

ARCADIA FNAL meeting

26-09-24

S.Ciarlantini, C. Pantouvakis, M. Rignanese, A. Zingaretti
INFN Sezione di Padova



TB analysis: status

Assumptions done for the tracking:

- just time coincidence, NO spatial check of any kind
- consider events with just one clz per plane

What's new

Tracking

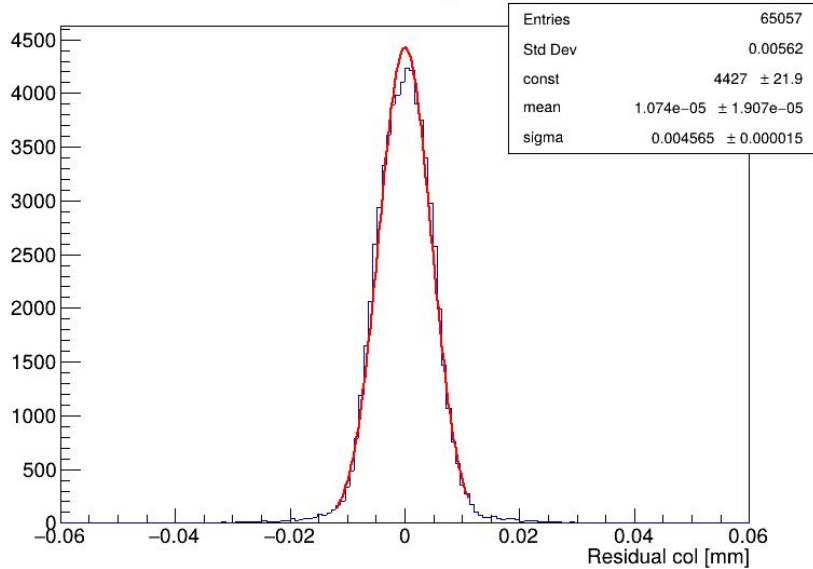
- Fixed kinky clz removal
- Study on residual histograms tails
- Study of resolution vs multiplicity

Residuals after tilt correction

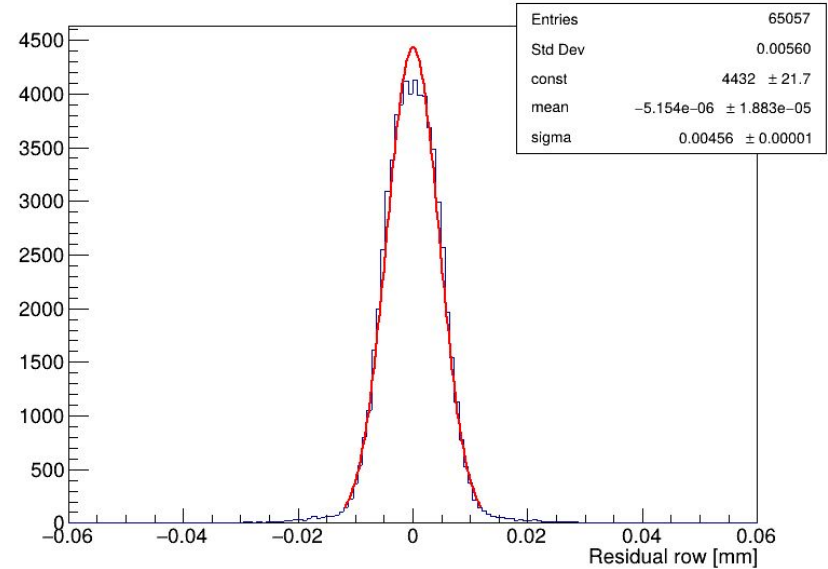
Residuals on single pixel clz on det1 data (multiplicity = 1)

long default run
VCASN = 5

Residual with col det1-det2 aligned and tilt correction



Residual with row det1-det2 aligned and tilt correction



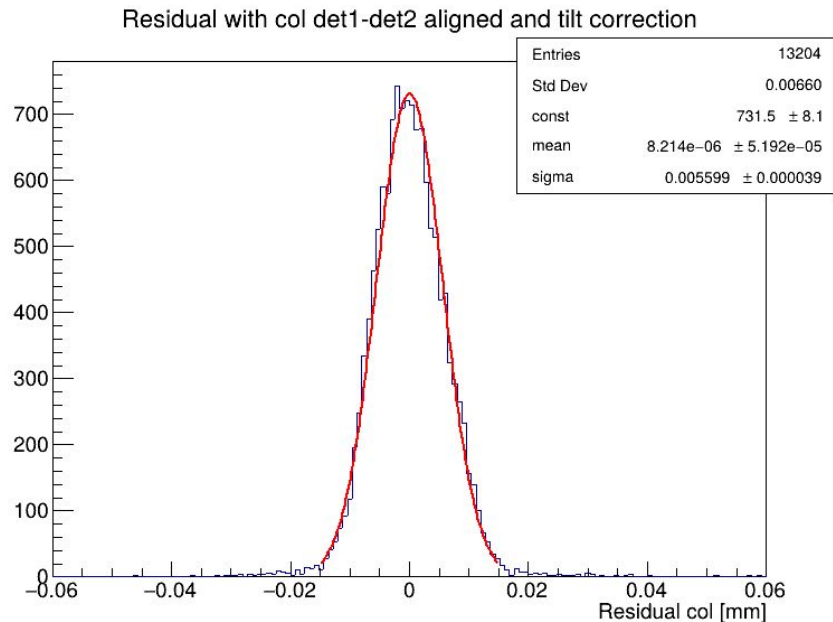
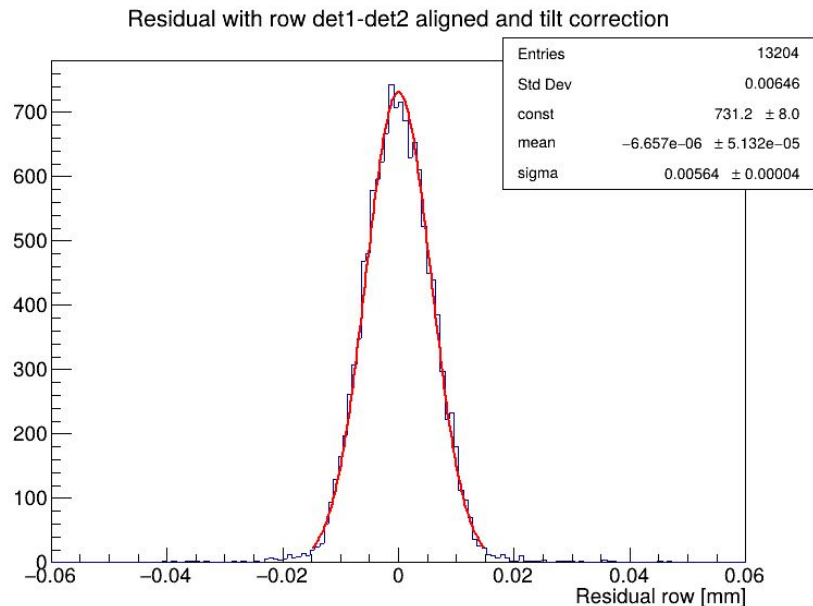
ARCADIA single
point resolution
 $25/\sqrt{12} = 7.2 \text{ um}$

still ~5 um of sigma for single pixel clz
could be that at this high threshold they are not “really”
single pixel clz?

Residuals after tilt correction

Residuals on single pixel clz on det1 data (multiplicity = 1)

Same study but at **lower** threshold \rightarrow VCASN = 20

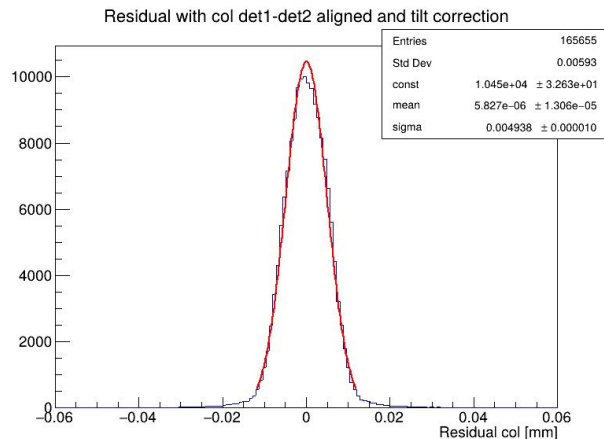


sigmas considering all clz on det1
row = 0.005329
col = 0.005321

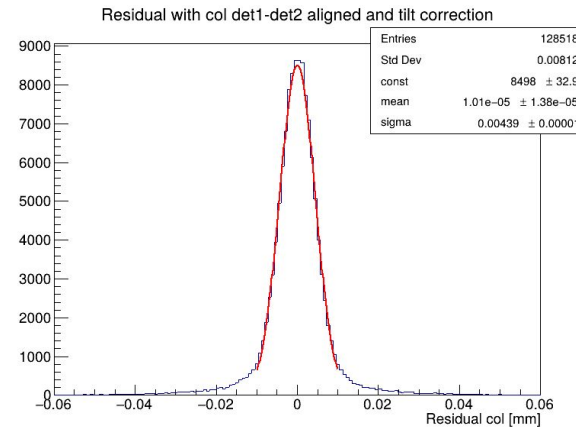


Still not equal to single point resolution
but higher sigmas

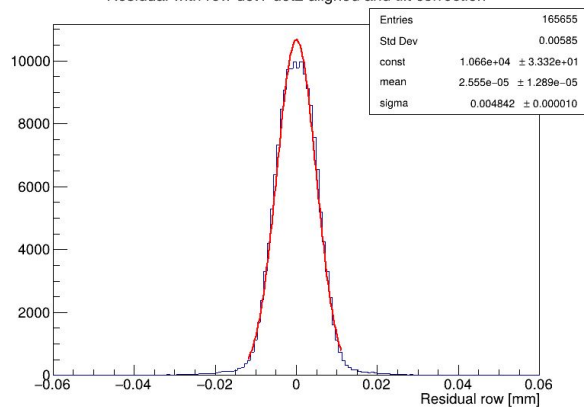
multiplicity = 2



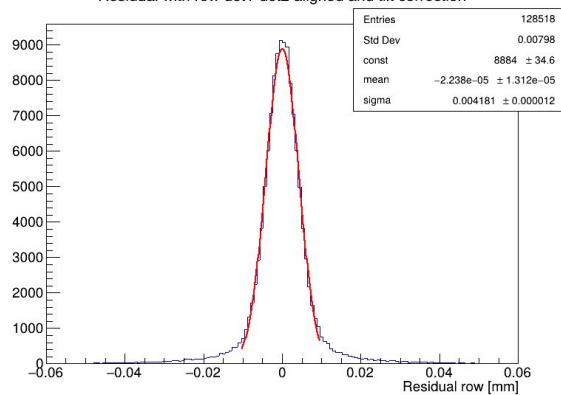
multiplicity > 2



Residual with row det1-det2 aligned and tilt correction

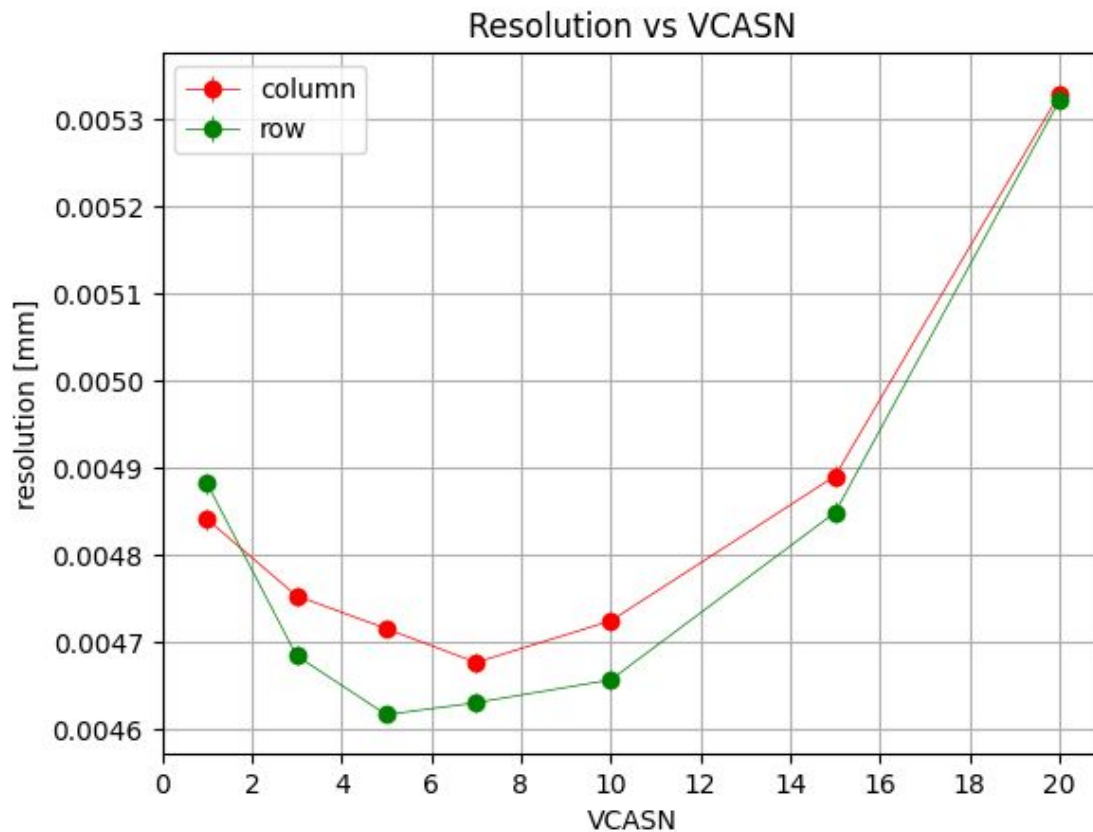


Residual with row det1-det2 aligned and tilt correction



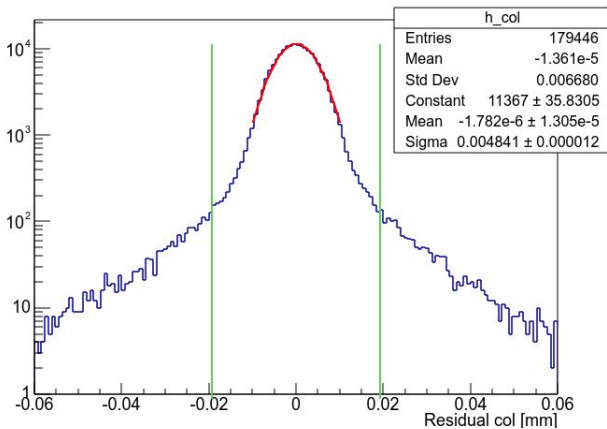
improvement of
resolution as
multiplicity (charge
sharing) increases

Resolution vs threshold (VCASN)

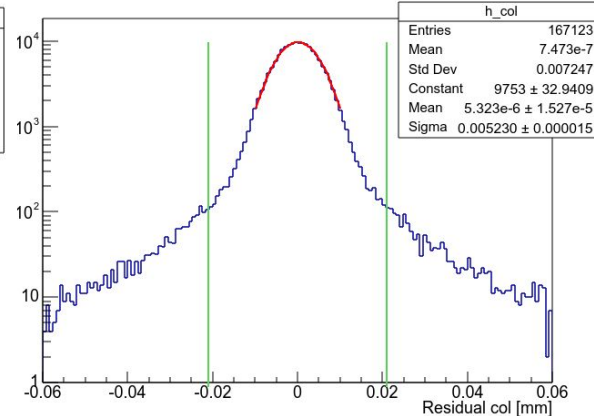


Study on Tails of residual histograms

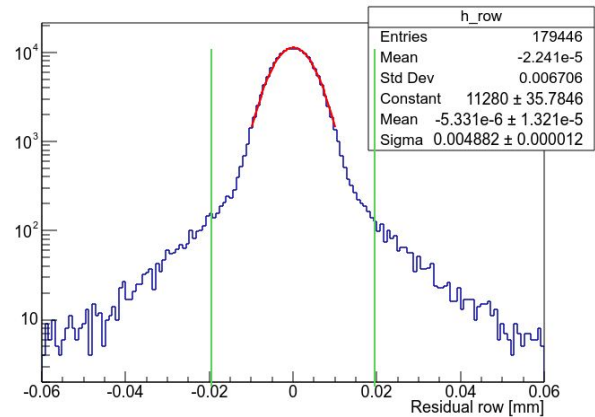
Residual on col VCASN = 1



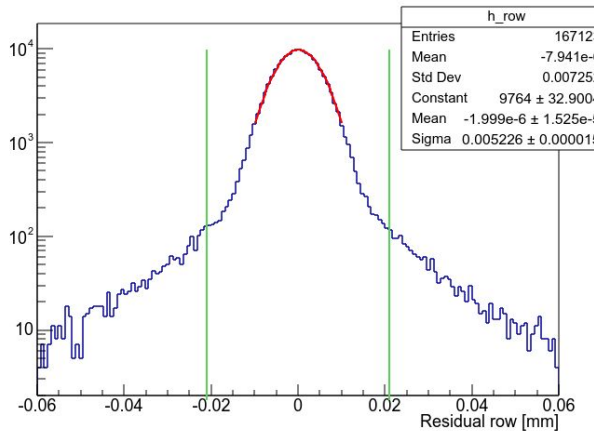
Residual on col VCASN = 20



Residual on rows VCASN = 1



Residual on rows VCASN = 20



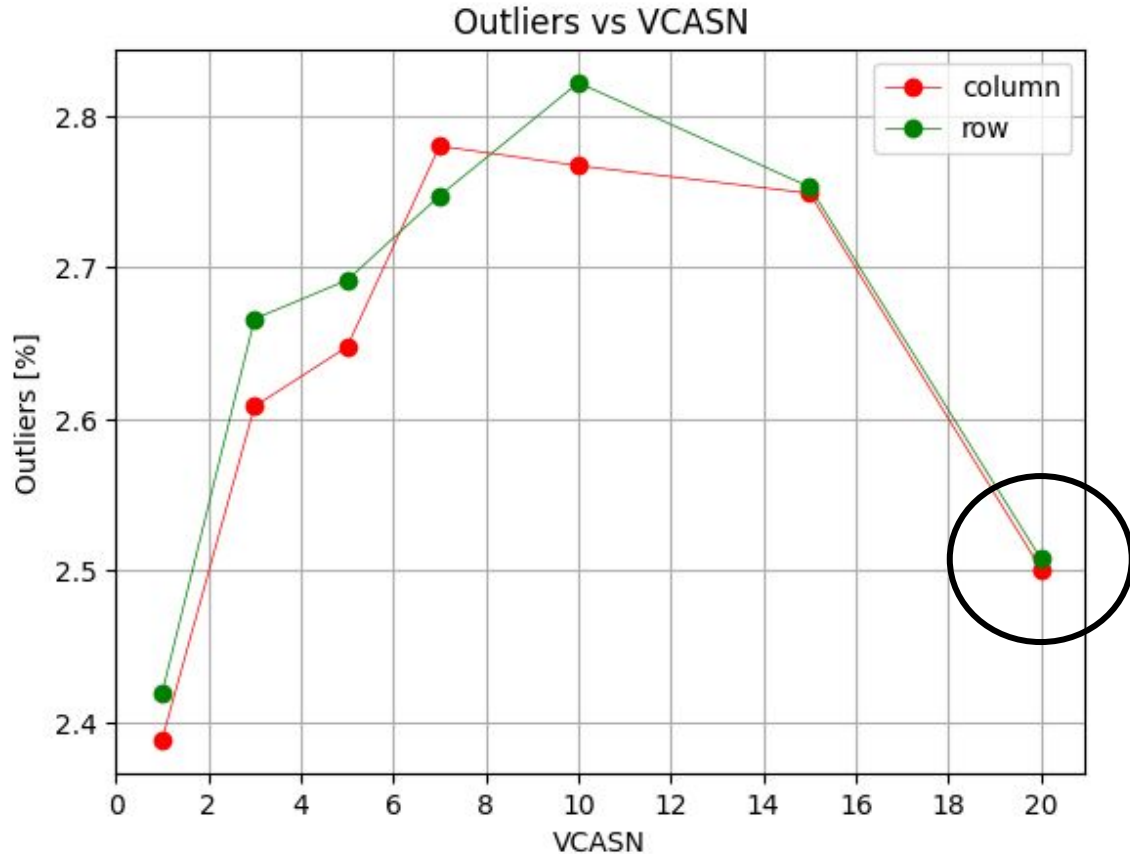
Residual histograms in log scale to enhance tails



Outliers of the histogram calculated as:
residuals $>$ mean + 4*sigma

Mean and sigma come from gaussian fit of residuals with det1-det2 aligned and tilt correction

Study on Tails of residual histograms



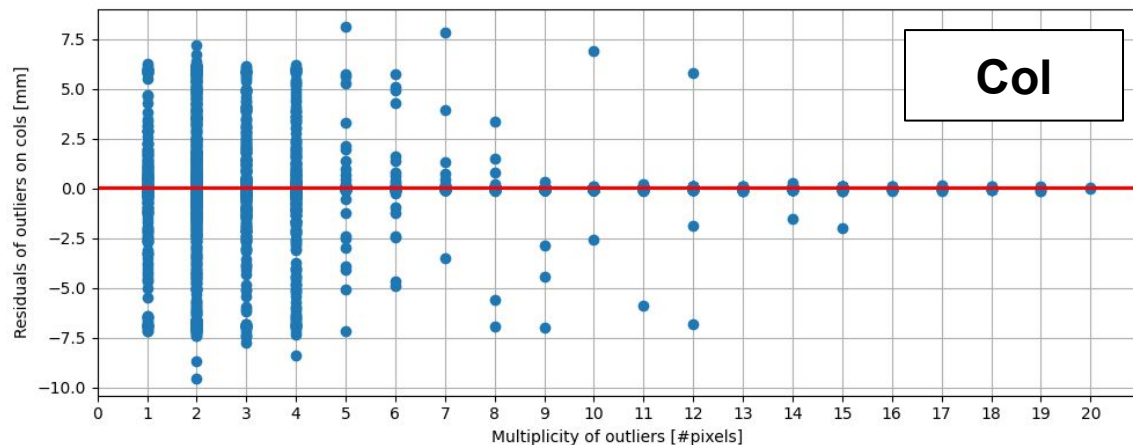
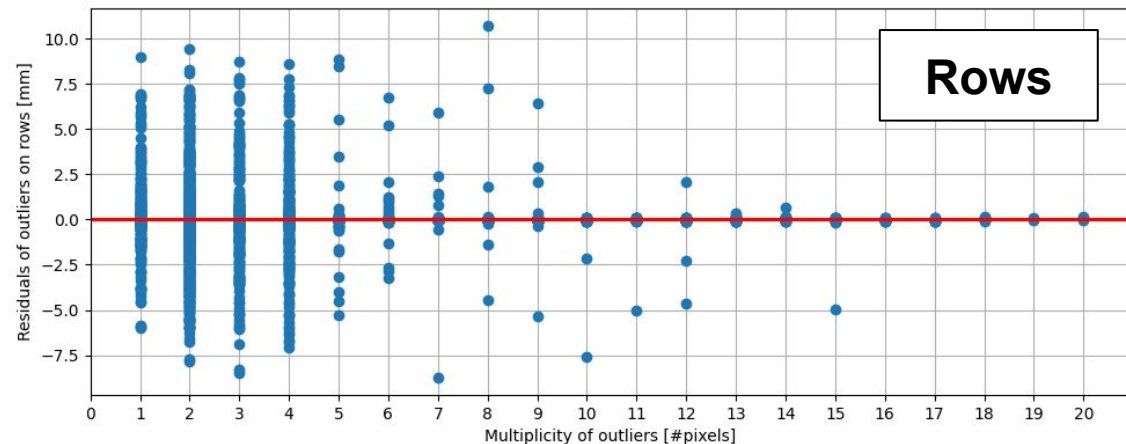
Amount of outliers (in %) per each VCASN



Outliers % grows with VCASN

VCASN = 20 distribution is broader (sigma ~ 0.0053 respect to ~0.0047 @ VCASN = 5) so outliers % is smaller

Study on Tails of residual histograms

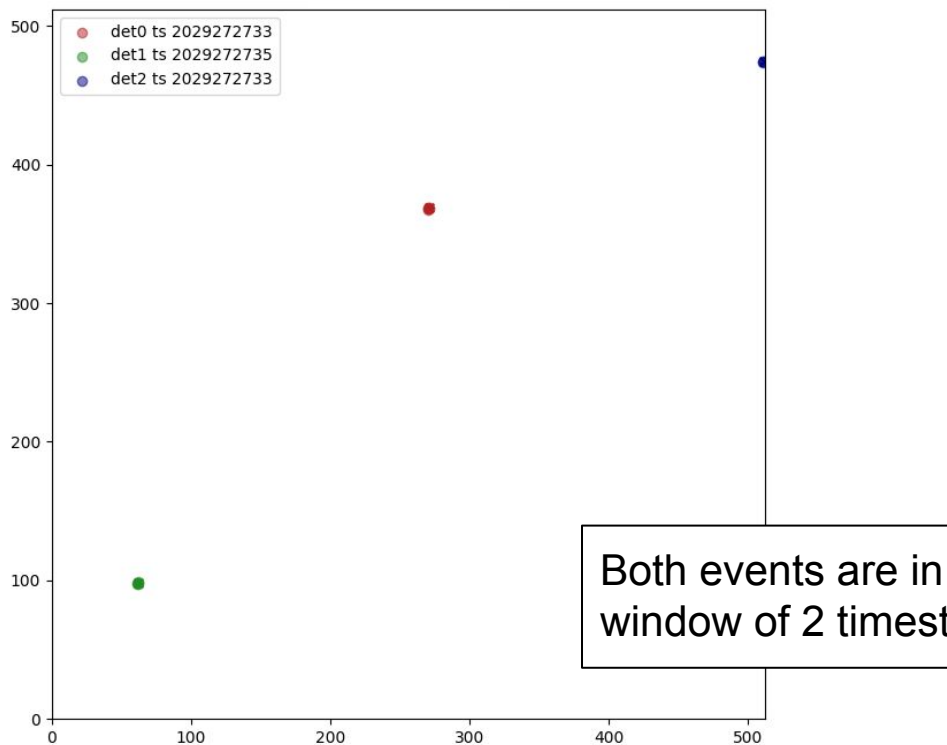


Residuals of clz with high multiplicity (>10) are not so distant from the threshold. They do not contribute heavily to tails

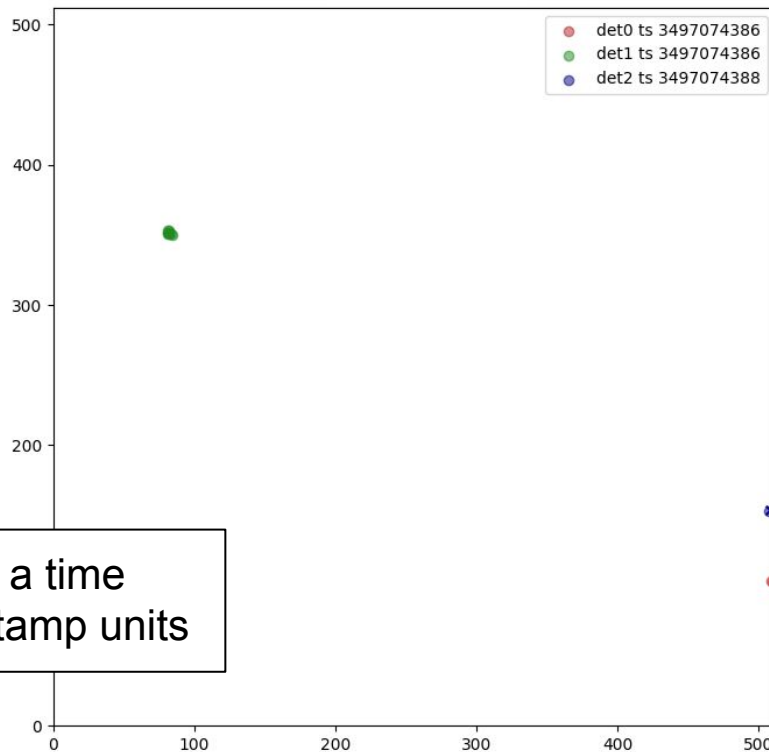
Study on Tails of residual histograms

Single event display

Hitmaps of event with
residual = 8.45 mm on col



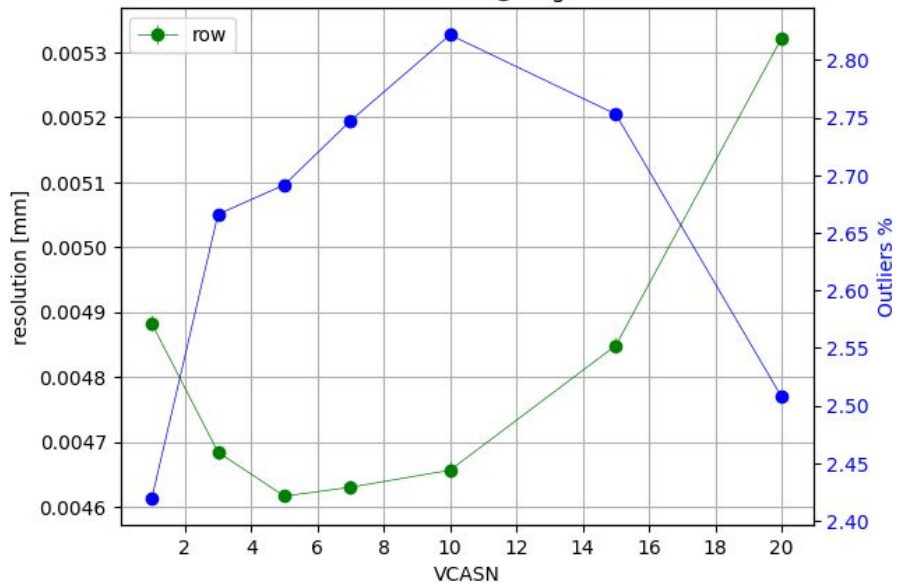
Hitmaps of event with
residual = 10.71 mm on rows



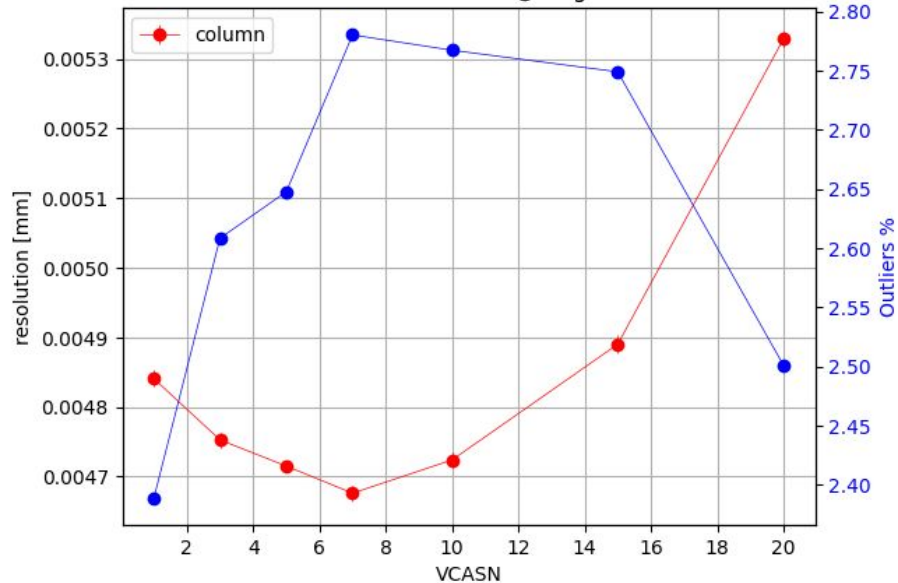
Both events are in a time window of 2 timestamp units

Resolution vs threshold (VCASN) and % of outliers

Row resolution @ angle 0°

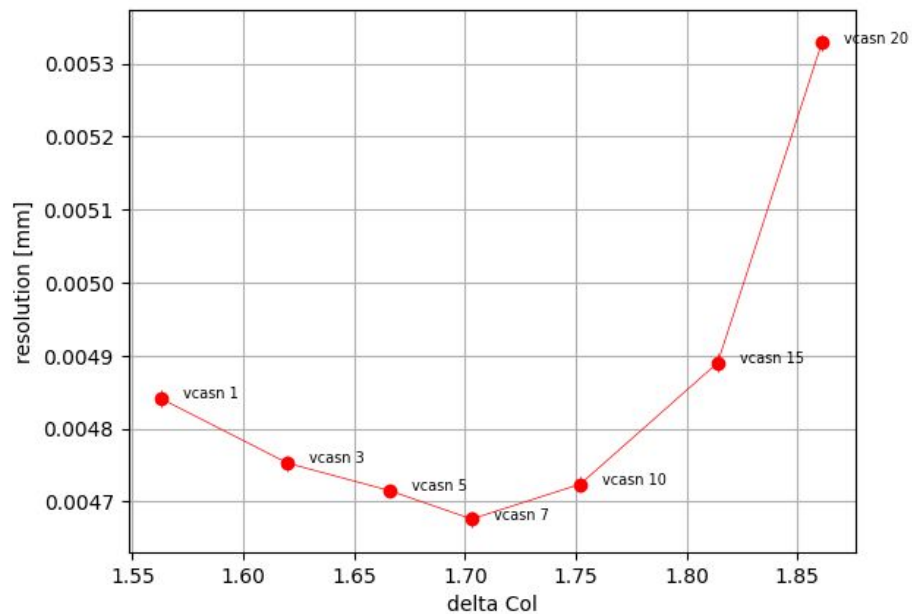


Column resolution @ angle 0°

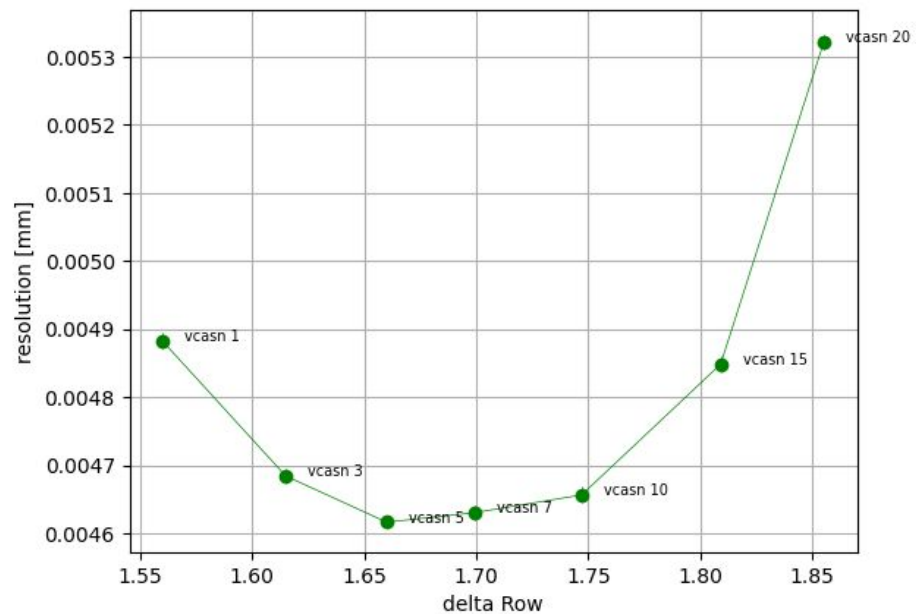


Resolution vs clz extension (delta row, delta col)

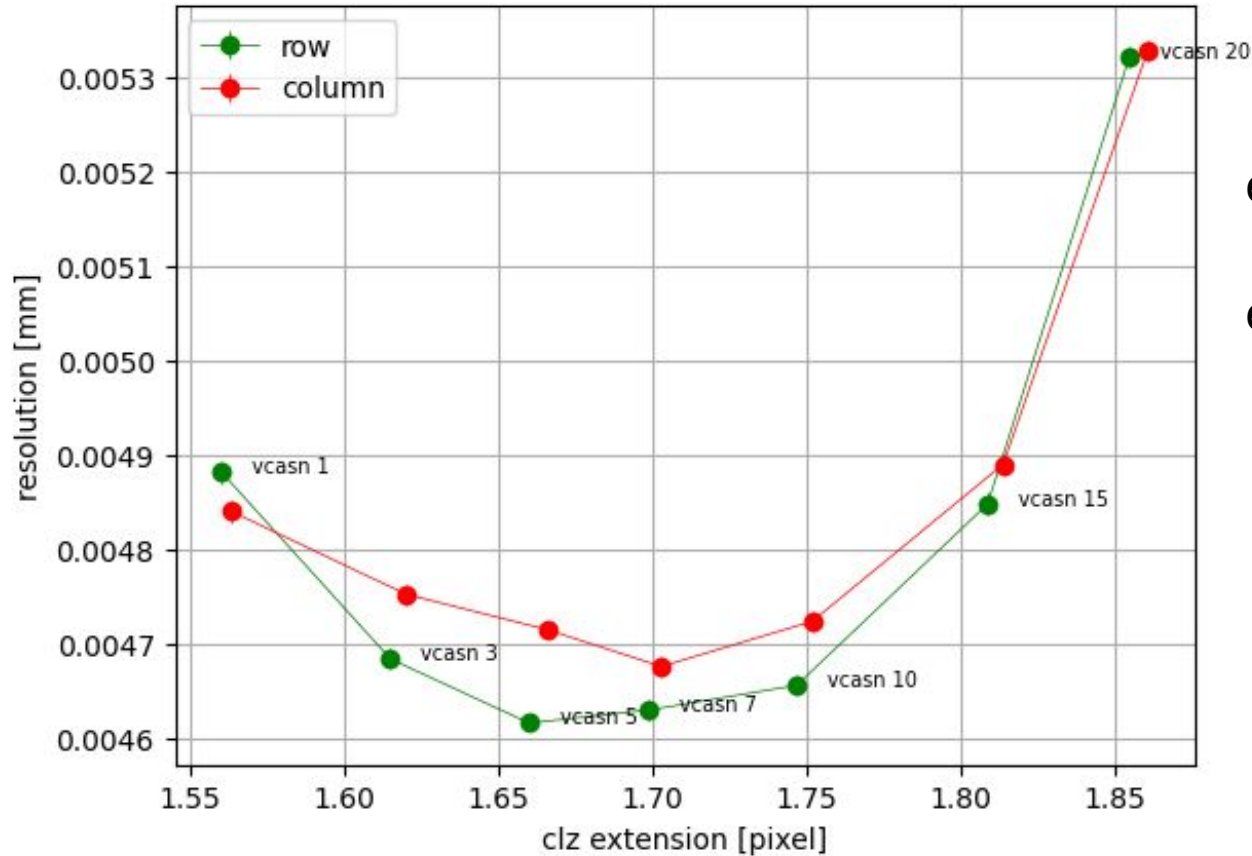
Resolution on col vs delta col



Resolution on row vs delta row



Resolution vs clz extension (delta row, delta col)

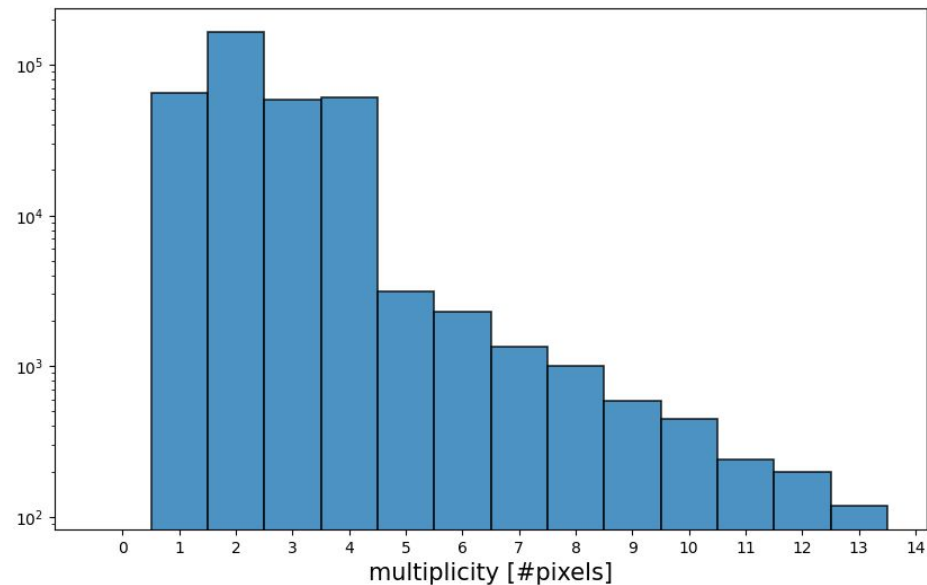
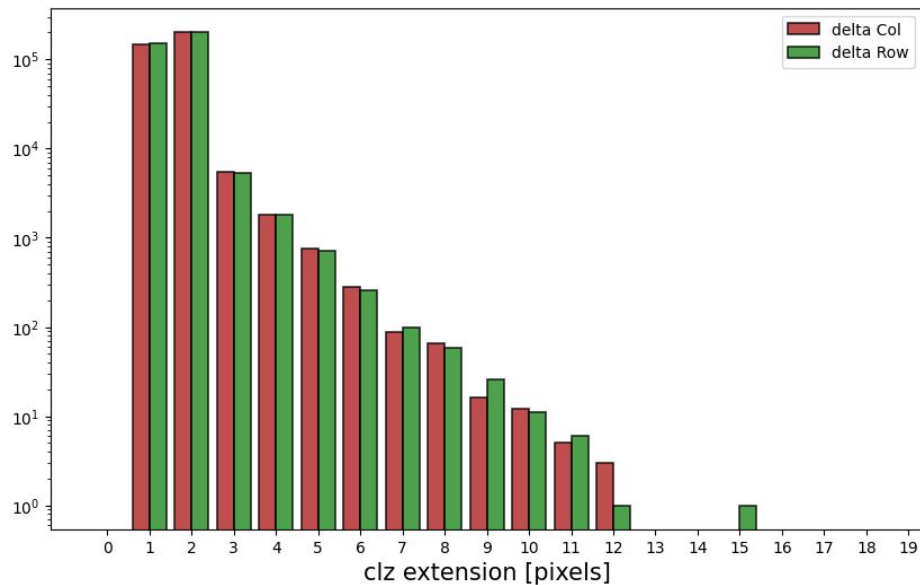


extension on row: **delta Row**

extension on col: **delta Col**

Distribution of det1 clusters used in tracking

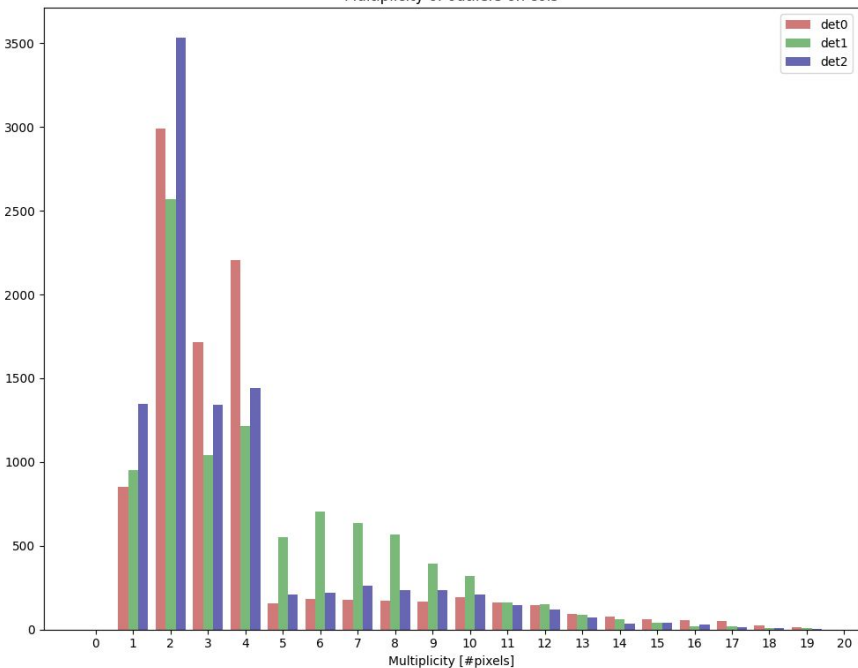
Number of det1 clz used in tracking: 359230



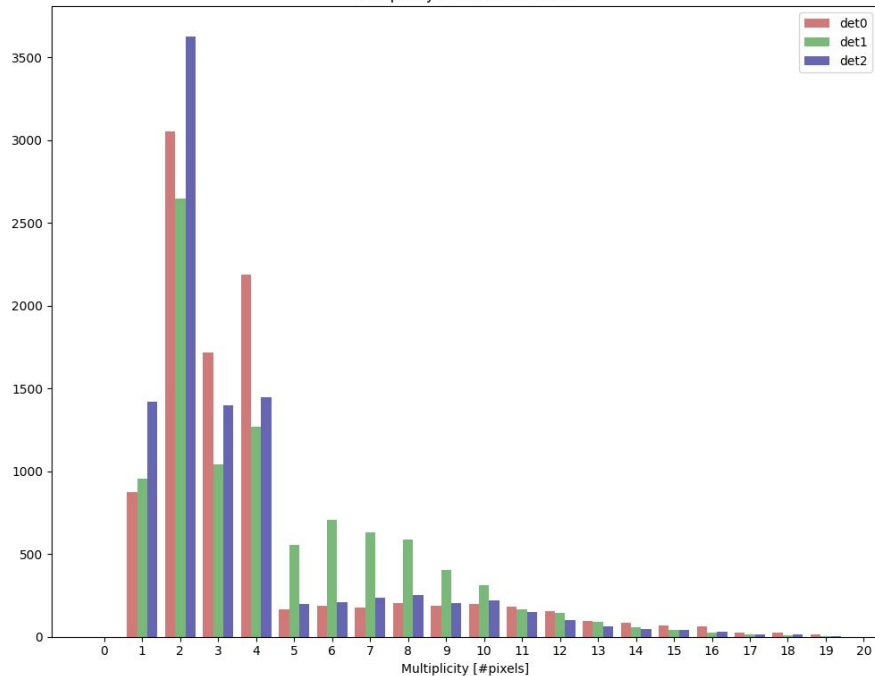
BACK UP

Study on Tails of residual histograms

Multiplicity of outliers on cols

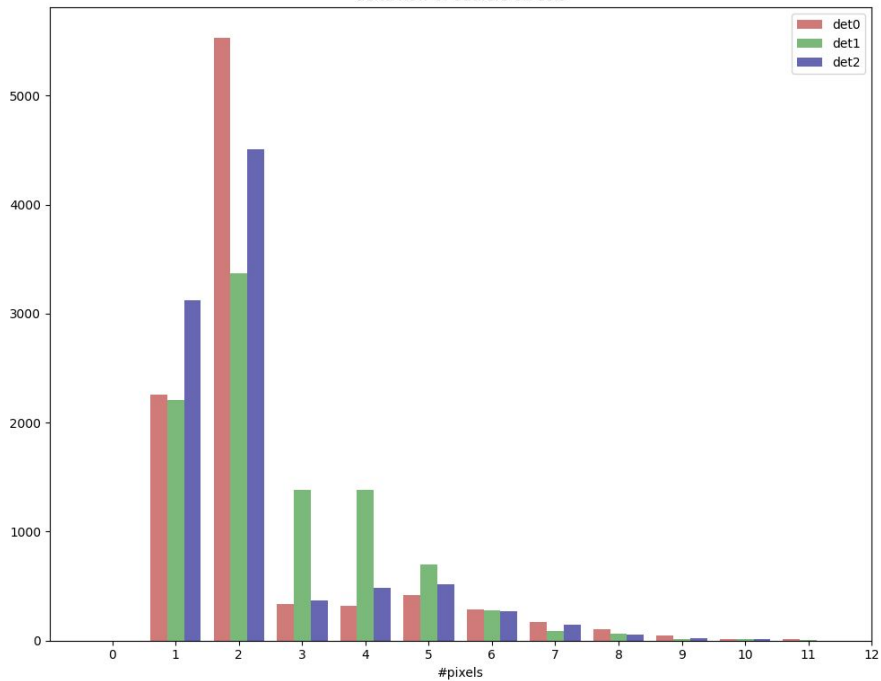


Multiplicity of outliers on rows



Study on Tails of residual histograms

delta Row of outliers on cols



delta Row of outliers on rows

