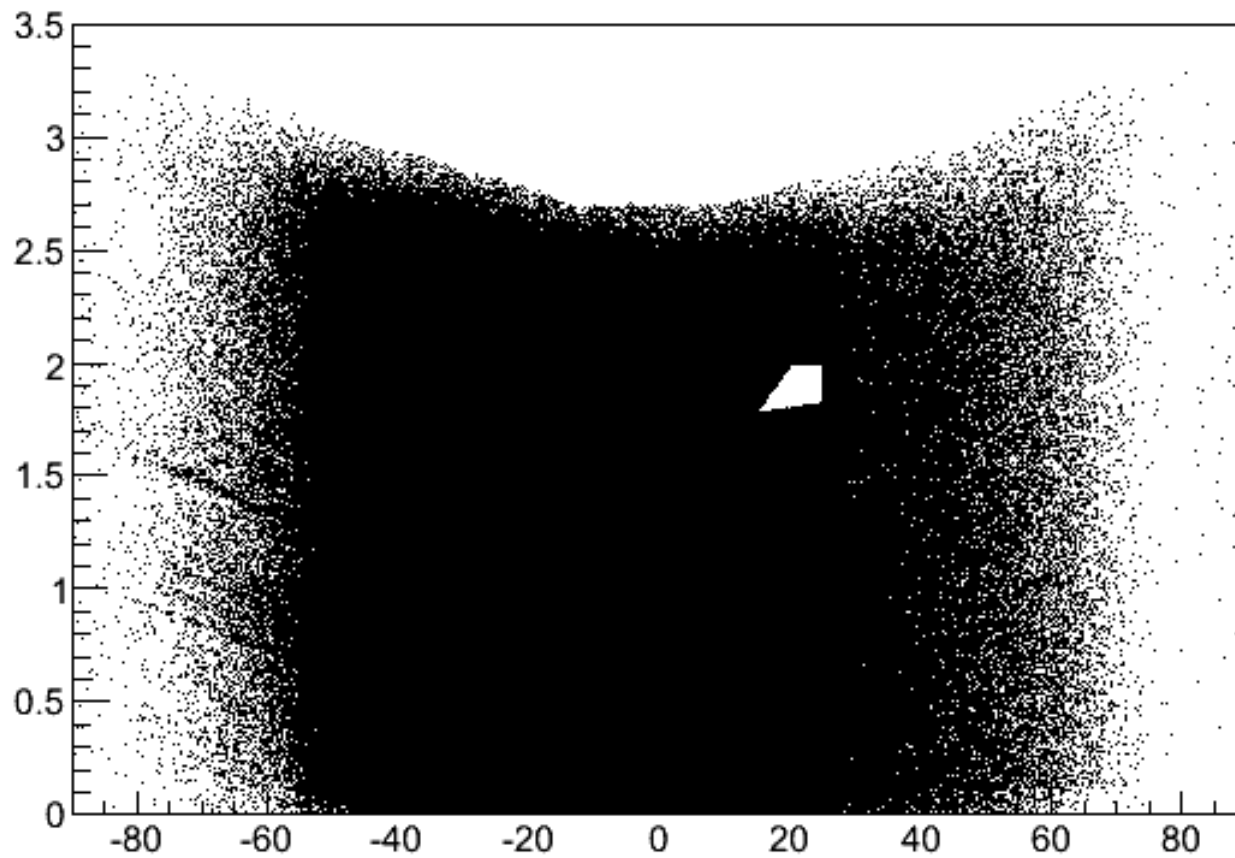


# Garfield Correction vs Attach Angle with $700 \text{ ns} < \text{Drift Time} < 710 \text{ ns}$

Dpthch1:Adjch1 [Mom > 0.030 && Mom < 0.600 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Dtch1 > 700 && Dtch1 < 710]



Dpthch1:Adjch1 [Mom > 0.030 && Mom < 0.600 && Ch2 < 100 && Nch1 > 0 && Nch1 < 9 && Dch1 > 0 && Dch1 < 800]

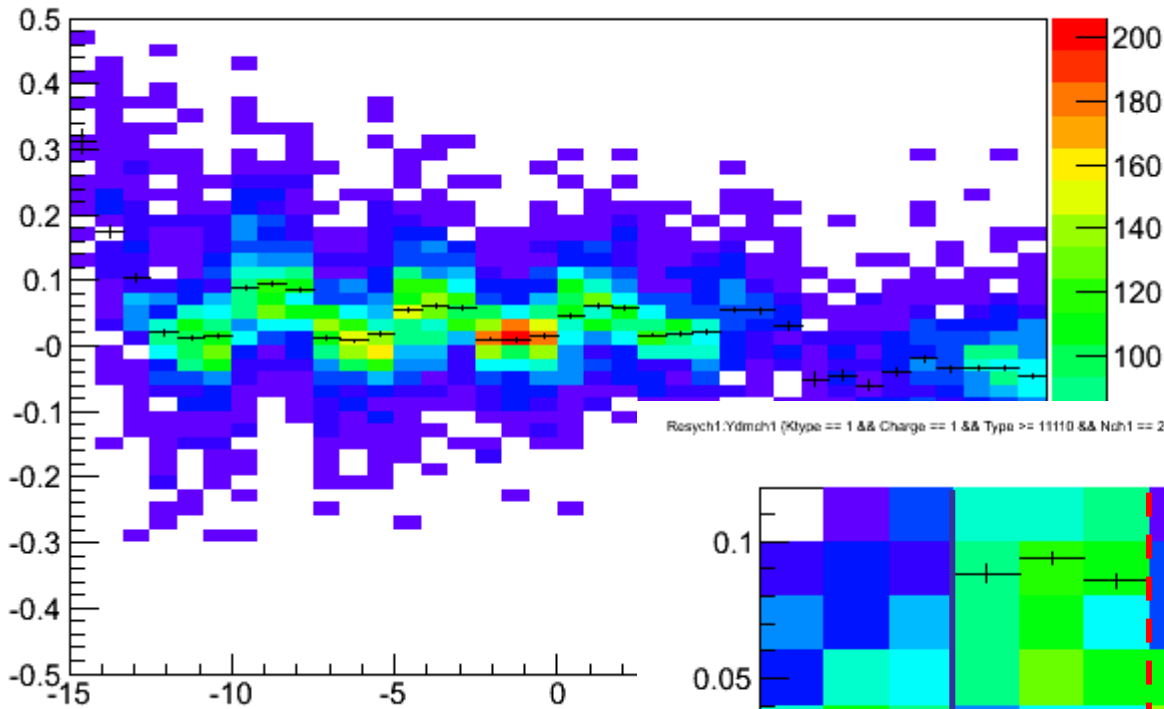




# $\Phi$ residual vs anodic wire

Resych1.Ydmch1 (KType == 1 && Charge == 1 && Type == 11110 && Nch1 == 2 && Ch2 < 100 && Mom > 0.230 && Mom < 0.240)

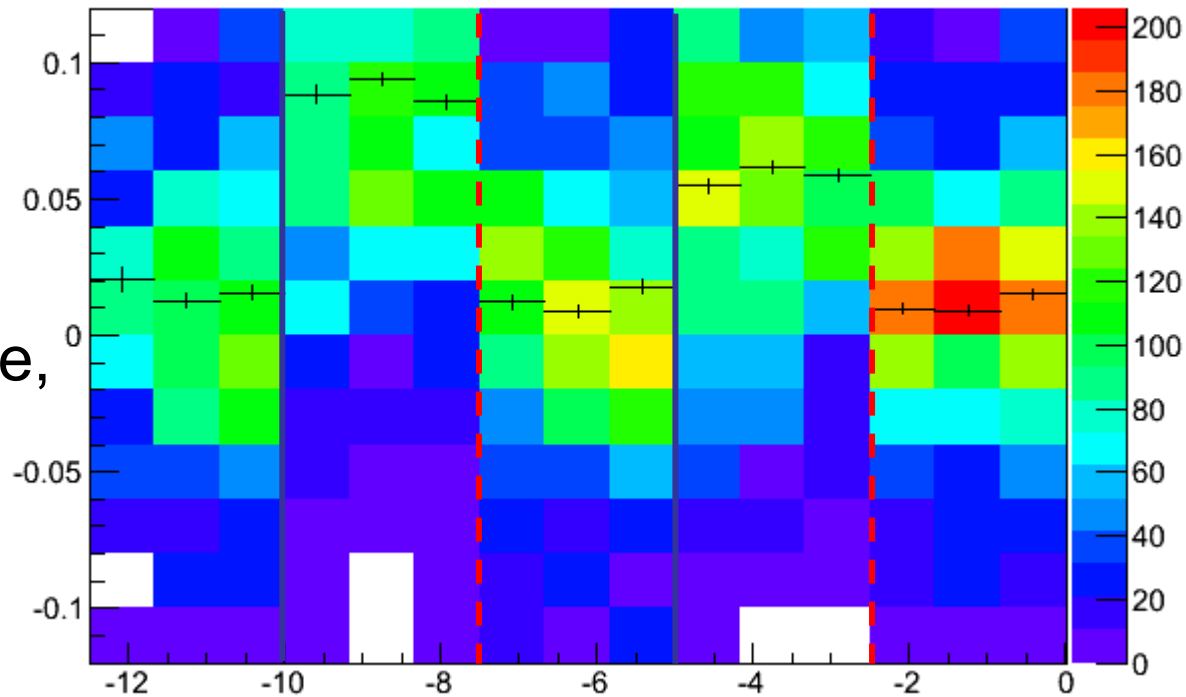
Entries 18903



Resych1.Ydmch1 (KType == 1 && Charge == 1 && Type == 11110 && Nch1 == 2 && Ch2 < 100 && Mom > 0.230 && Mom < 0.240)

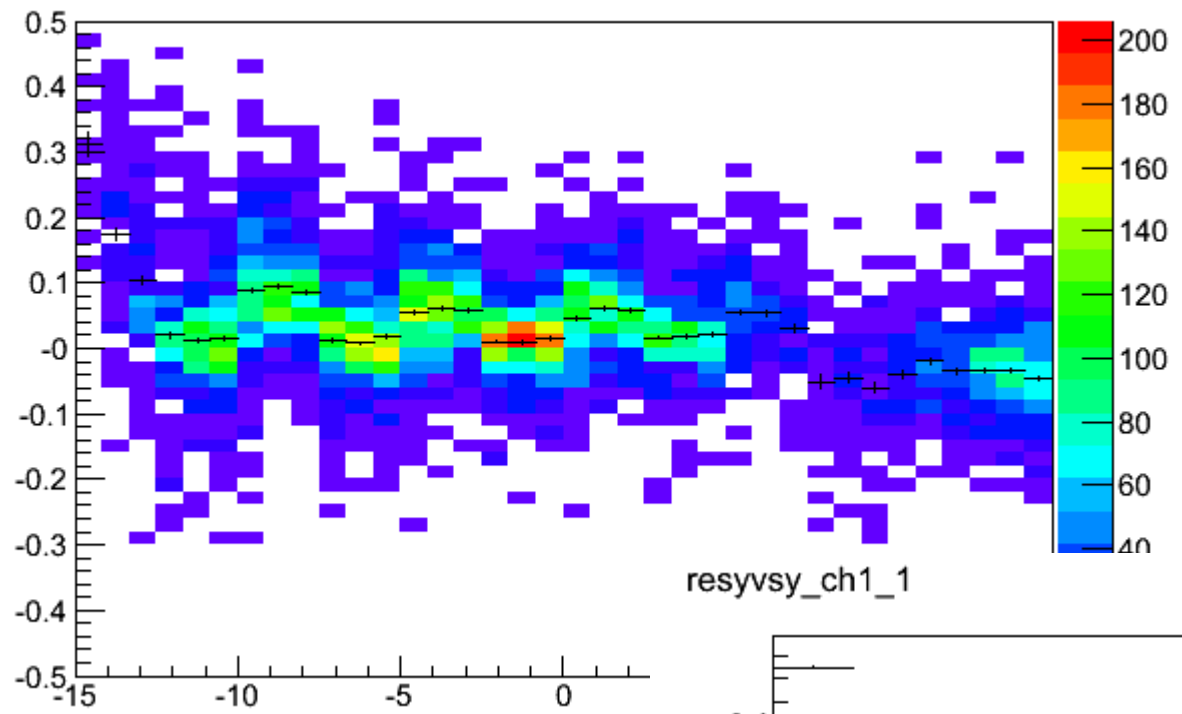
Entries 18903

Cell-by-cell dependence,  
enough uniform inside  
the cell

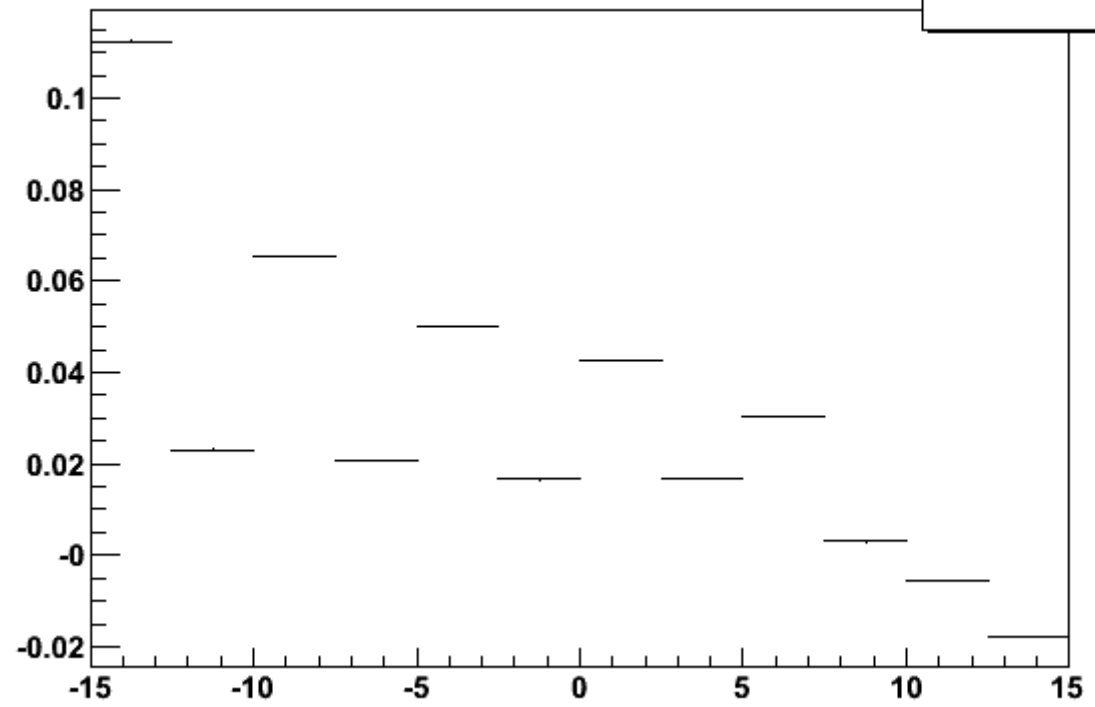


ResyCh1.YdmCh1 {KType == 1 && Charge == 1 && Type == 11110 && Nch1 == 2 && Ch2 < 100 && Mom > 0.230 && Mom < 0.240}

Entries 18903

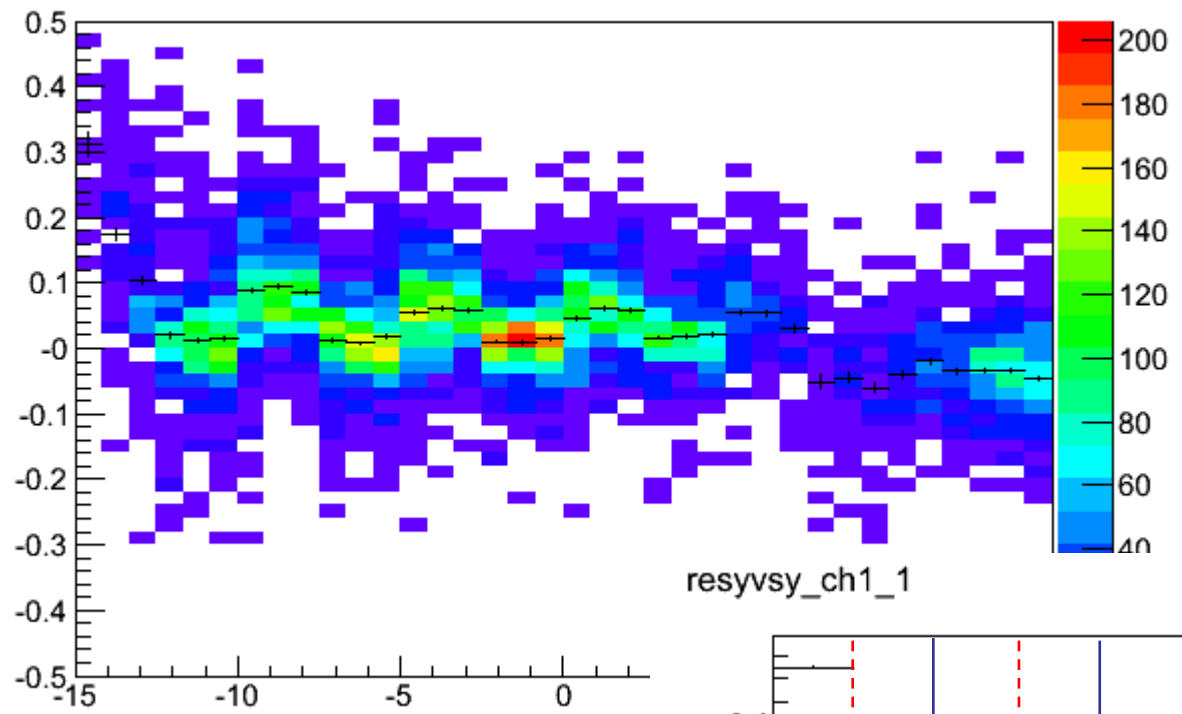


Entry 512675



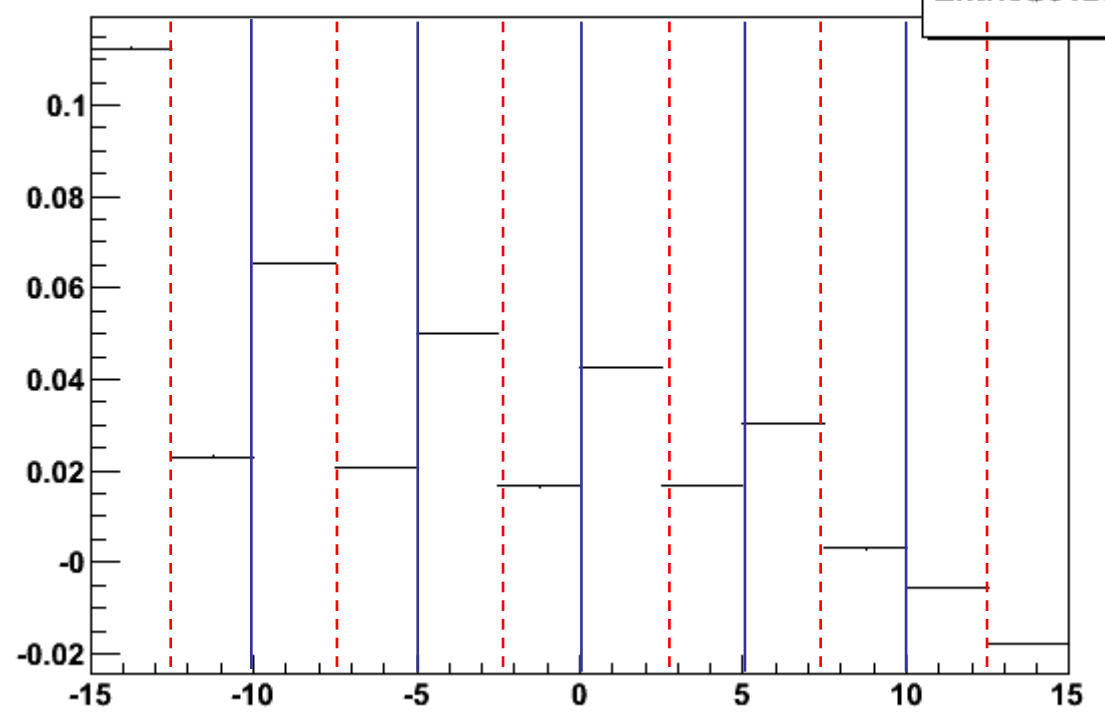
ResyCh1.YdmCh1 (kType == 1 && Charge == 1 && Type == 11110 && Nch1 == 2 && Ch2 < 100 && Mom > 0.230 && Mom < 0.240)

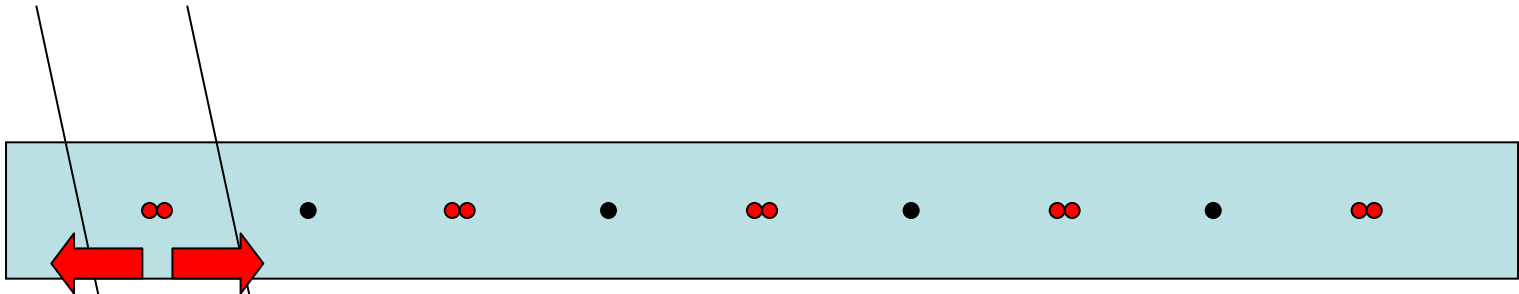
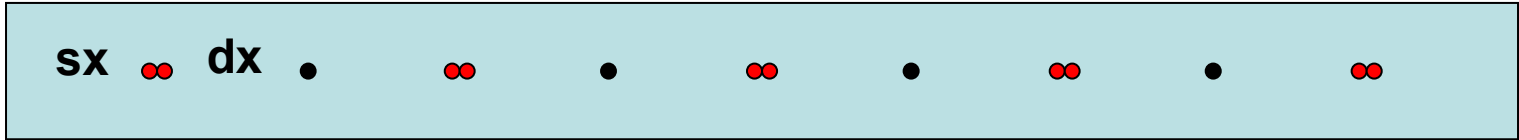
Entries 18903



resyvsy\_ch1\_1

Entries 512675

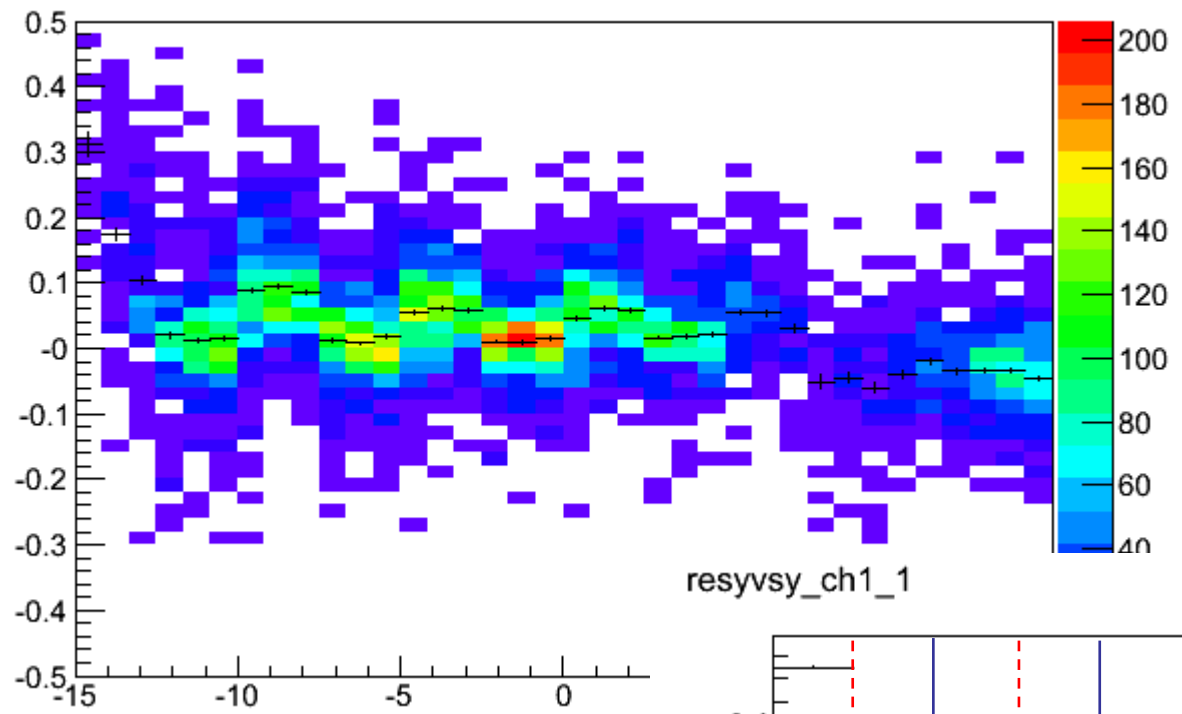




**sx**  
RES +

**dx**  
RES -

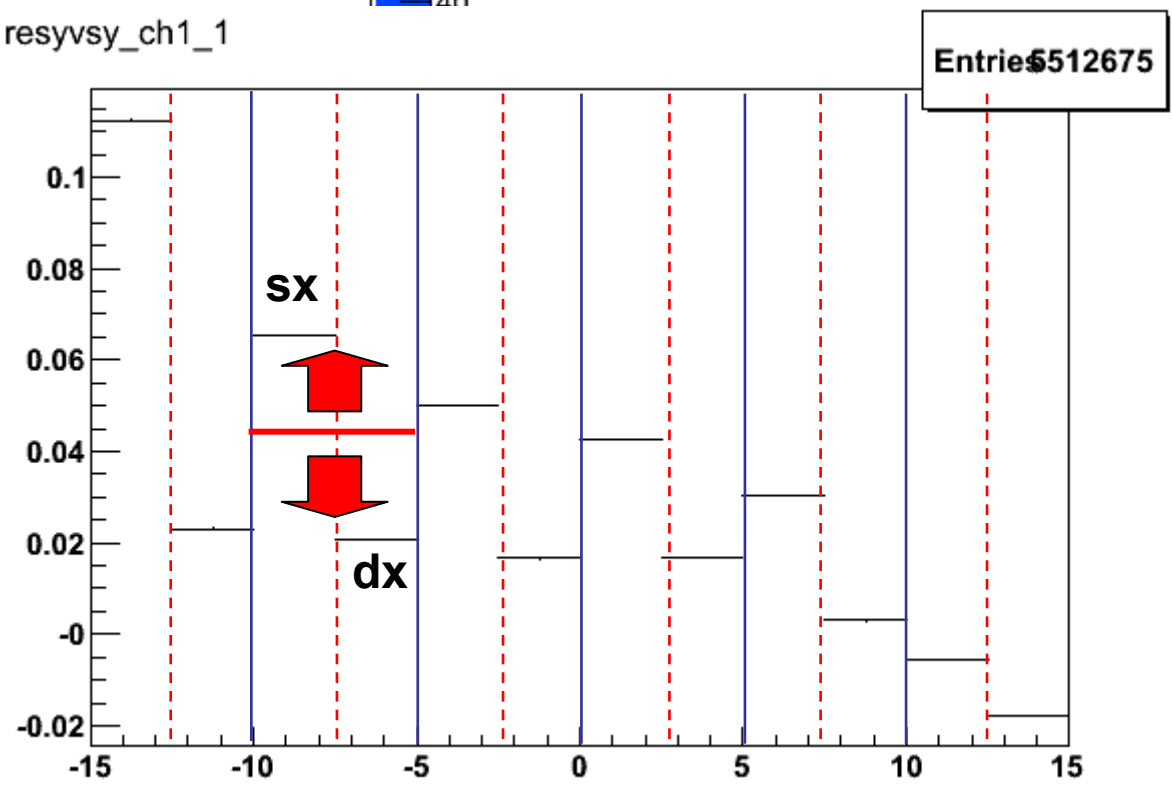
# Evaluation of Garfield correction effect on residual distribution



## Garfield correction effect on residual distribution

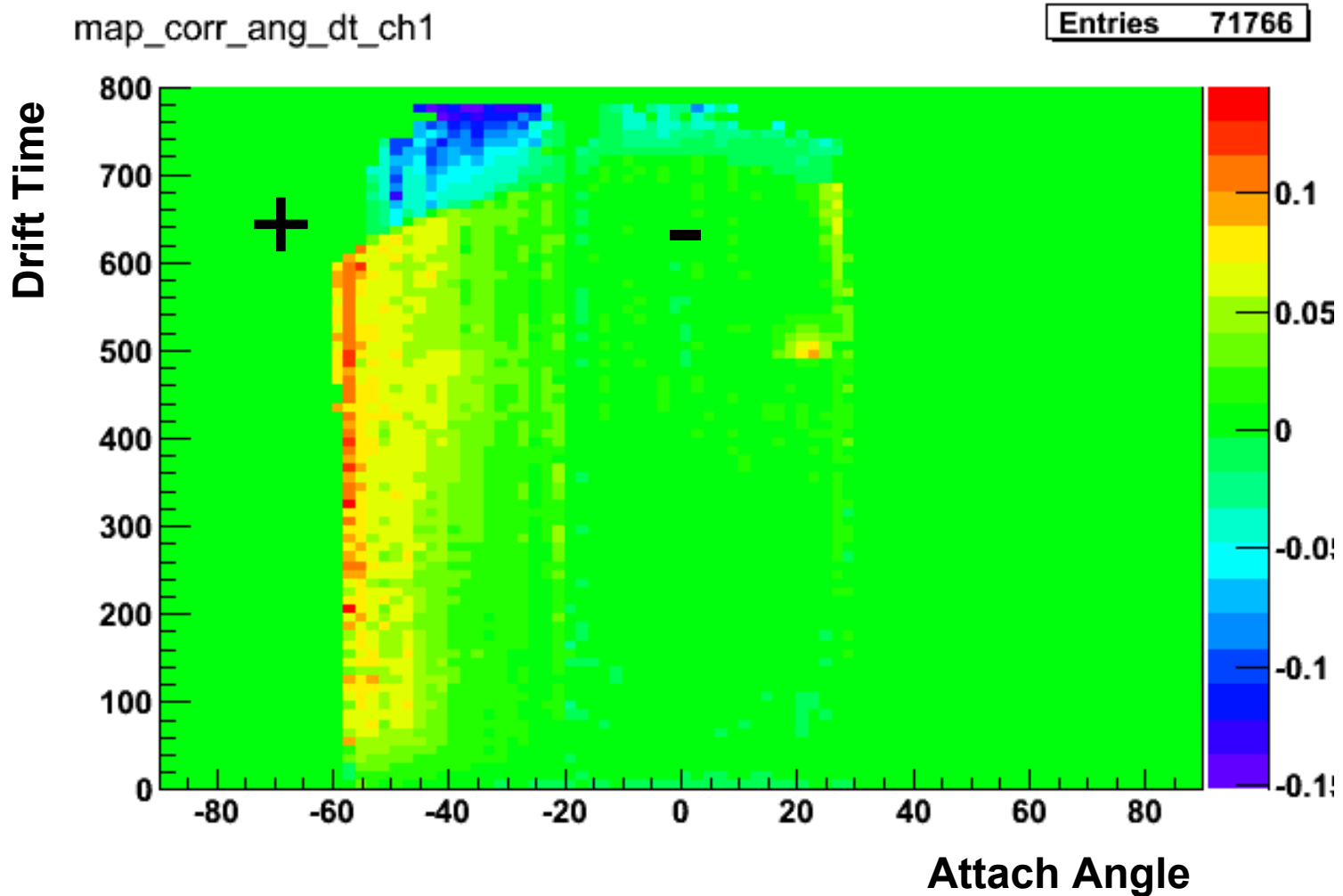
$$\frac{RES(SX) - RES(DX)}{2}$$

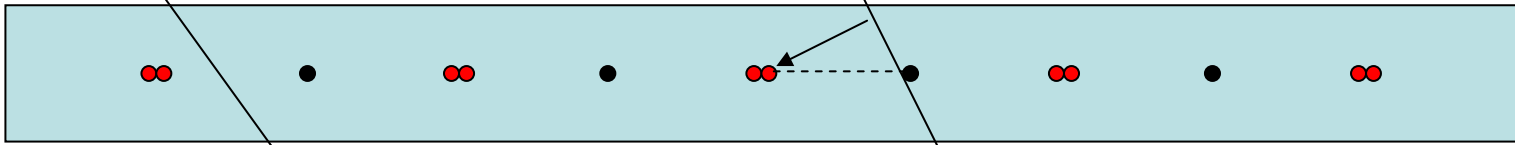
### VS Drift Time and VS Attach Angle For all Chambers



Entries 512675

# Map of Garfield correction effect on residual distribution vs Attach Angle vs Drift Time

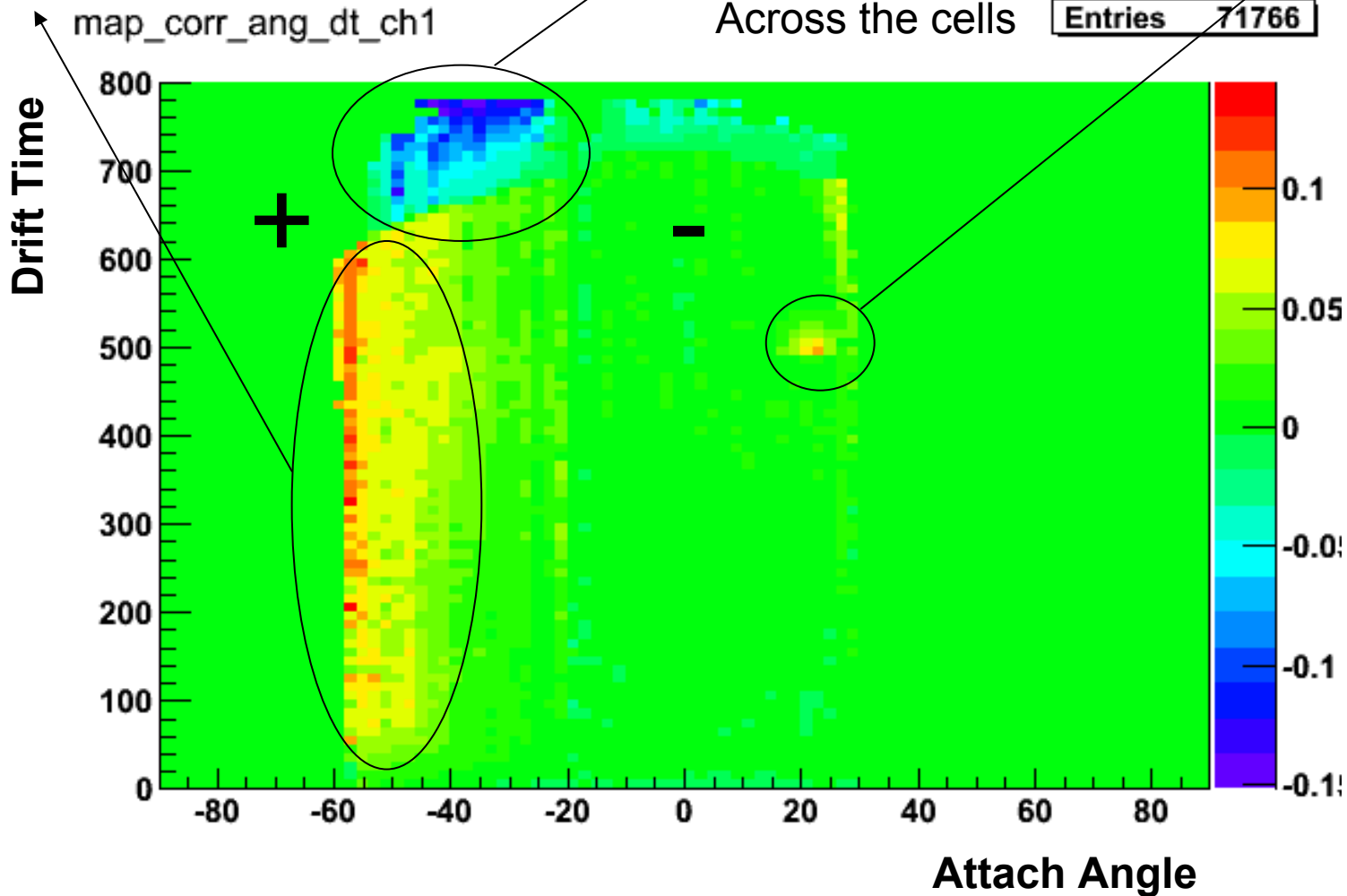




> 45° Garfield  
doesn't work properly

Long drift time =  
Across the cells

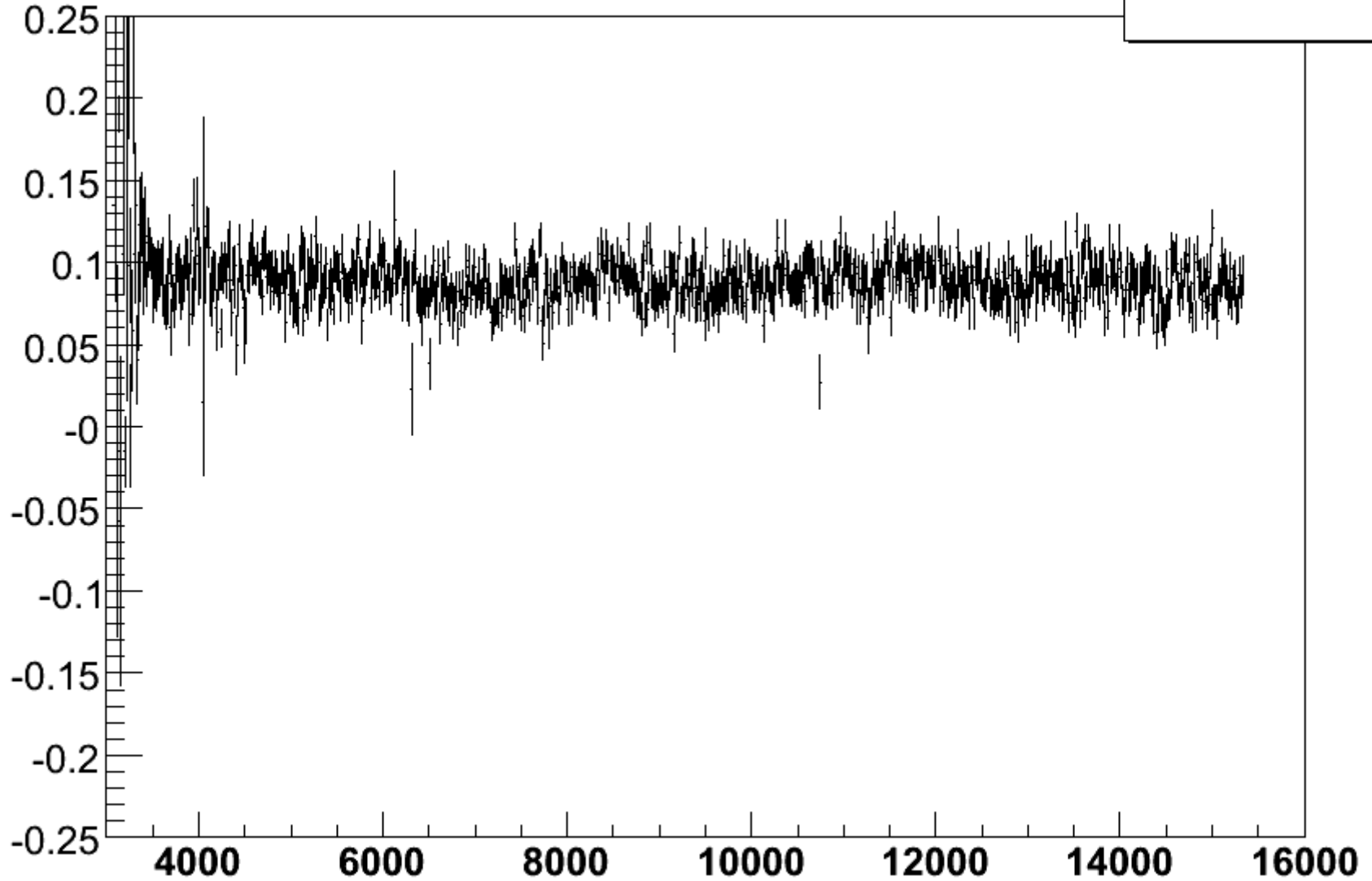
Wrong Garfield  
Table Entry



# Time dependence of $\Phi$ residual wire by wire

runvsresyvsy\_ch1\_0\_w\_00

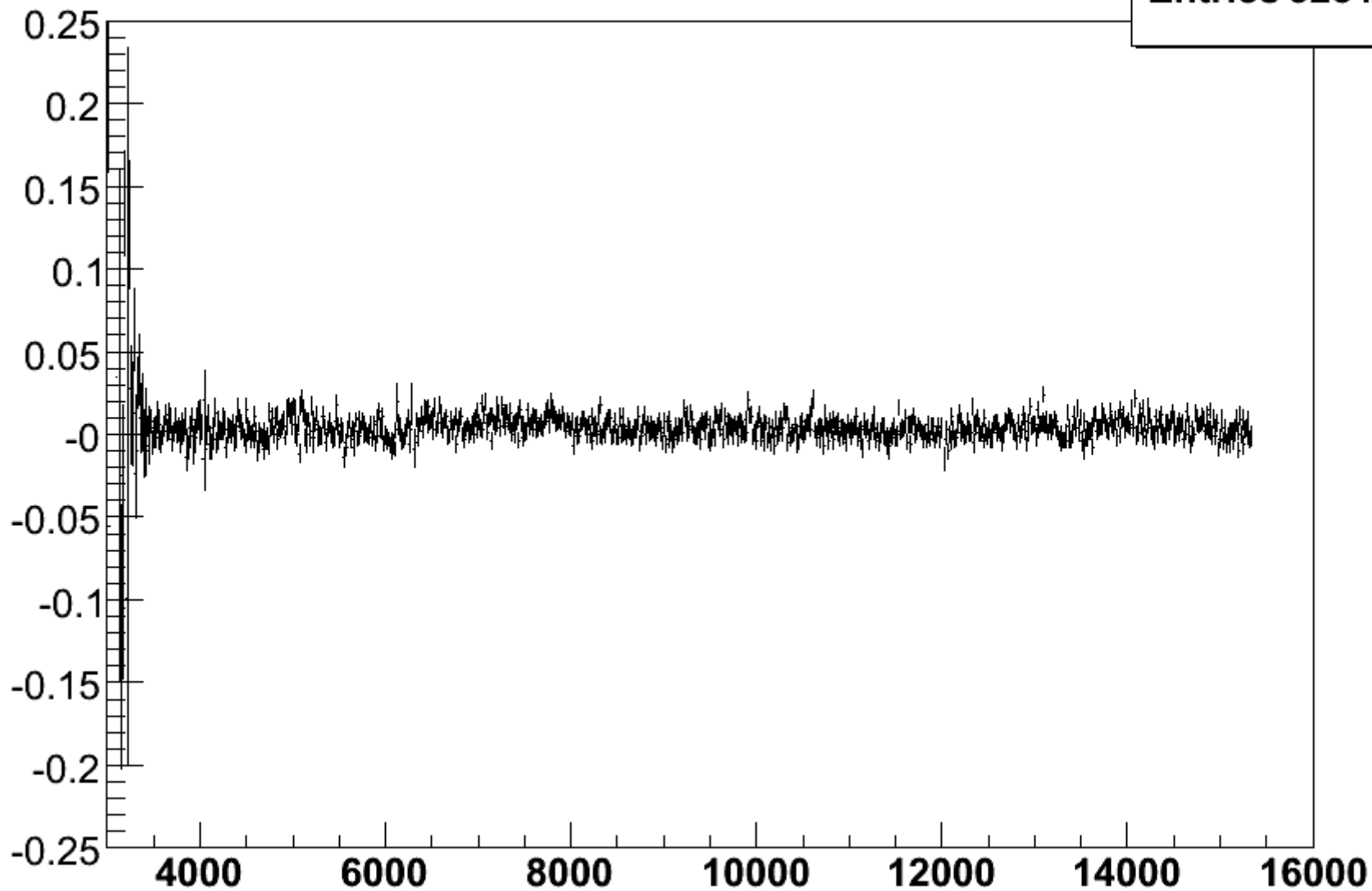
Entries 3871661





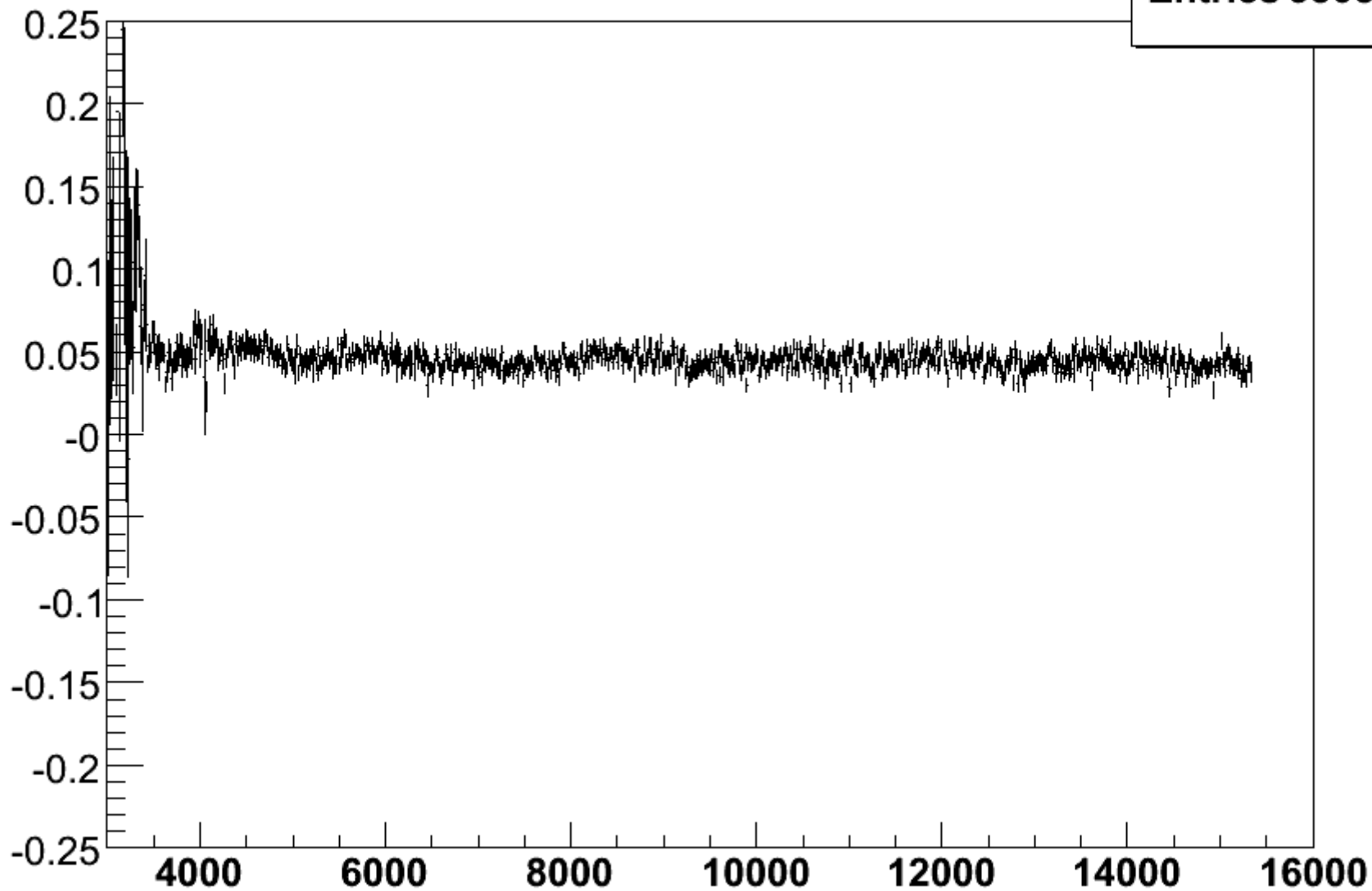
runvsresyvsy\_ch1\_0\_w\_01

**Entries 323168**



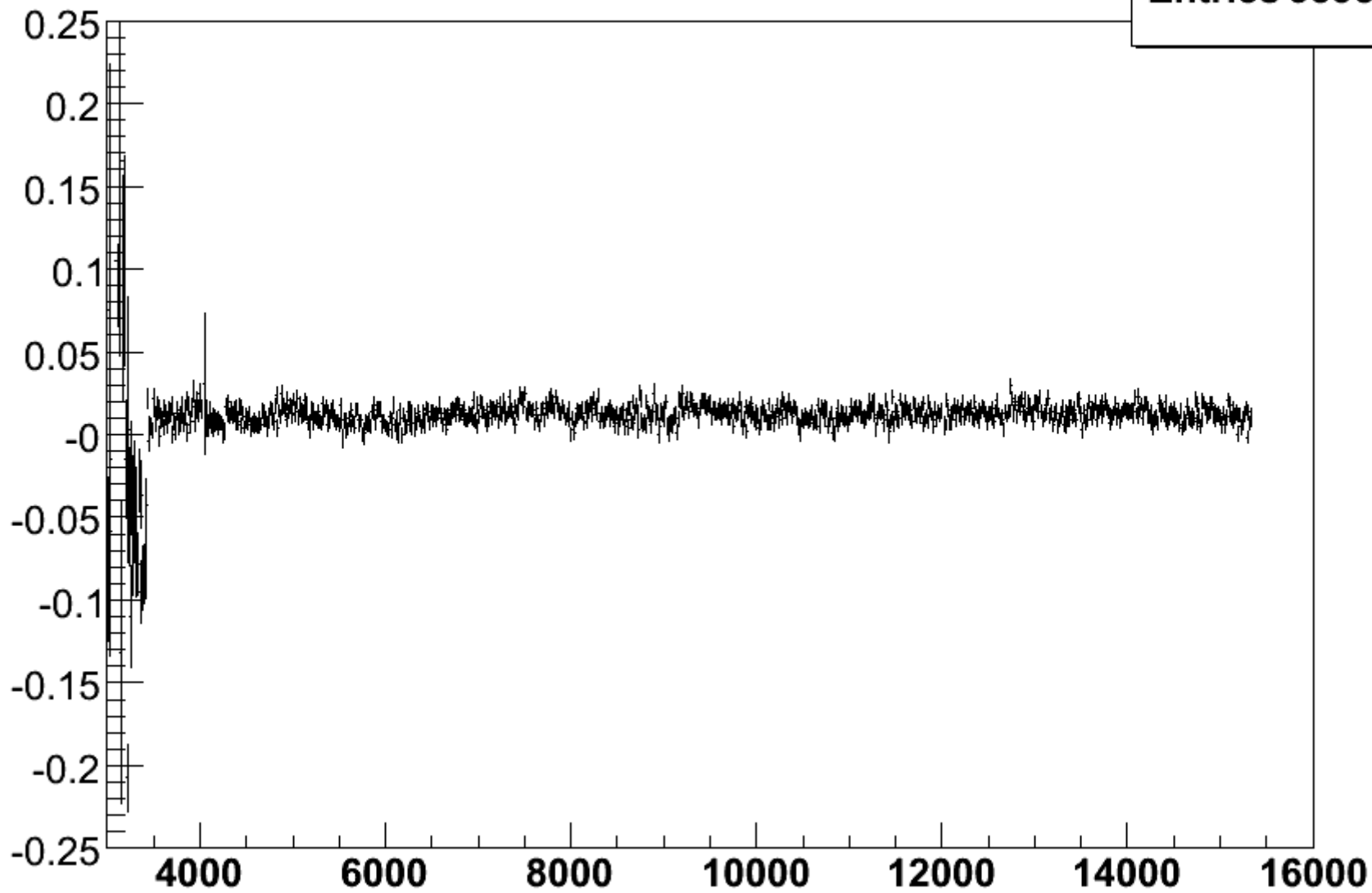
runvsresyvsy\_ch1\_0\_w\_02

Entries 350512



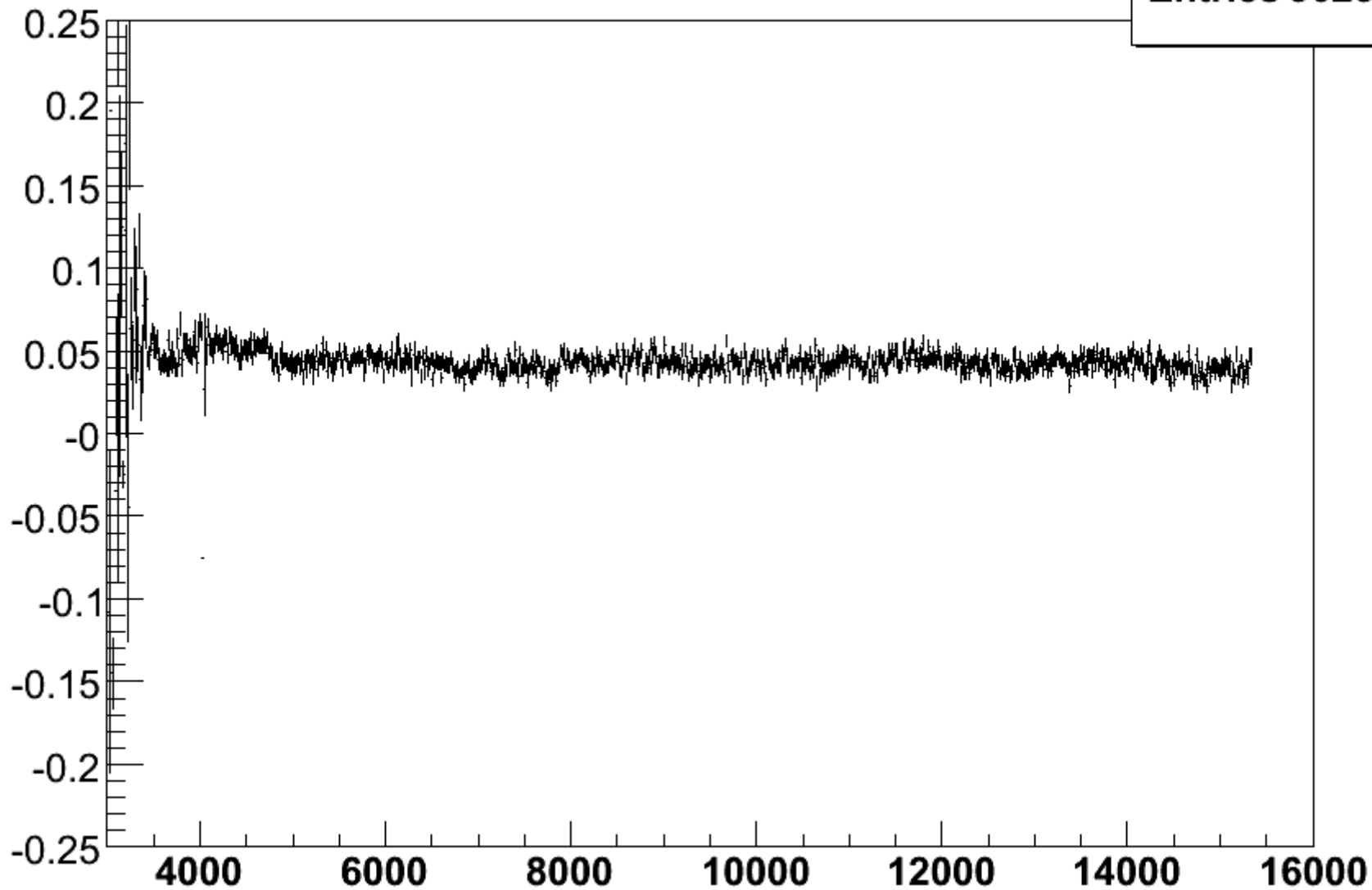
runvsresyvsy\_ch1\_0\_w\_03

Entries 359548



runvsresyvsy\_ch1\_0\_w\_04

Entries 362322

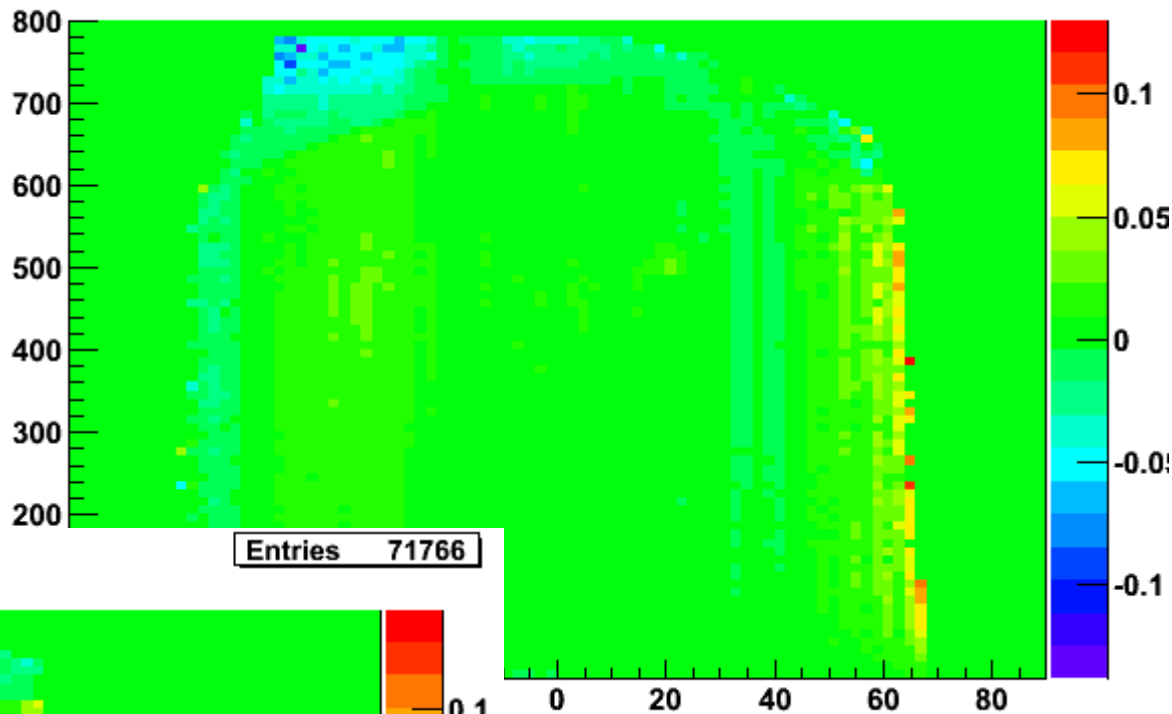


# Drift Chamber $\Phi$ Side ... the TDC SIDE of the moon

- Garfield doesn't work properly for attach angle  $>45^\circ$  (but now we have a correction map to add to Garfield !!!)
- No cluster on hit chamber reconstruction:
  - Wrong evaluation of  $\Phi$  coordinate when the track crosses two cells (long drift time)
- Residual are stable in time but they are different from zero (500  $\mu\text{m}$  – 1 mm)
  - Calibrations
  - Alignment

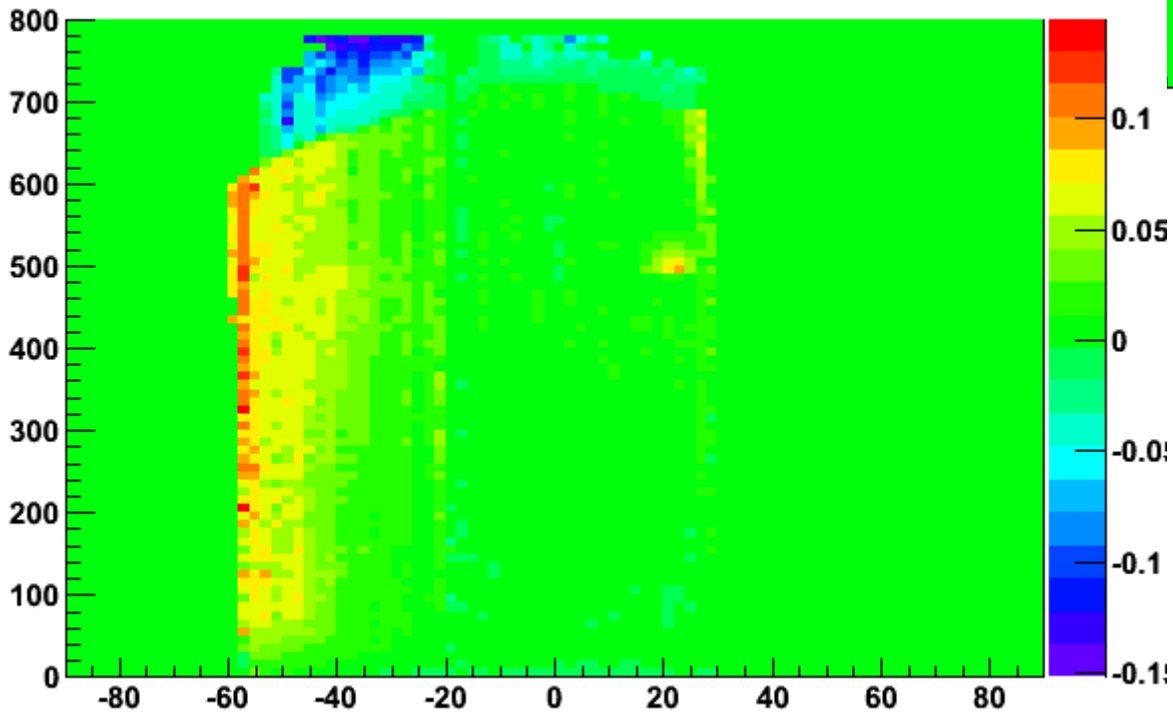
map\_corr\_ang\_dt\_ch1

Entries 152622



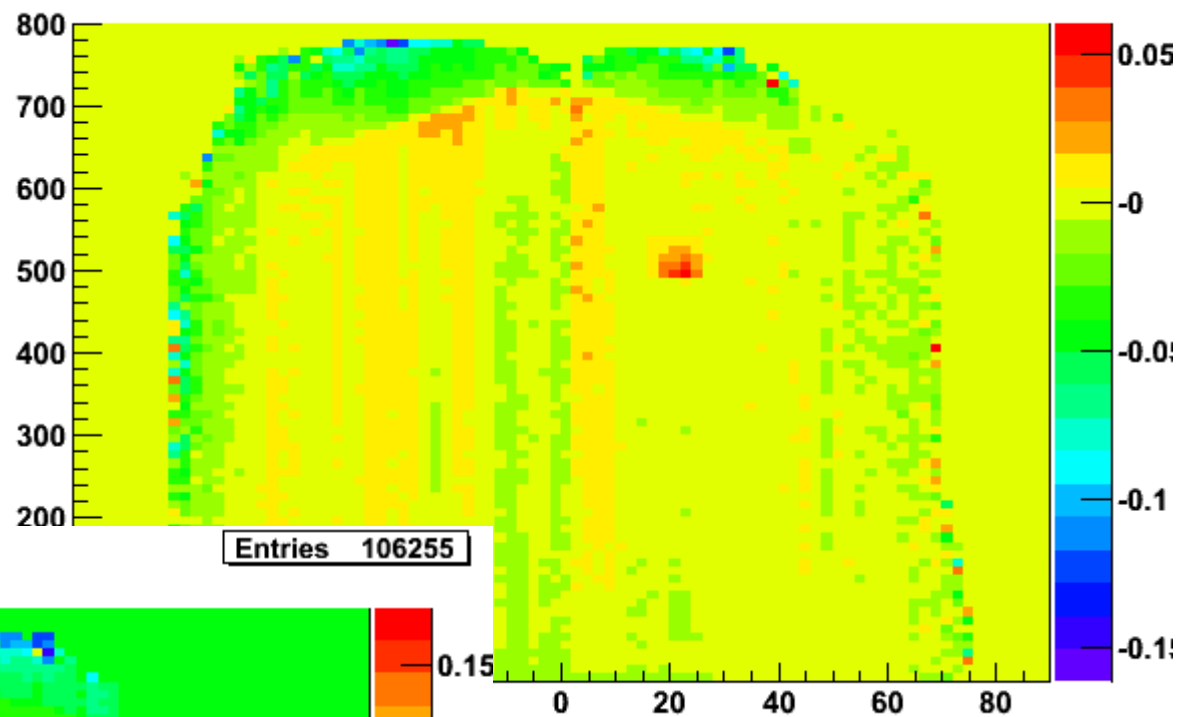
map\_corr\_ang\_dt\_ch1

Entries 71766



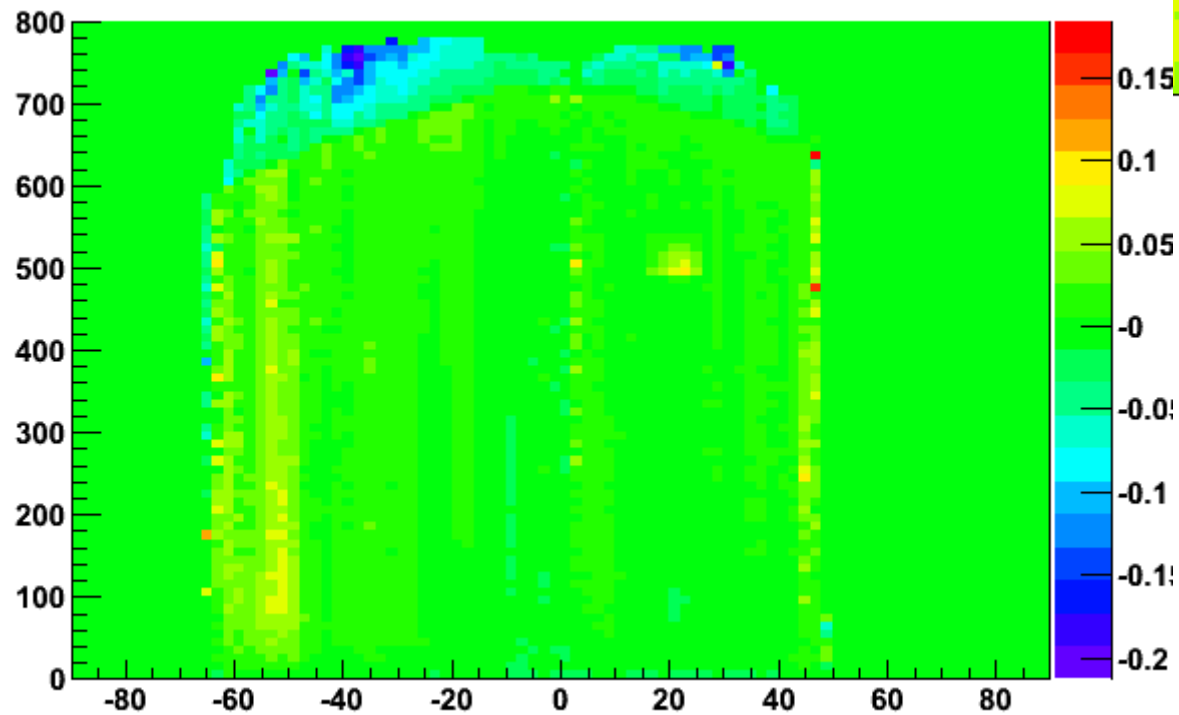
map\_corr\_ang\_dt\_ch2

Entries 222848



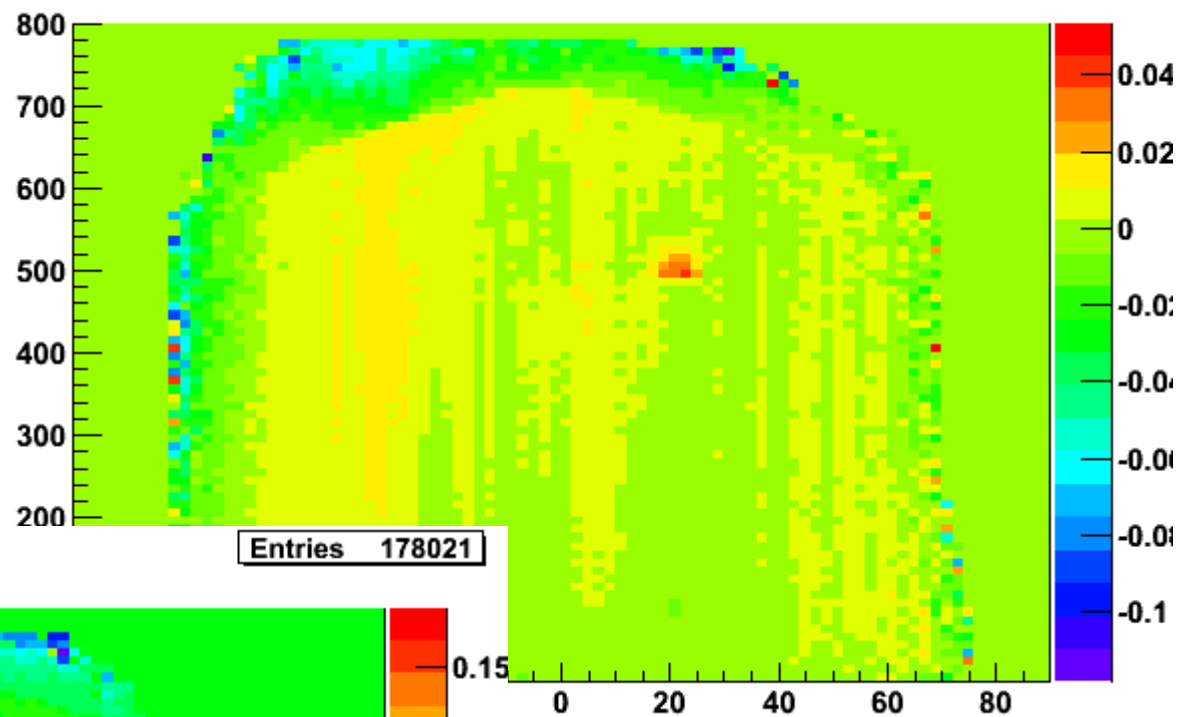
map\_corr\_ang\_dt\_ch2

Entries 106255



map\_corr\_ang\_dt\_chtot

Entries 375470



map\_corr\_ang\_dt\_chtot

Entries 178021

