

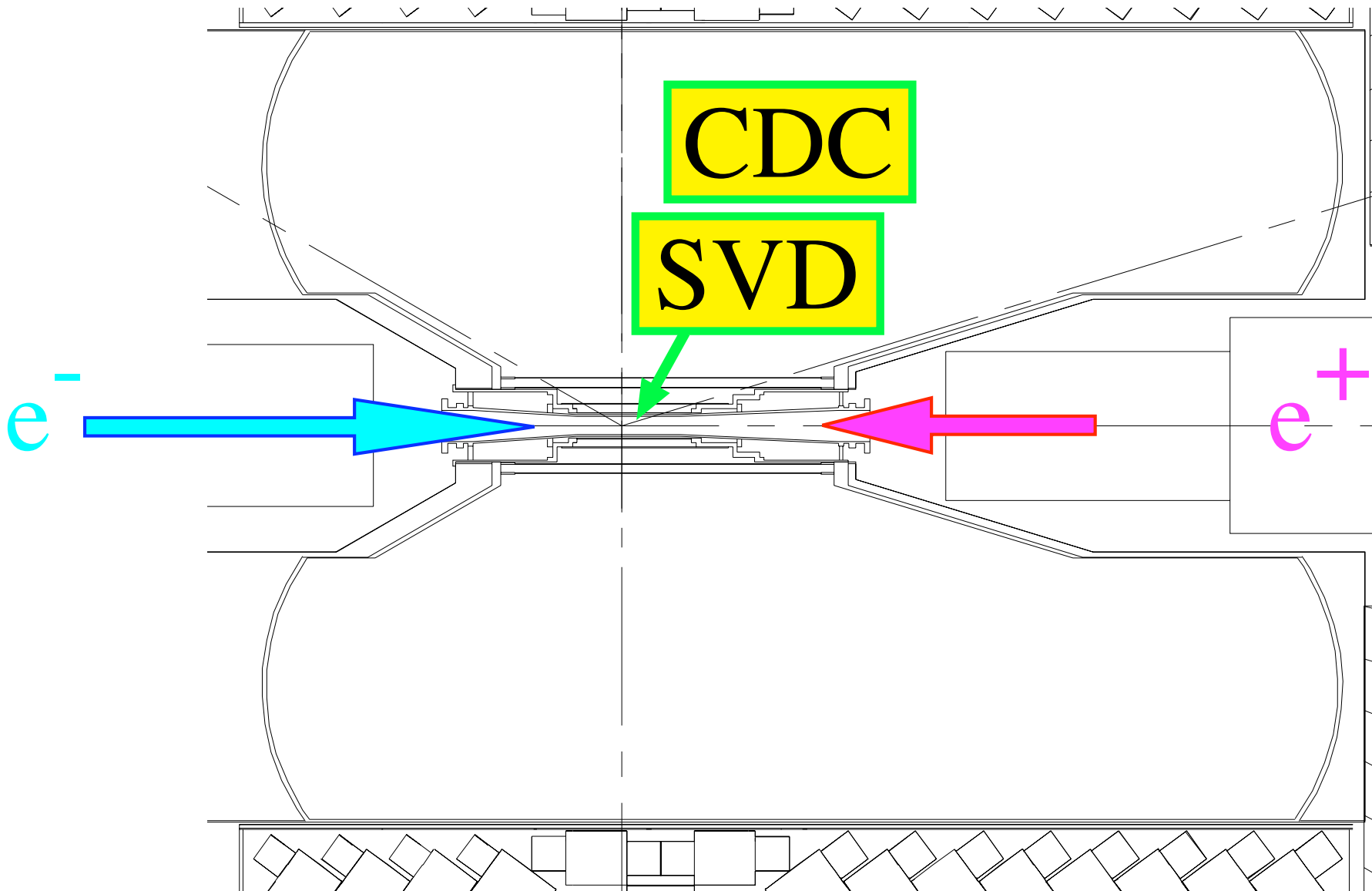
New shape dependent clustering algorithm

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Introduction

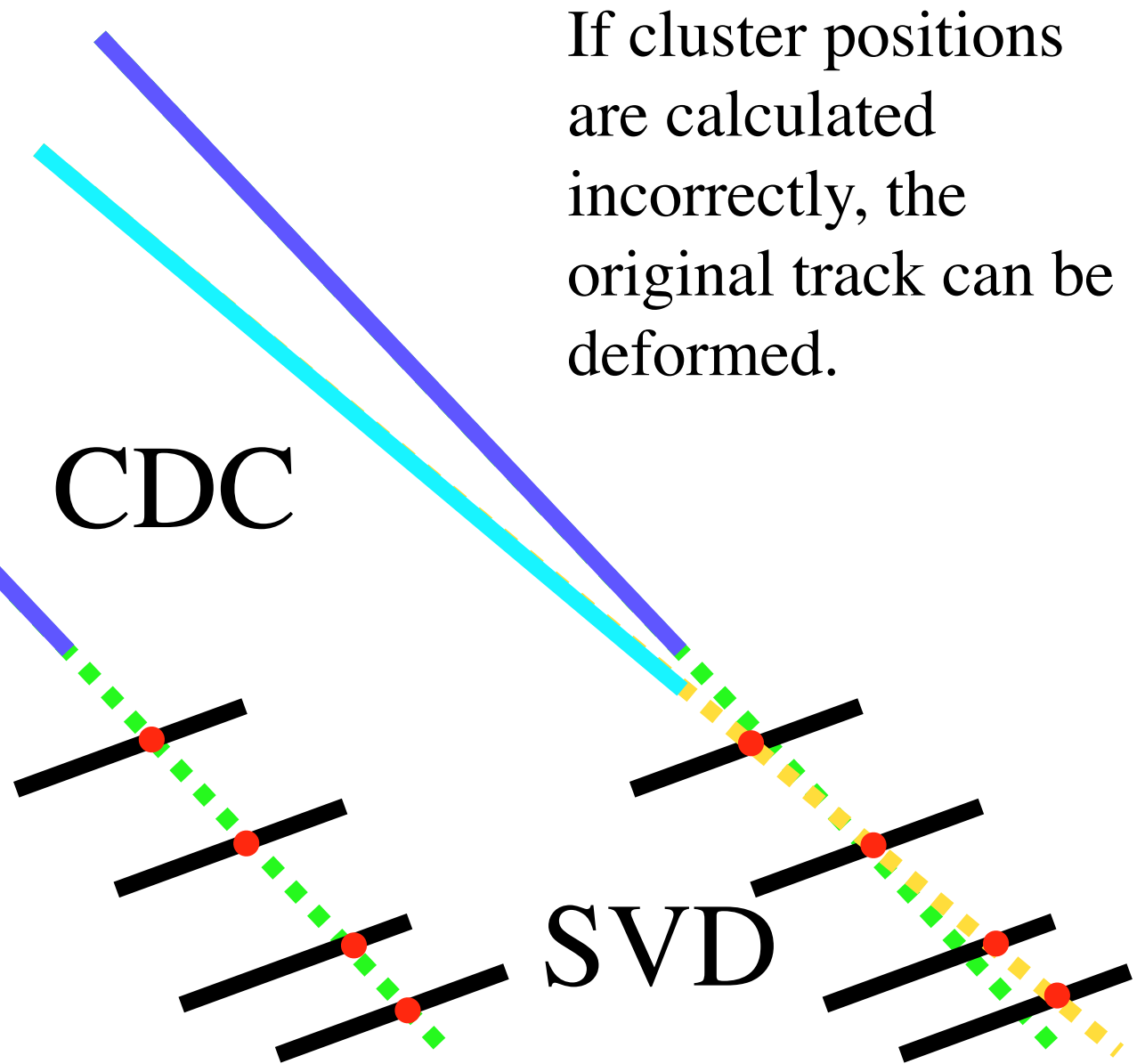
Drift chamber and vertex detector



Tracking is performed in the multiwire chamber (CDC). Silicon vertex detector (SVD) is situated inside this tracking chamber. Outer detectors are used for calorimetry and particle species identification.

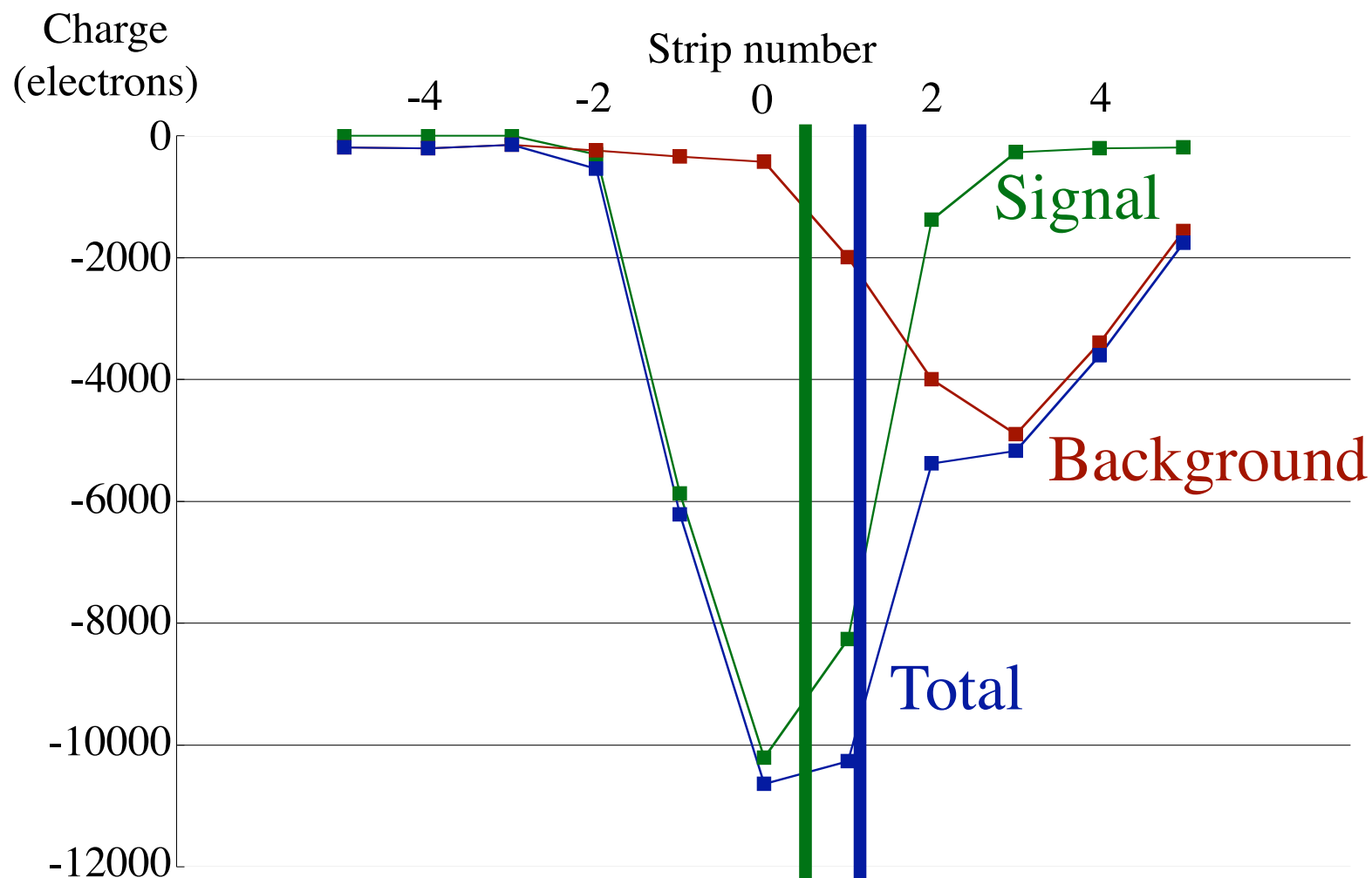
Tracking

1. Extrapolate track from CDC into SVD.
2. Look for nearby clusters in SVD.
3. Recalculate track parameters using new information from SVD.

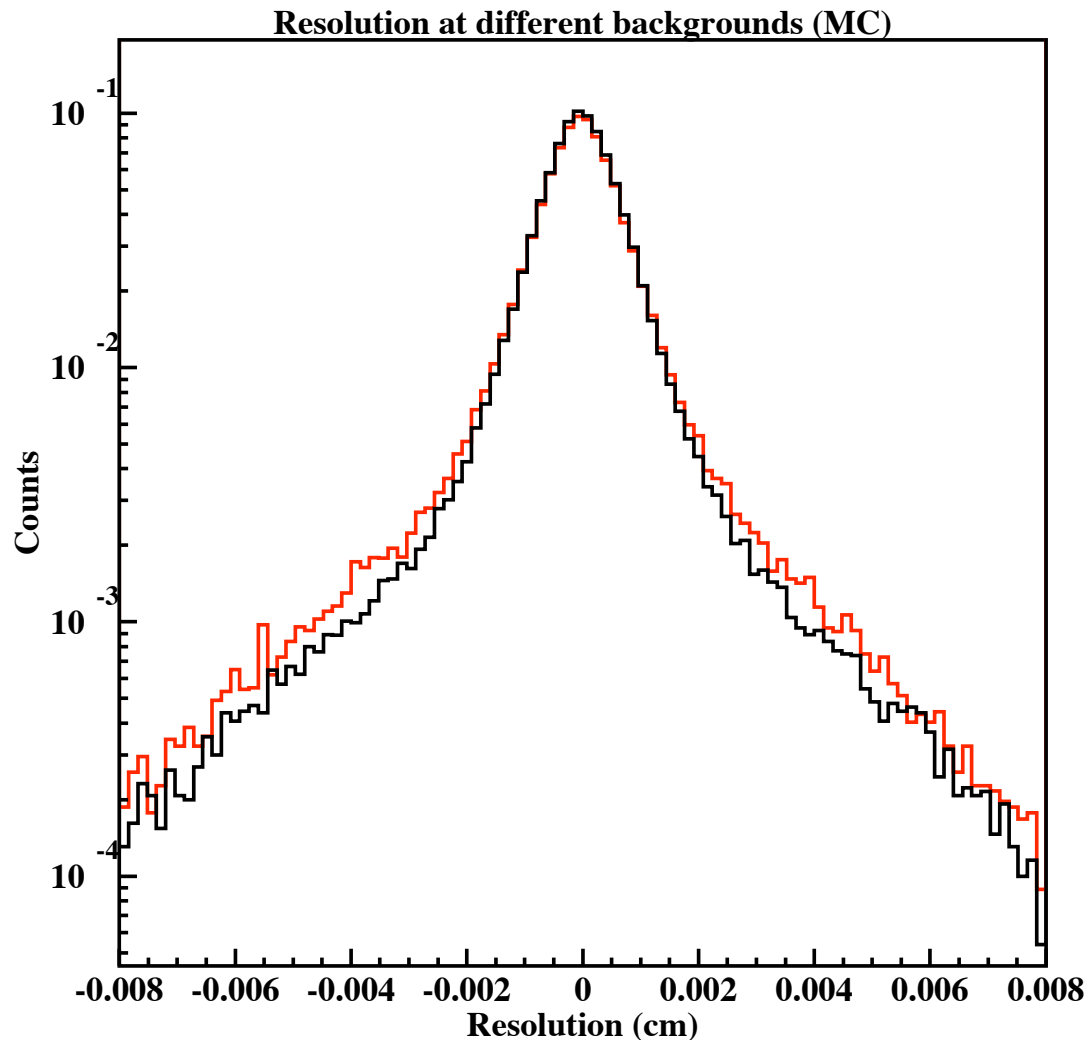


Cause of tracking degradation

1. Incorrect cluster used in tracking.
2. Cluster position is calculated incorrectly due to merging of true hit with background hit.



Merged clusters



normal background (x1)

higher background (x3)

Under higher background the some clusters are significantly deformed because of merging with the background.

Position of merged clusters can not be calculated correctly and results in degradation of SVD clustering resolution.

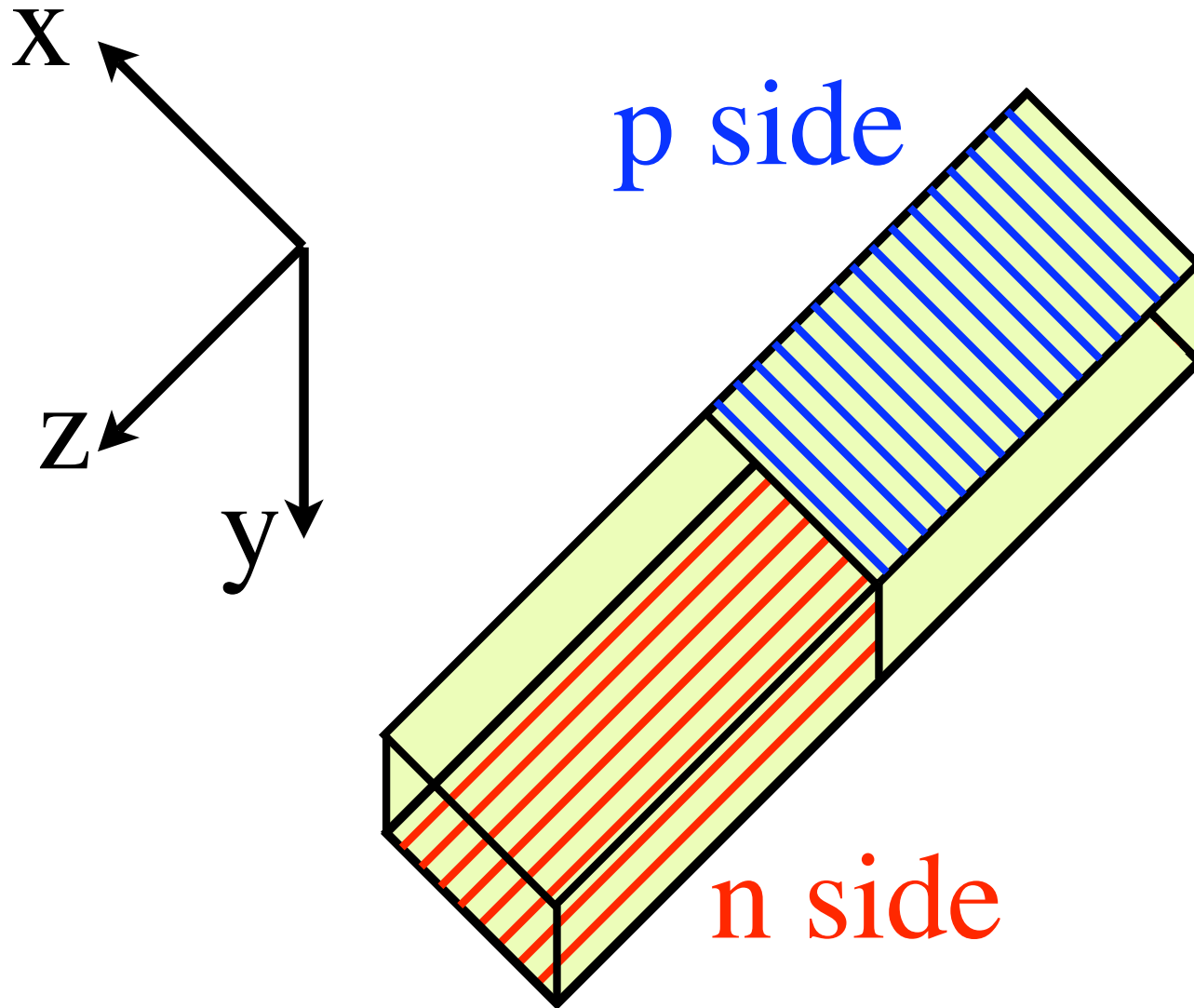
Aim

In the future, higher background will result in a degradation of clustering resolution.

- Aim:
1. Determine behaviour of cluster shape.
 2. Create a new clustering algorithm that takes differences in cluster shape into account.

Inter-strip Charge sharing

Local DSSD coordinates

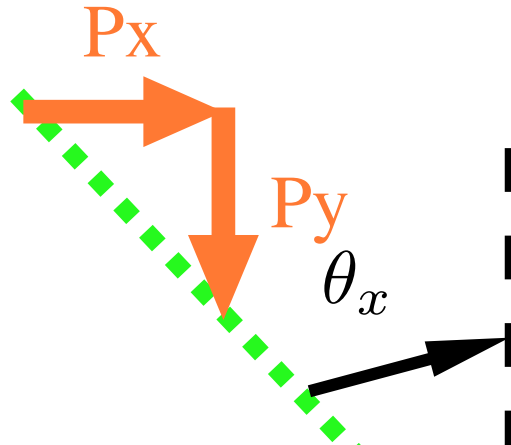


Latitudinal (p-side) and longitudinal (n-side) readout strips. Strips on each side are parallel. P-side and n-side readout wires are orthogonal.

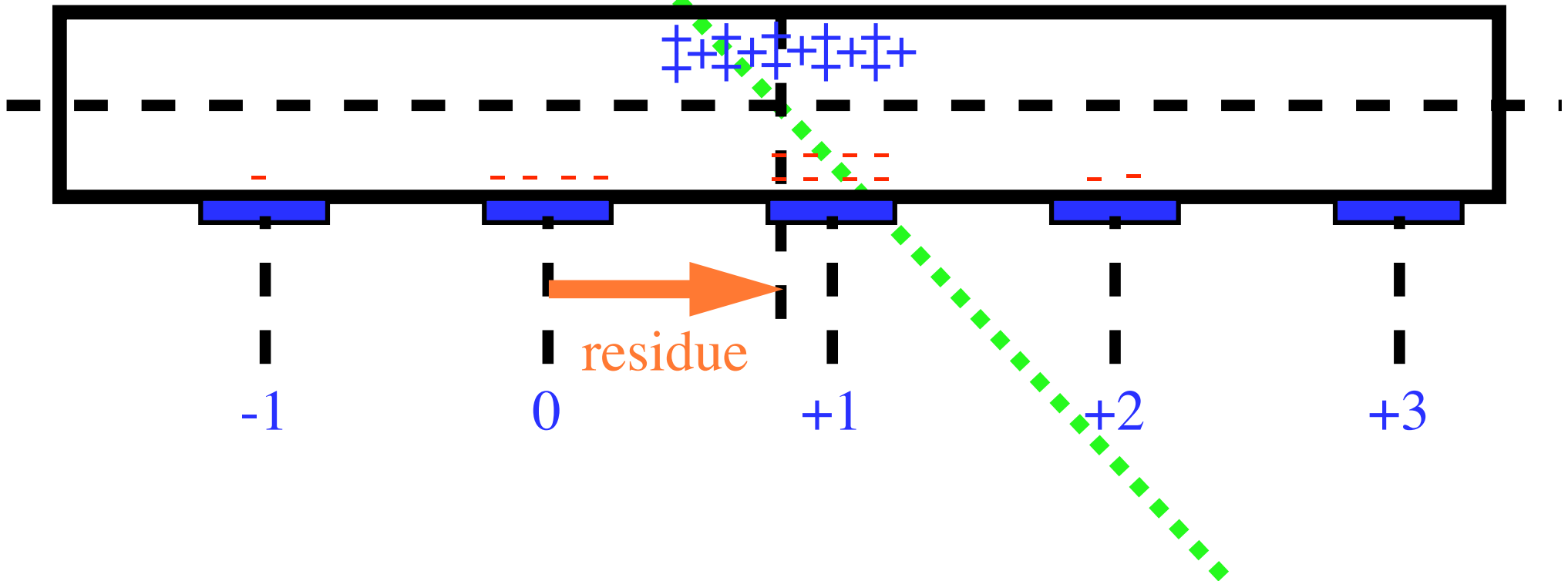
Definition of “incident angle” and “residual”

Extrapolate from
CDC to SVD :

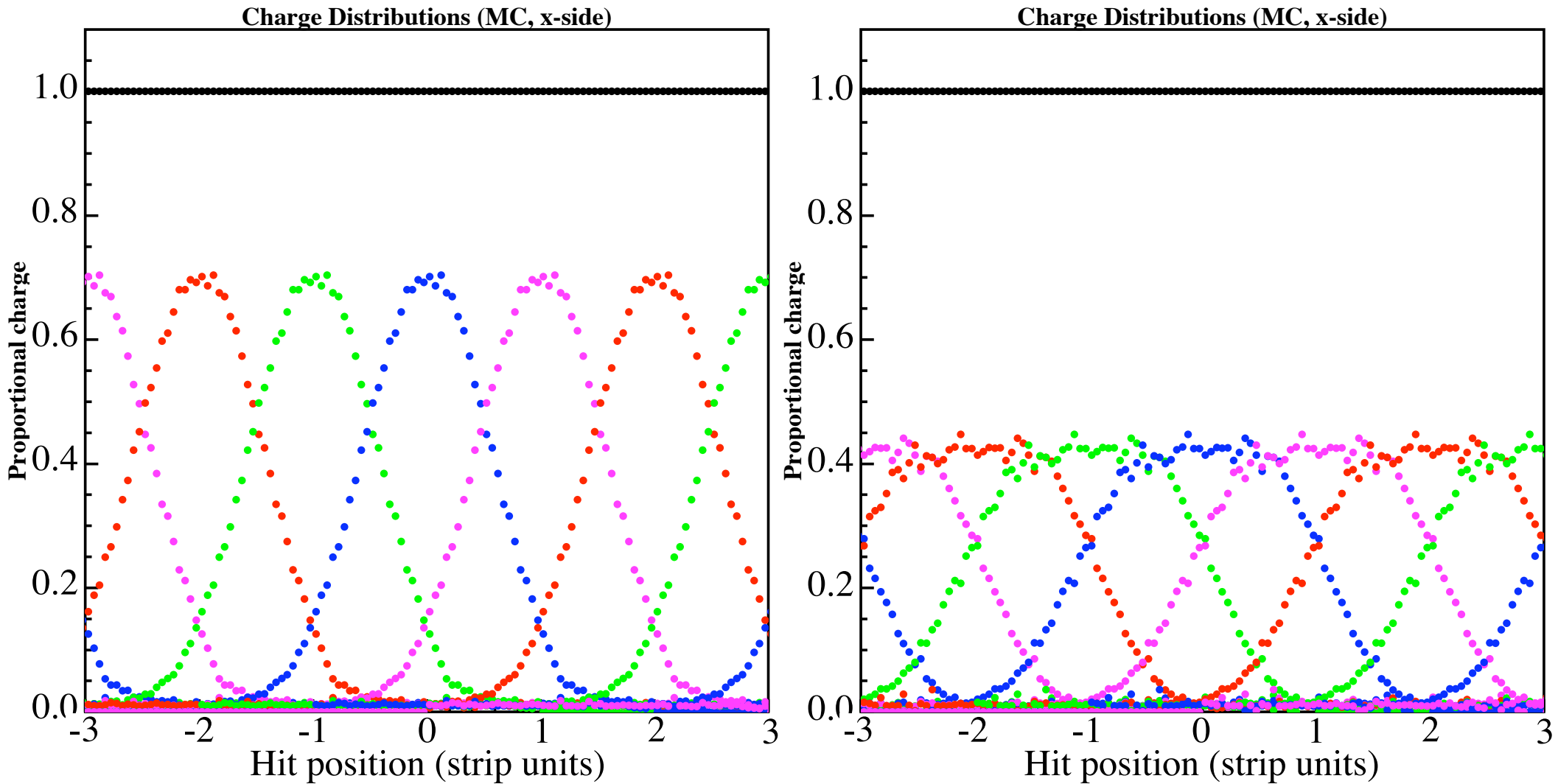
charge sharing behavior depends
on residual as well as incident
angle



define total charge as charge on
strips #-2 ~ #3



charge sharing distributions (MC)



$0 < \text{angle} < 5$

sum
strip# = -2,+2
strip# = -1,+3
strip# = 0,4
strip# = -3,+2

$20 < \text{angle} < 25$

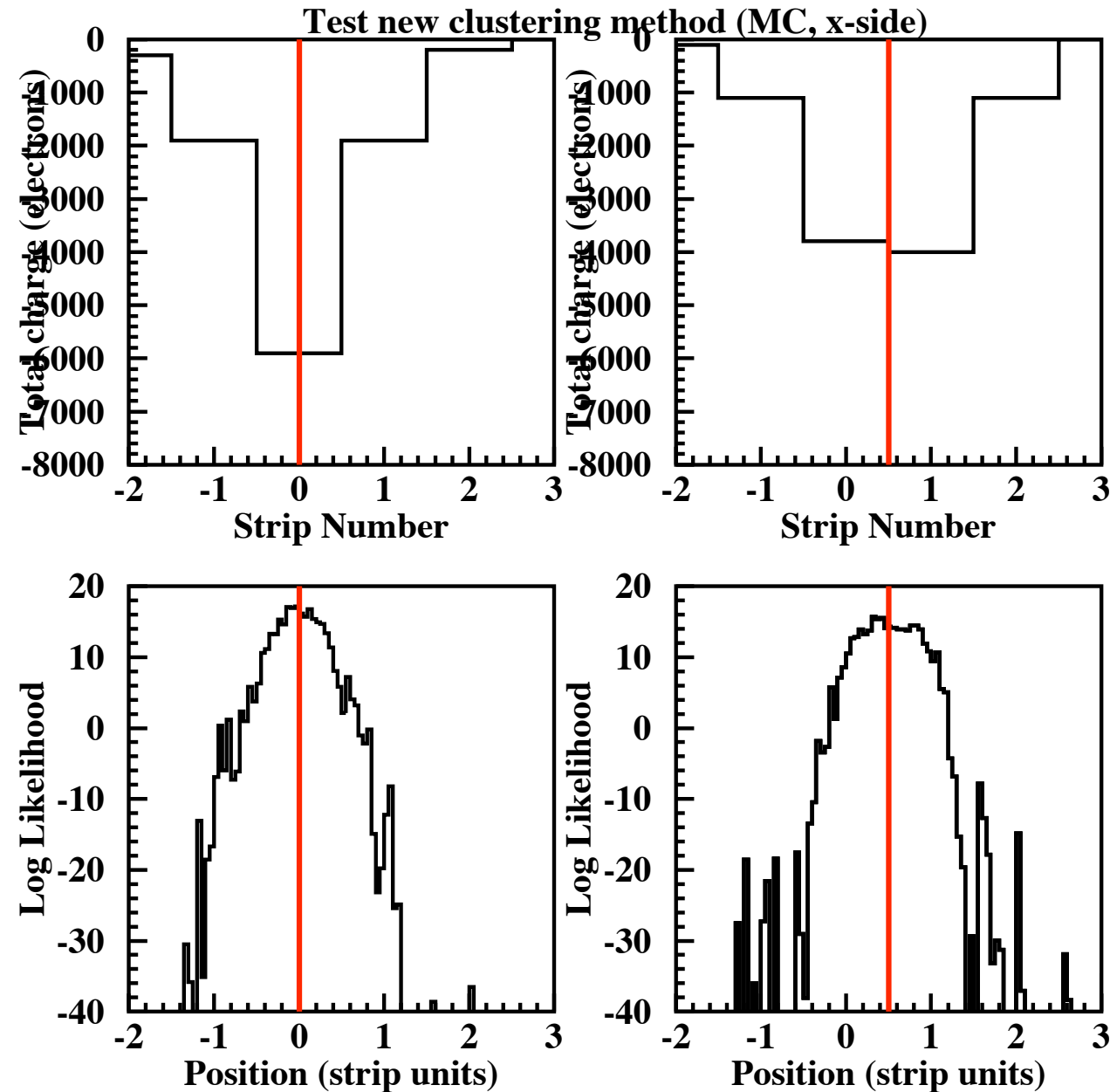
Interaction point resolution

Clear differences in the charge sharing behavior (cluster shape) can be seen for different residuals and incident angles.

Shallower incident angles produce wider clusters.

New Clustering Algorithm

new clustering method test

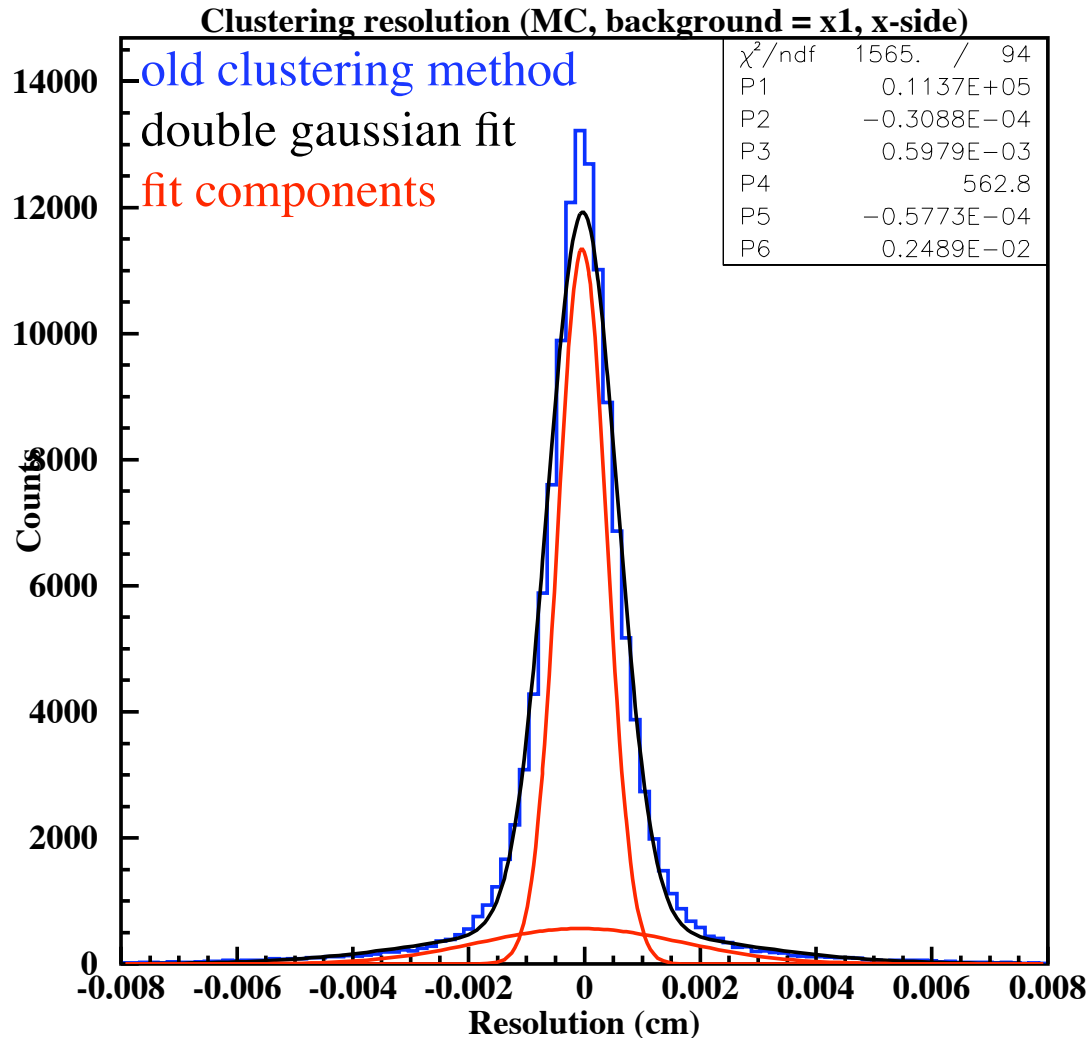


The new clustering algorithm was run on ideal clusters produced from the values in the charge distributions.

The resulting likelihood functions are peaked very close to the expected residual positions (red lines).

Results

Clustering position resolution 2

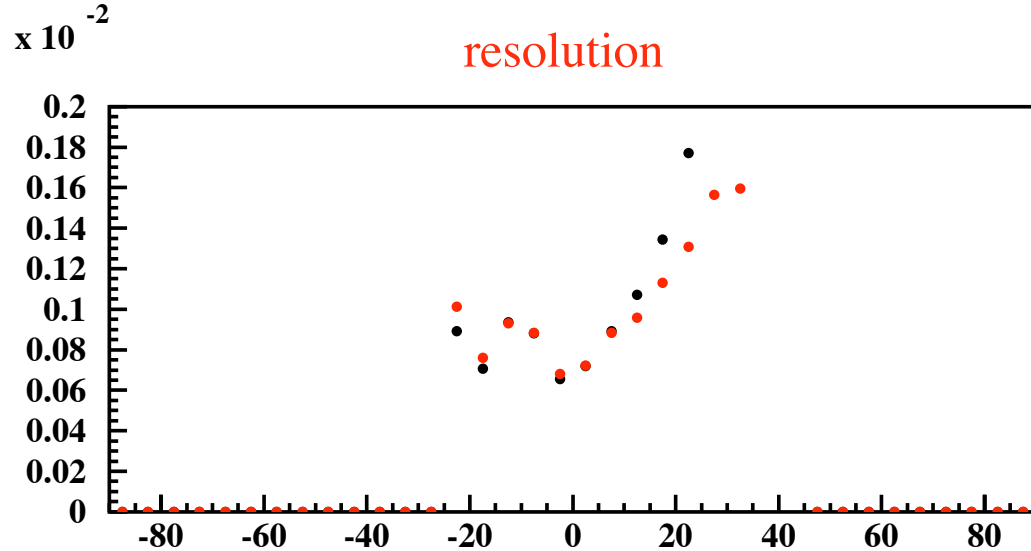


The distributions are fitted with a double gaussian function, the two components of which express the peak and the wider tail respectively.

The two variances are combined as follows to yield a RMS value that represents the combined distribution.

$$f(x) = A_1 \exp\left(\frac{-x^2}{\sigma_1^2}\right) + A_2 \exp\left(\frac{-x^2}{\sigma_2^2}\right)$$
$$\sigma_{\text{comb}}^2 = \sigma_1^2 \left(\frac{A_1 \sigma_1}{A_1 \sigma_1 + A_2 \sigma_2} \right) + \sigma_2^2 \left(\frac{A_2 \sigma_2}{A_1 \sigma_1 + A_2 \sigma_2} \right)$$

Resolution, old vs. new (r/phi)

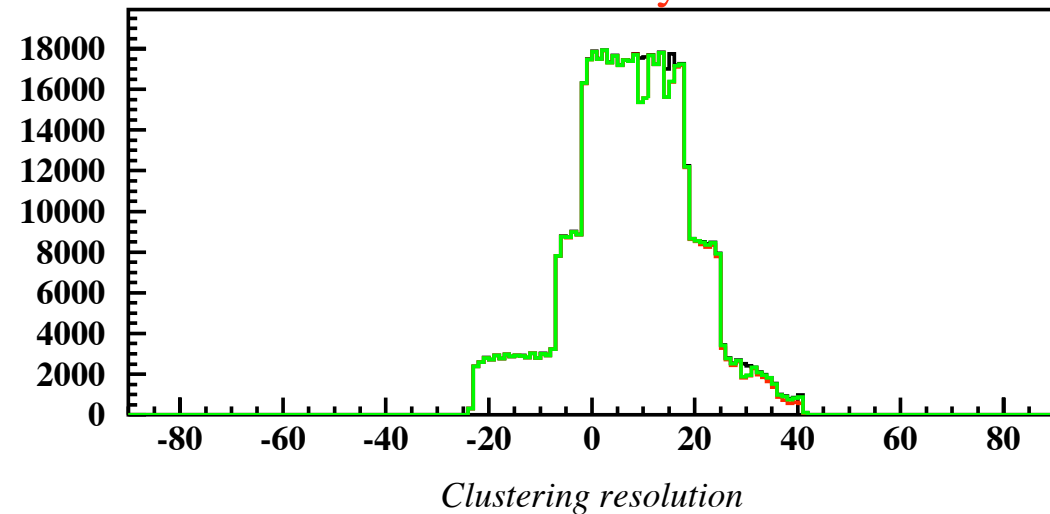


Similar or better resolution.
Better efficiency with new method.

Same true for z side.

Use this new clustering method in tracking to check overall resolution.

efficiency

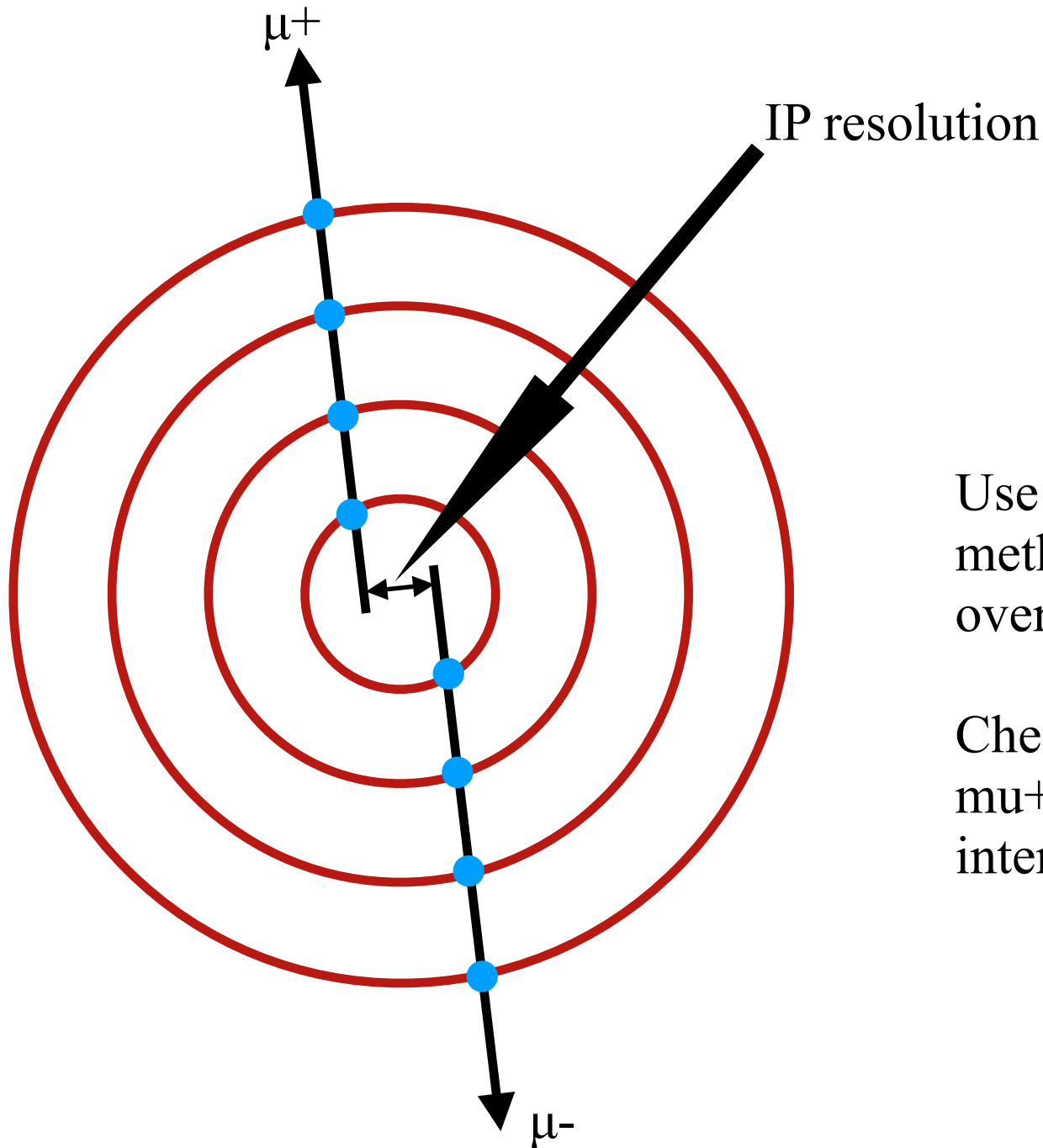


X : local incident angle

Y (top) : RMS

Y(bottom) : number of counts.

Interaction point resolution



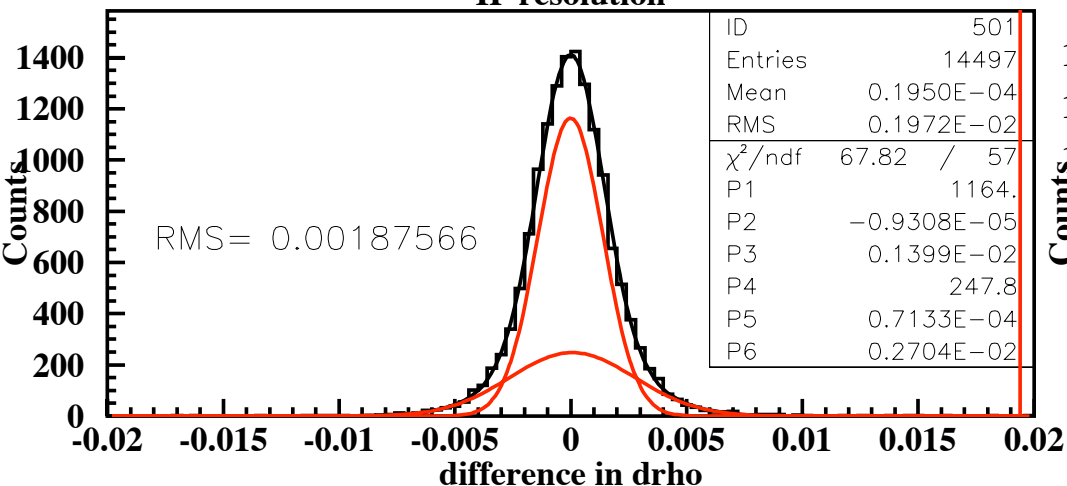
Use this new clustering method in tracking to check overall resolution.

Check distance between μ^+ and μ^- tracks at the interaction point.

IP resolution - MC - resolution

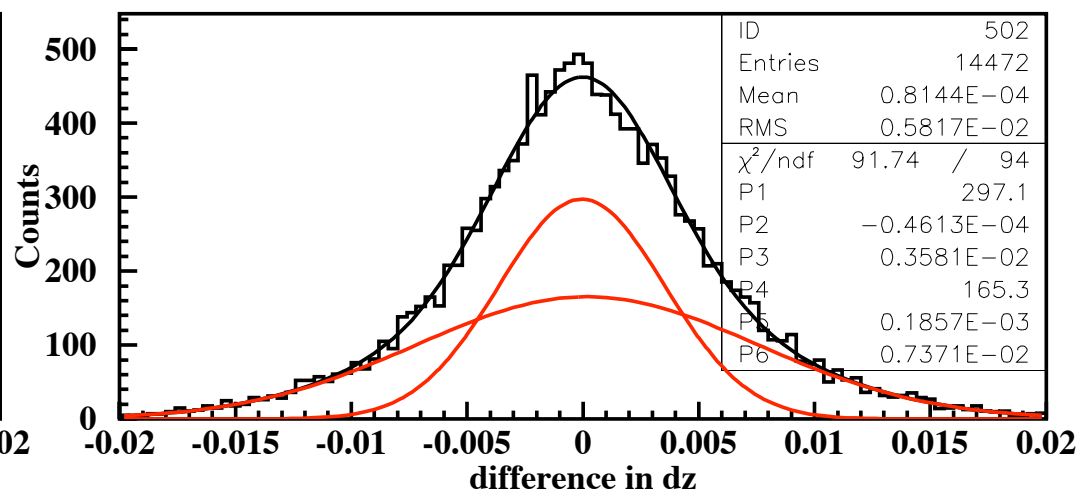
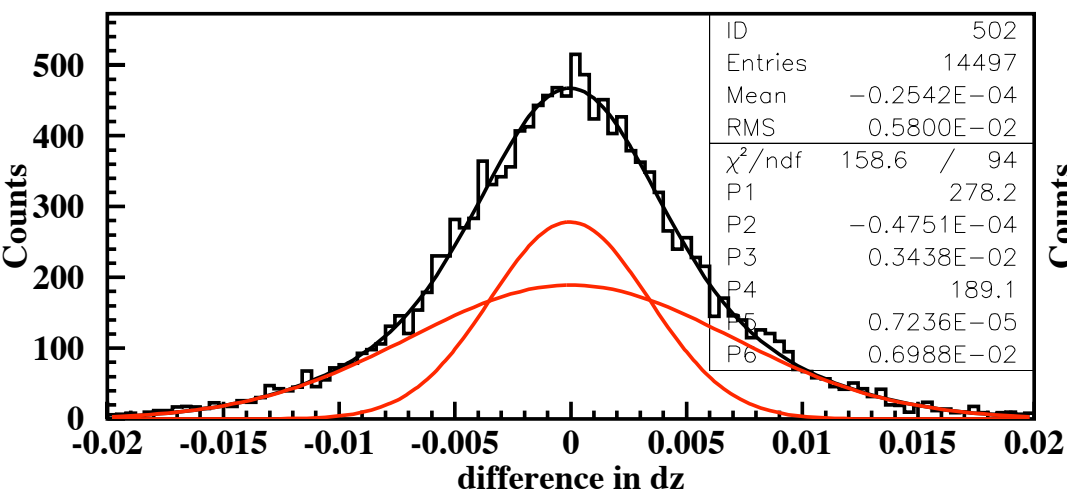
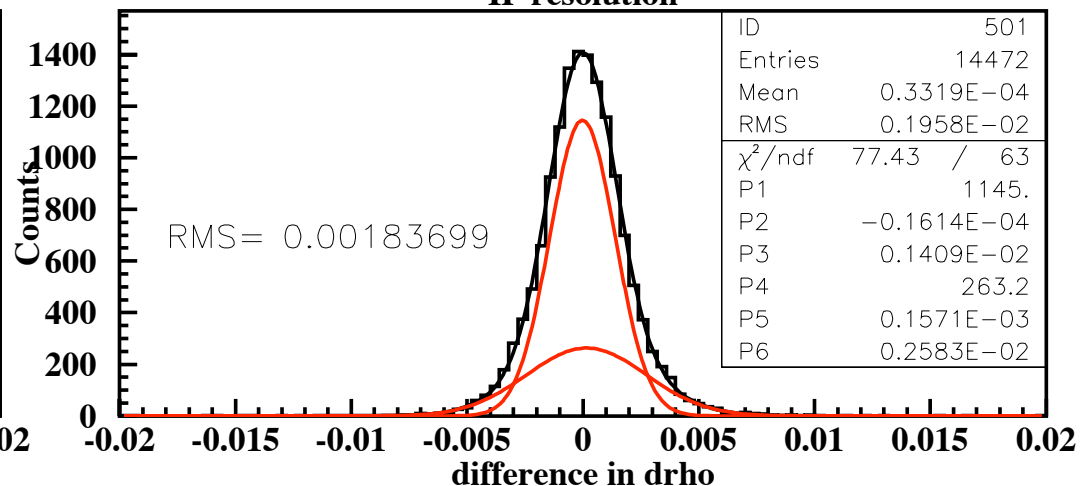
recsvd (old clustering method)

IP resolution



reclus (new clustering method)

IP resolution

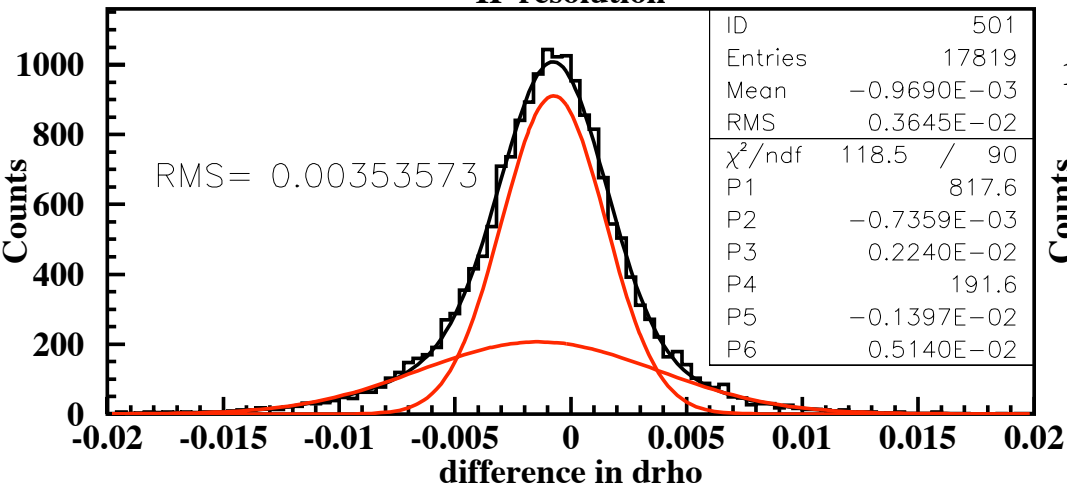


Small reduction in RMS, sigma of peak and tail components of double gaussian fit, and weighted mean of double gaussian fit.

IP resolution - data (Exp 37) - resolution

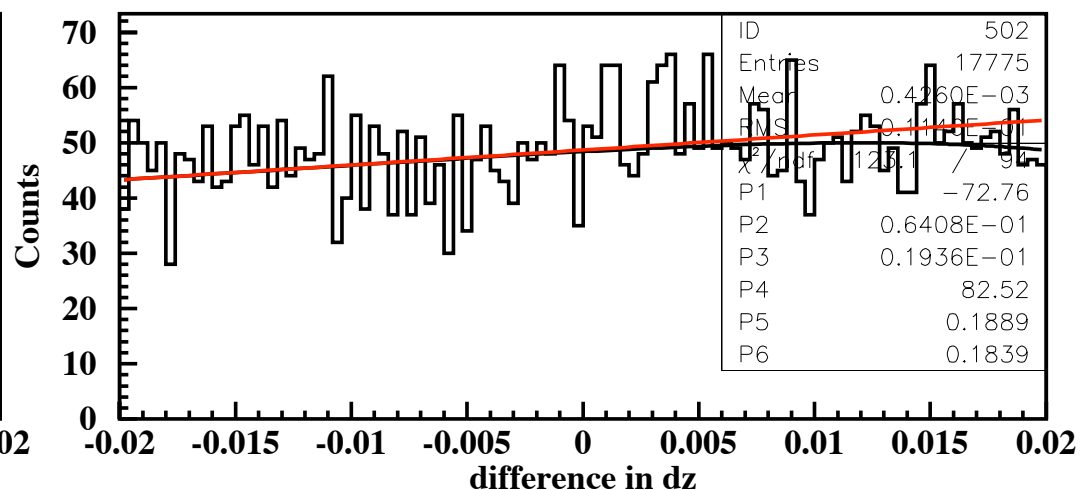
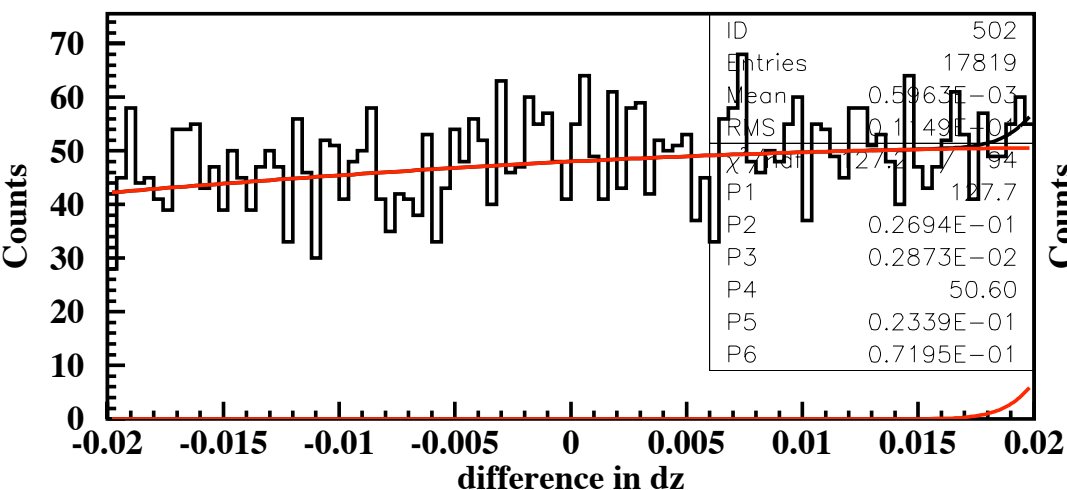
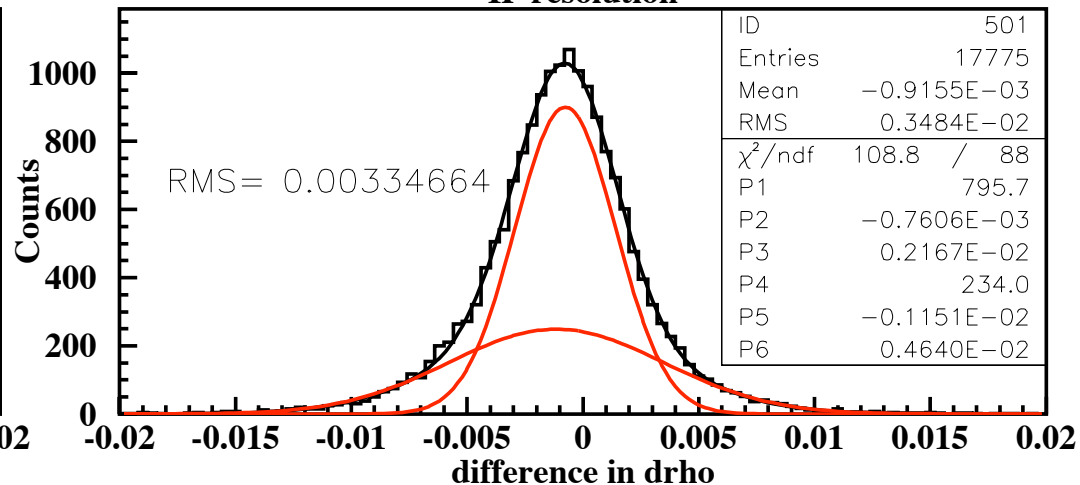
recsvd (old clustering method)

IP resolution



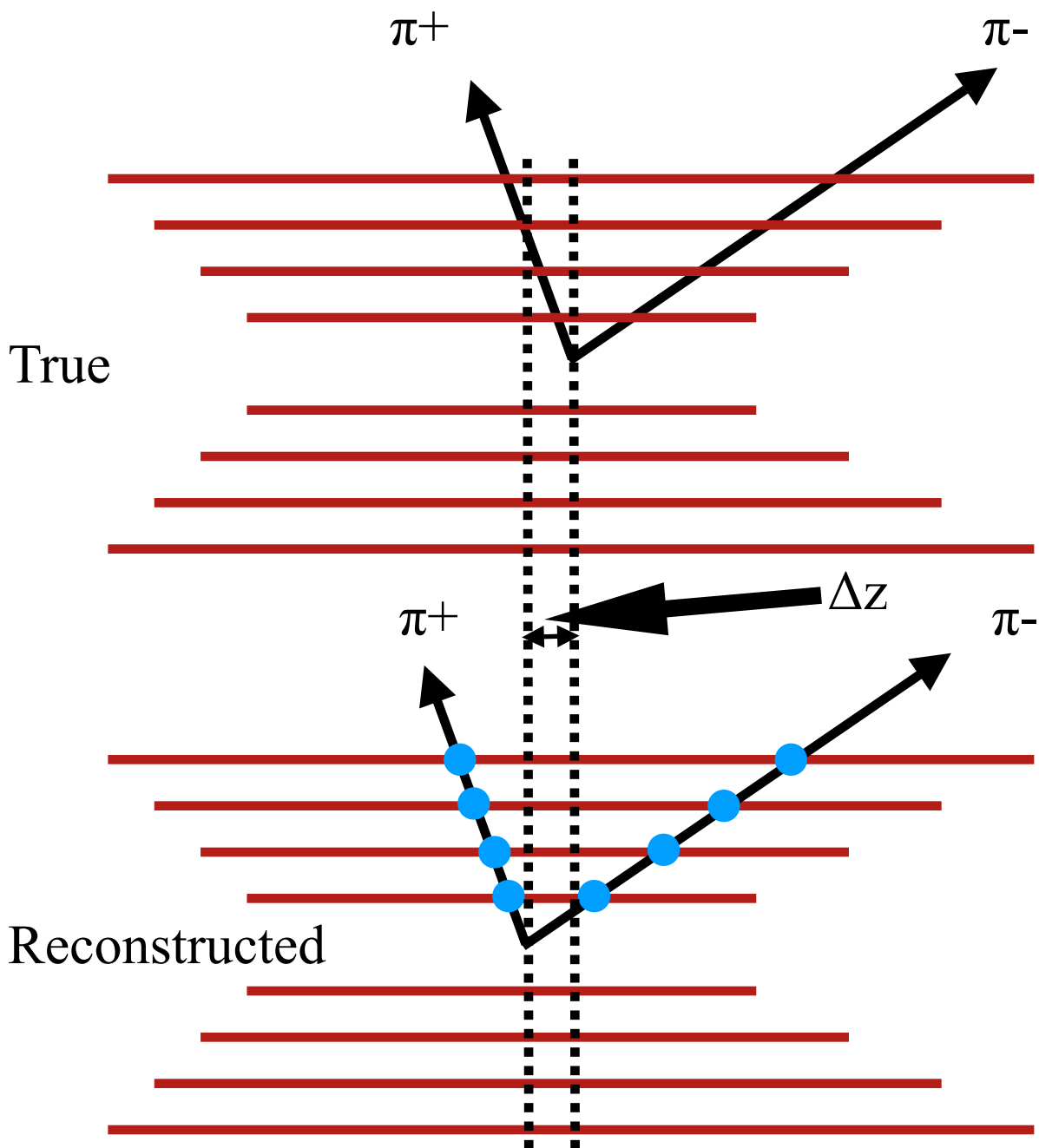
reclus (new clustering method)

IP resolution



Small reduction in RMS, sigma of peak and tail components of double gaussian fit, and weighted mean of double gaussian fit.

resolution for hadronic CP events



Final goal : to test new method on hadronic B decays, and check reconstructed Δz .

Use MC simulation

$Y(4S) \rightarrow BB$

$B \rightarrow \pi^+ \pi^-$

$B \rightarrow \text{generic}$

Summary

1. Charge distribution depends on incident angle and residual. Shallower incident angles have wider spread.
2. Based on these differences, a new clustering algorithm was created to find cluster positions based on the maximum likelihood method.

Clustering resolution is similar for old and new method for MC.

IP resolution is better for dimuon events for both MC and data.

Currently working on checking resolution using hadronic events.