



Overlap-hits under the microscope

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- ▶ `HIPAlignmentAlgorithm` properly treats the correlation between x and y , assuming that these are both continuous random variables with Gaussian errors.
- ▶ Both x and y distributions have structure: narrow Gaussian peak when the muon passes between strips/wire groups, flat distribution when the muon passes through strips/wire groups.
- ▶ Separation between y wire groups is much greater than our desired alignment precision (1.5 cm in ME+2/1, 5 cm in ME+2/2), so granularity in y could cause serious artifacts.
- ▶ These propagate into x because the strips are angled.

Exploratory study

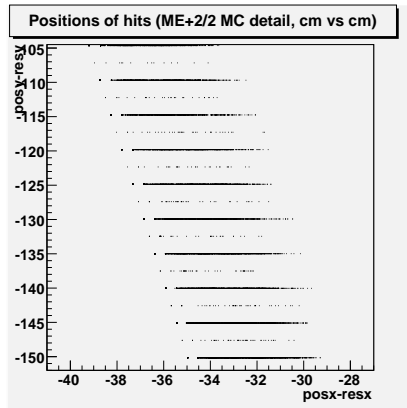
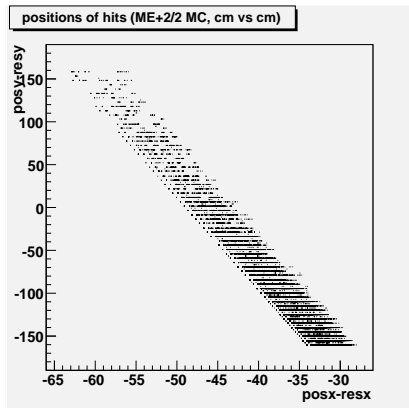
- ▶ Made an ntuple of hit positions and residuals; looked at what we're starting with

Distribution of hits

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- ▶ Edge of trapezoid is clearly visible
- ▶ As are the wire groups in y
- ▶ I don't understand the gap on the far left edge or the small half-integer peaks
- ▶ This isn't discovery: CSCRecHits are constructed to have this distribution
- ▶ ME+2/2 has 5 cm between wire group centers

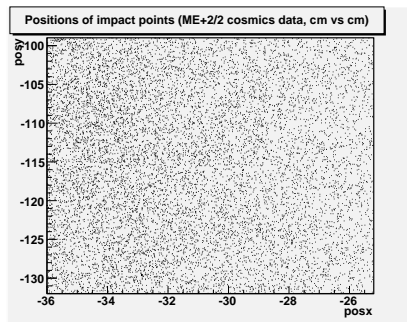
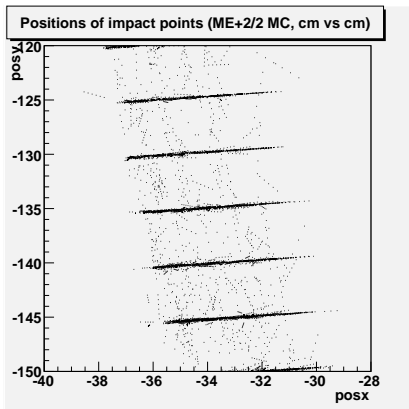


Distribution of impact points

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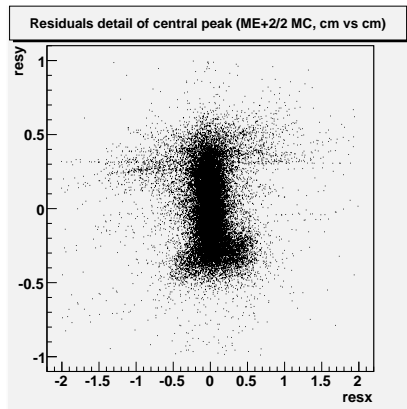
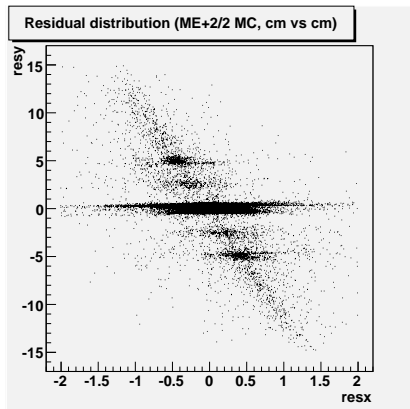


- ▶ Propagate tracks from reference chamber, plot them in local coordinates of target chamber
- ▶ Clearly see the angle between the two chambers
- ▶ Some broadening because tracks are not perfectly parallel
- ▶ In *cosmics* data, complete smearing because tracks are nowhere near parallel





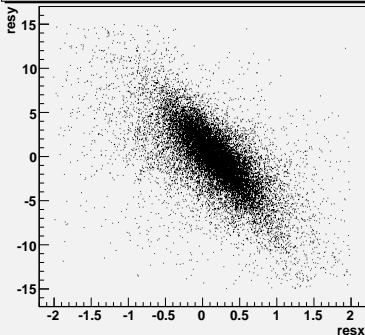
- ▶ 2-D residual distribution: expect
 - ▶ correlation between x and y because x can only be determined from strip if y is known
 - ▶ peaks in y corresponding to $\pm 0, \pm 1, \pm 2 \dots$ differences in wire groups from reference to target
- ▶ Observed both features: slope is 0.1, central peak has vast majority of the events
- ▶ Inner $|r_y| < 2$ mm is featureless



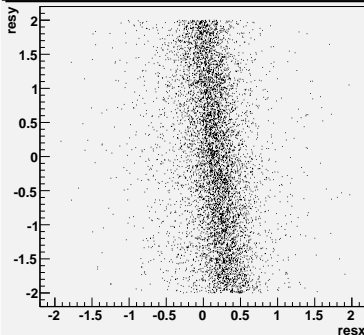


- ▶ Look at the same distributions in CRUZET data
- ▶ Same correlation, but structure is entirely washed out
- ▶ More to do with the fact that these are cosemics than the fact that these are data
- ▶ (Yes, we only look at one chamber at a time in data. This one has the most hits.)

Residuals in ME+2/2 chamber 25 (data, cm vs cm)

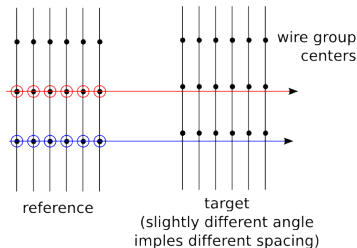


Residuals in ME+2/2 chamber 25 detail (data, cm vs cm)

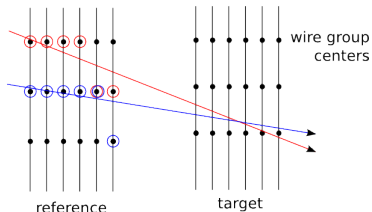


Why are cosmics smeared?

Side view (beam-halo):



Side view (cosmics):



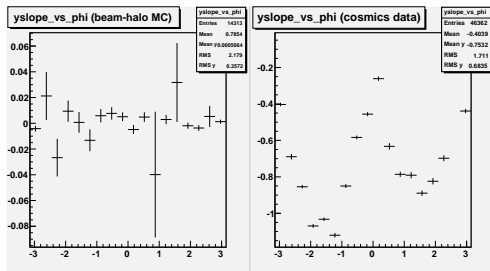
More continuous distribution of extrapolated tracks

Quantify angles of tracks with local $\partial y/\partial z$

Close to zero in beam-halo MC

Clear shape in cosmic ray data

- ▶ cosmic rays steeply angled relative to beamline
- ▶ preference for negative $\partial y/\partial z$, points to interaction point (ALCT patterns)
- ▶ slope of 1 means a new wire group every two layers

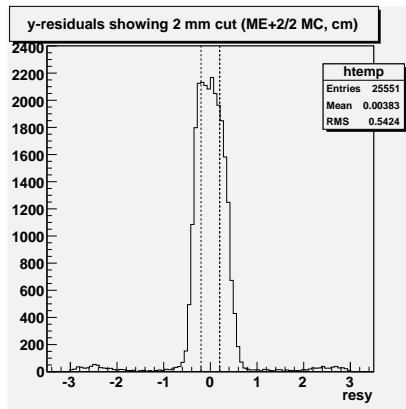
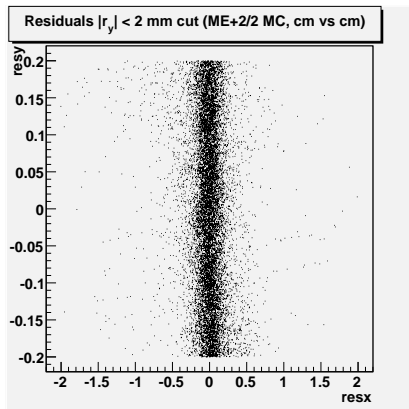


Granularity in beam-halo (1/2)

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- ▶ Can deal with granularity in the beam-halo ME+2/2 case (biggest separation between peaks) by selecting events in good region of the $\Delta y = 0$ peak
- ▶ In this region, we have a flat probability distribution in y
- ▶ (Not as important right now as the cosmic-ray case)

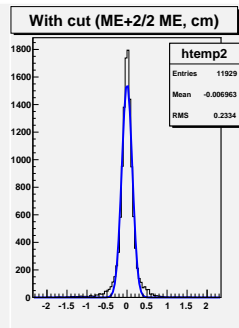
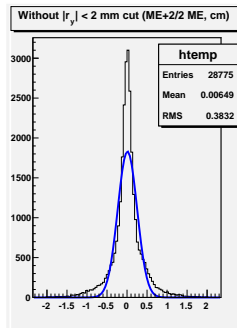
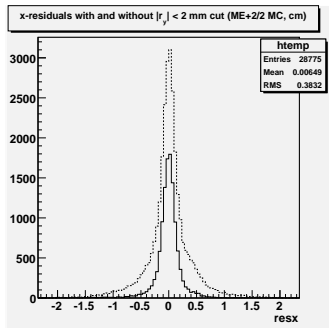
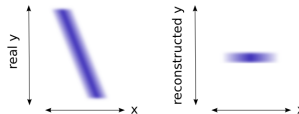


Granularity in beam-halo (2/2)

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- ▶ Lose a factor of 2, but x residuals become cleaner, better understood
- ▶ Resolution: (half y -gap) \times slope \oplus intrinsic = $\frac{5 \text{ cm}}{2} \times 0.1 \oplus 300 \mu\text{m} = 2.5 \text{ mm}$
- ▶ Can't do better than this in x because
 - ▶ we have no knowledge about y except that it's closest to this wire-group center
 - ▶ determining x from the strip measurement is a linear function of y
 - ▶ hmmm... why isn't the x residual distribution a rounded box?



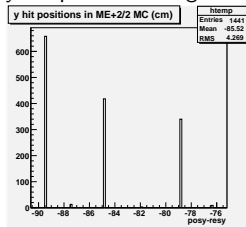
Inner ring vs. outer ring

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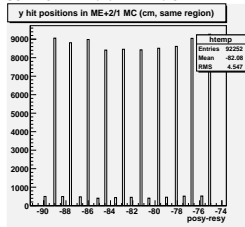


- ▶ Wire groups are 1.6 cm apart in ME+2/1, as opposed to 5 cm
- ▶ Smaller spacing washes out structure; harder to see ± 0 , $\pm 1 \dots$, peaks
- ▶ Slope in x - y correlation is 0.35, consistent with 20° trapezoidal chamber design

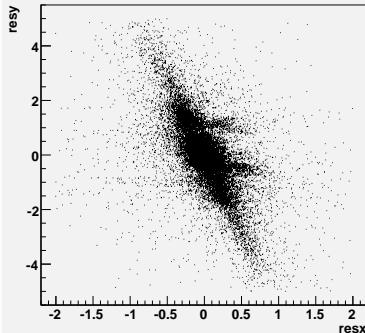
y hit position histograms



same window width



Residual distribution (ME+2/1 MC, cm vs cm)

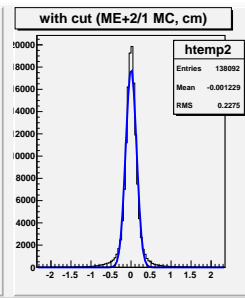
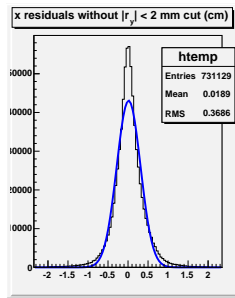
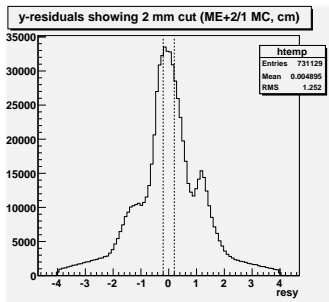


Same trick in inner ring

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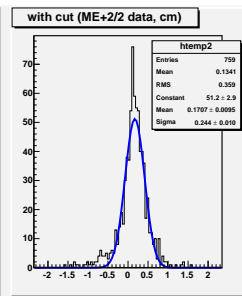
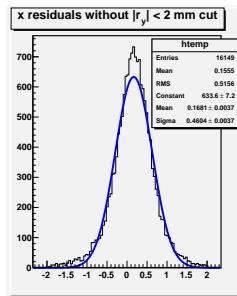
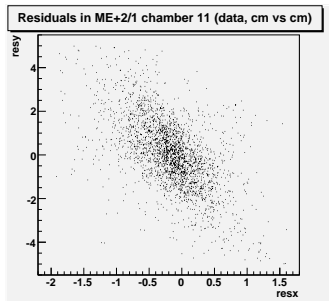


- ▶ Much greater loss in statistics due to greater probability of $\Delta y = 1$ or 2
- ▶ Only a marginal benefit, but final resolution is the same:
 $(\text{half } y\text{-gap}) \times \text{slope} = \frac{1.6 \text{ cm}}{2} 0.35 = 2.5 \text{ mm}$ (by design?)
- ▶ When y is very granular, it pays to take advantage of this and select the clean central region, and if y were completely continuous, we wouldn't need to do anything special. But what about the middle case?



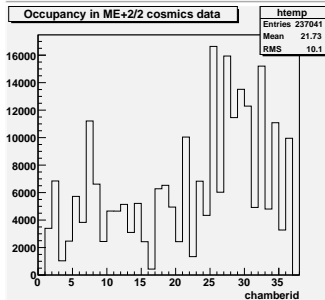
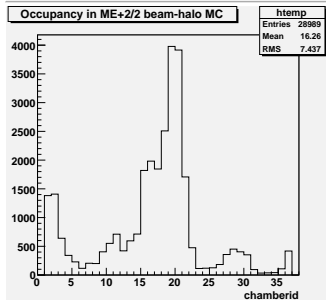
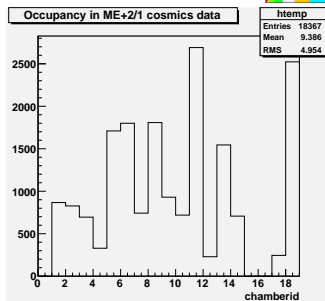
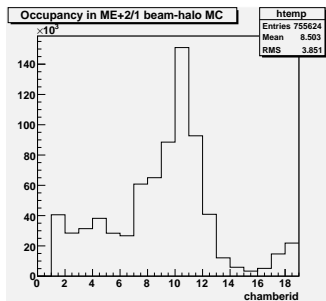


- ▶ Like I said, it's the fact that they're cosmic rays with non-horizontal impact angles (and the fact that there's a spread in impact angles can only help)
- ▶ Cutting central region trick does very little good here
- ▶ The distribution looks smooth: do you suppose it really is smooth? Can we just assume that the Gaussian correlations-handling in HIPAlignmentAlgorithm will take care of it?
- ▶ Layer misalignments can also smooth the y residuals, but 6 is a rather small number for such featurelessness
- ▶ Alignment artifacts we see in beam-halo MC will not necessarily affect cosmic data, so the MC is not a very helpful guide



Occupancies: not very flat

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- ▶ No sign of structure in cosmic ray residuals: they're just broad, as would be expected from granularity plus large uncertainty as to which wire group you're on
- ▶ Maybe we can just call the cosmic ray case Gaussian and move on? (That is, interpret any failure to close as a *real* radial displacement of all chambers. . .)
- ▶ We'll need to specially handle granularity effects with beam-halo, but not yet
- ▶ Inner ring (ME+2/1), being intermediate between “large gaps” and “nearly continuous,” could be a difficult case
- ▶ No need to modify the code now if we don't worry about granularity in cosmic rays
- ▶ We should not use beam-halo MC as a predictor of cosmic ray data!
- ▶ Running a one-step-at-a-time alignment from ME+2/2 chamber 34 in both directions to chamber 16 (which has the least statistics) to check closure in the simplest algorithm