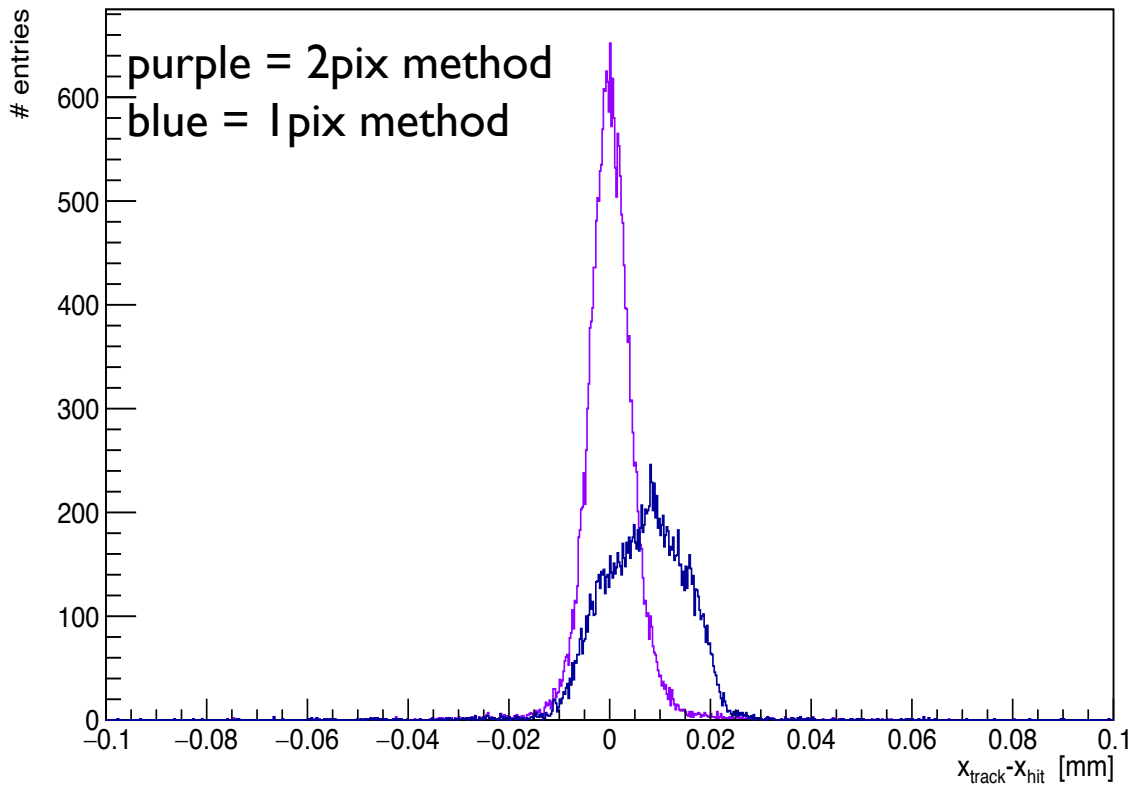


# CUT SCANS FOR DESY TB RECO

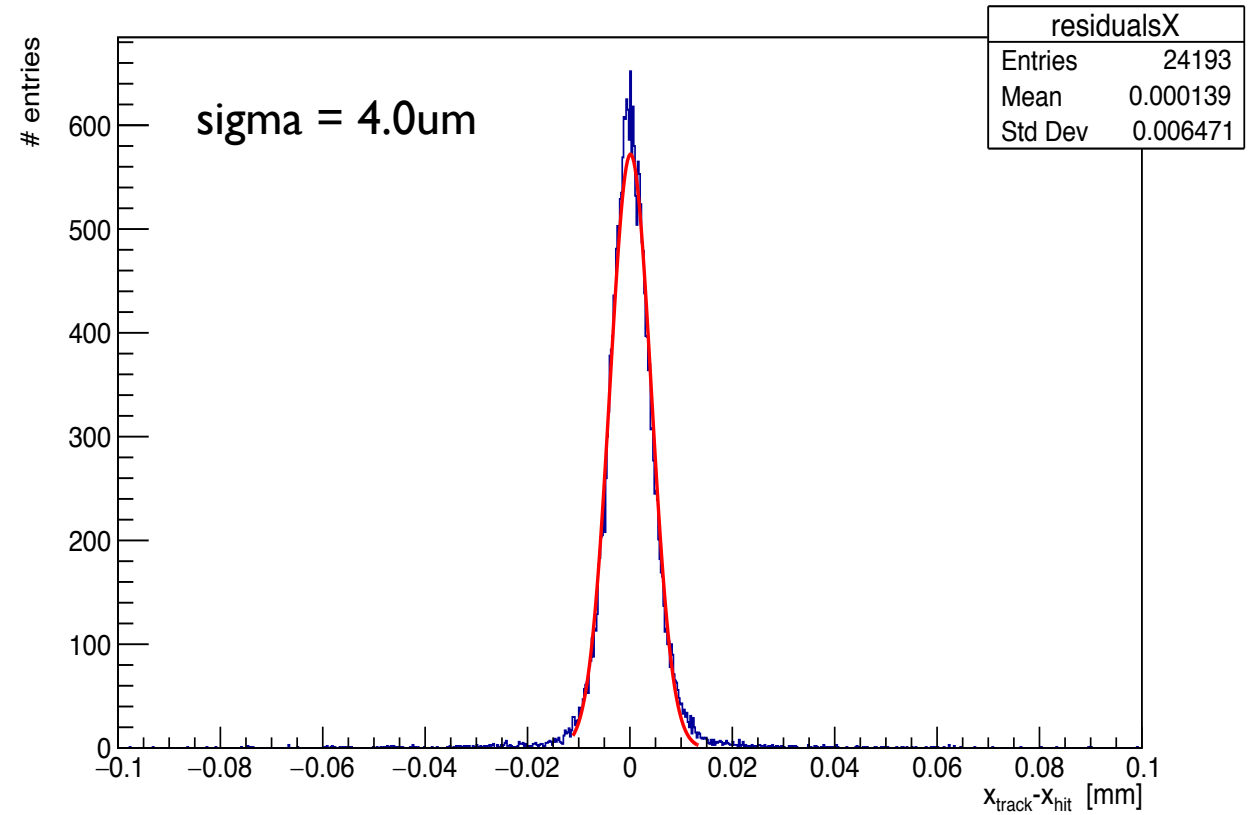
1/7/2020

# IMPLEMENTED 2-PIXEL UNIT CELL

residualsX

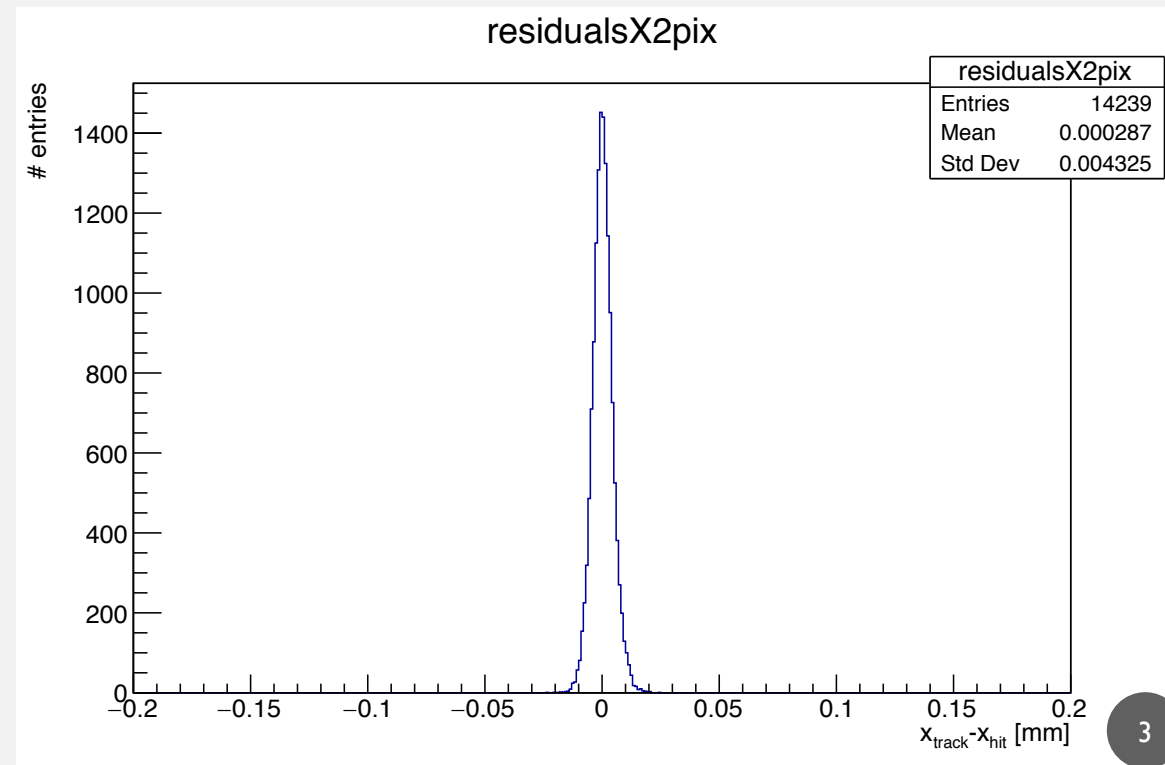
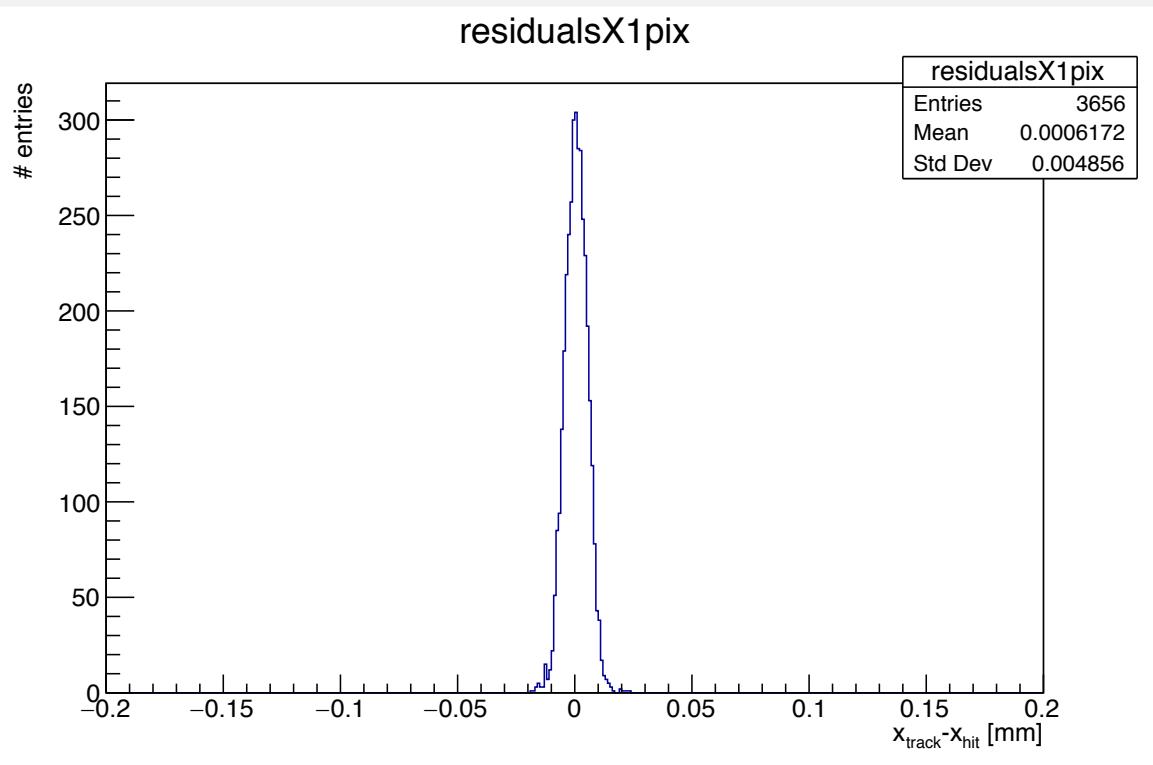
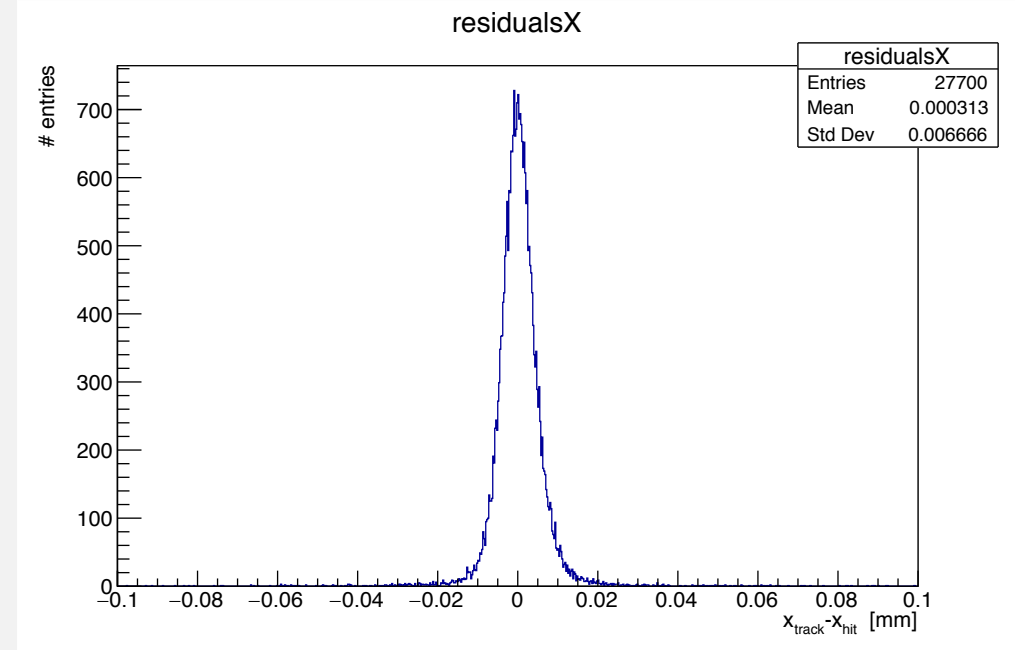


residualsX

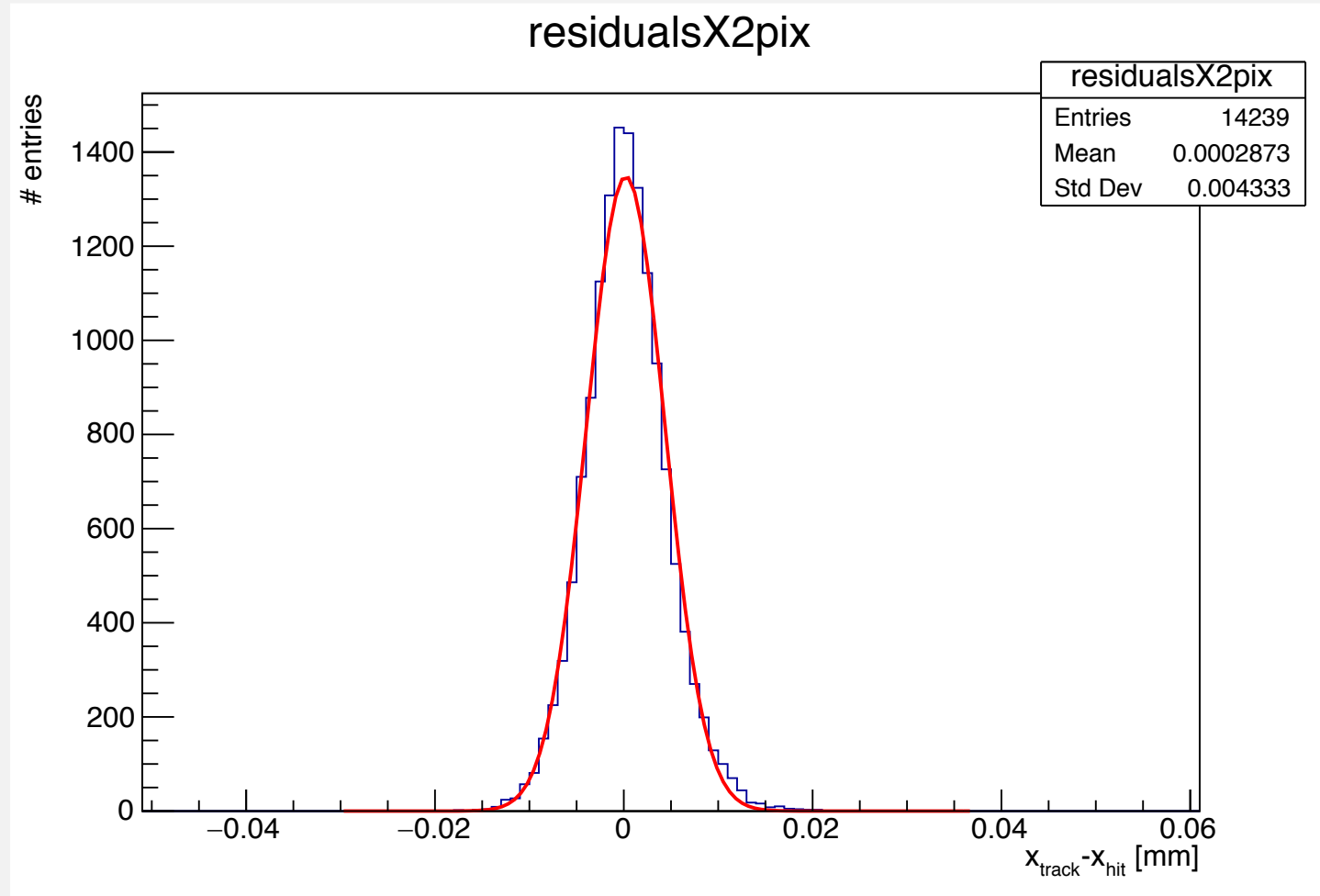


# RESIDUAL FOR DIFFERENT CLUSTER SIZES

	All clusters	1 pix clusters	2 pix clusters
Gauss mean	0.15 $\mu$ m	0.66 $\mu$ m	0.19 $\mu$ m
Gauss sigma	3.94 $\mu$ m	4.68 $\mu$ m	4.12 $\mu$ m



# SLIGHT ASYMMETRY OF 2 PIX CLUSTERS



# SPATIAL TRACKING CUT

Want to determine the spatial cut that should be used during track formation.

Timepix3 plane has the widest residual, with a gauss sigma of  $\sim 20\mu\text{m}$ .

Mimosa planes have residuals  $\sim 2.5\mu\text{m}$ .

Alignment error in X and Y is  $\sim 1\mu\text{m}$  and  $1^\circ$  per plane.

Z alignment error is  $\sim 0.2\text{mm}$ .

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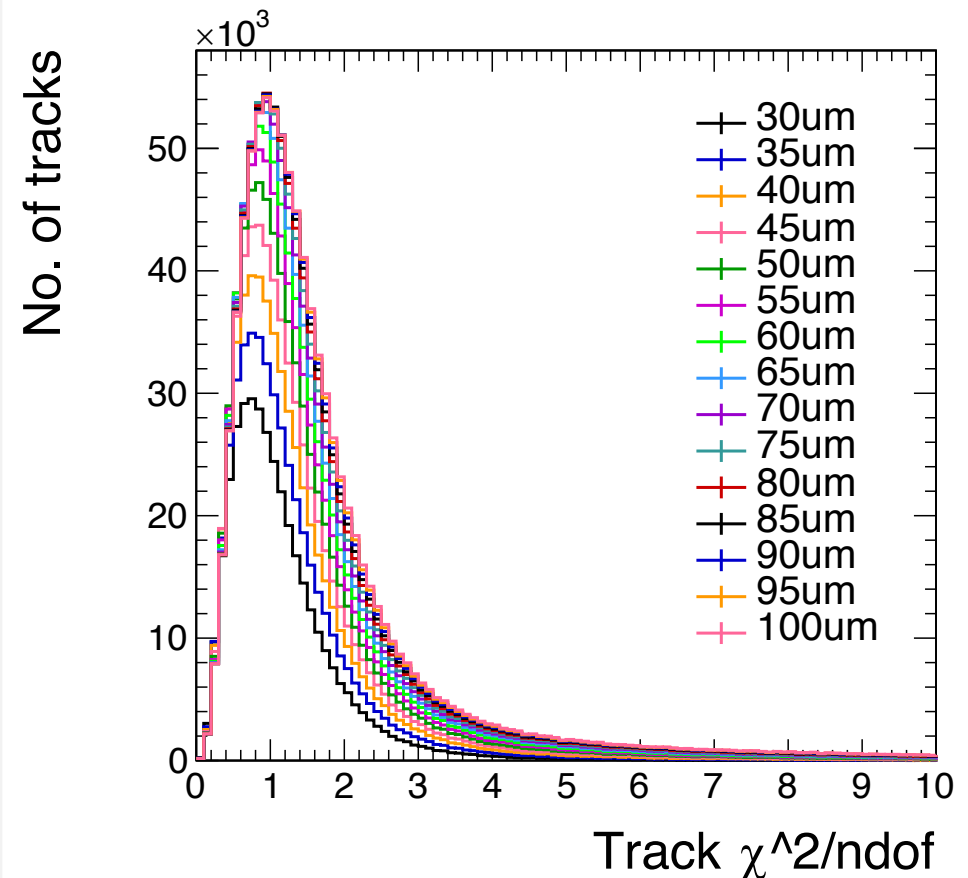
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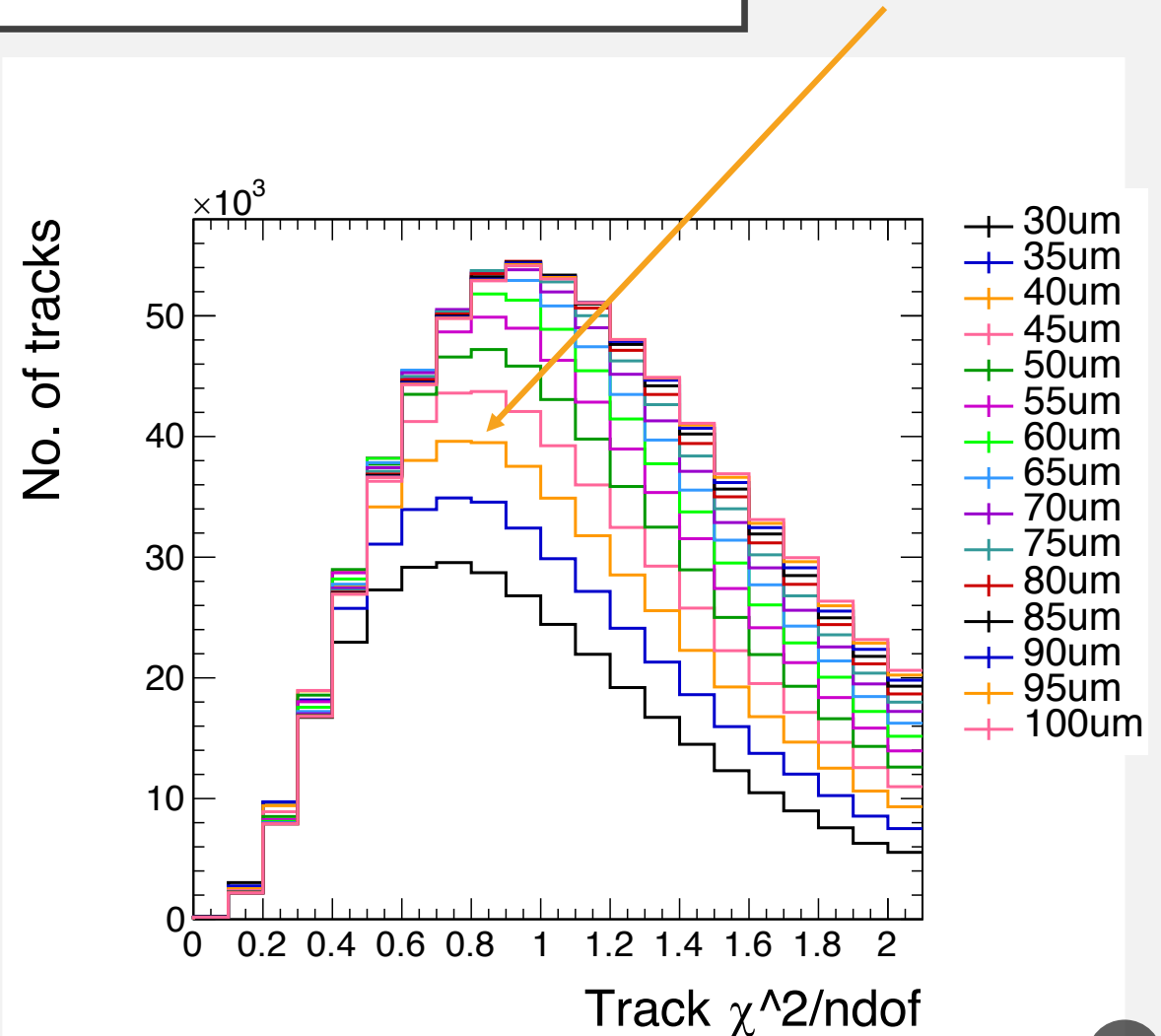
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To check the effect of different cuts, did a spatial tracking cut scan and plotted the  $\chi^2/\text{ndof}$  distribution.

A tracking cut of  $40\mu\text{m}$  has a peak  $\chi^2/\text{ndof}$  at  $\sim 0.9$ .



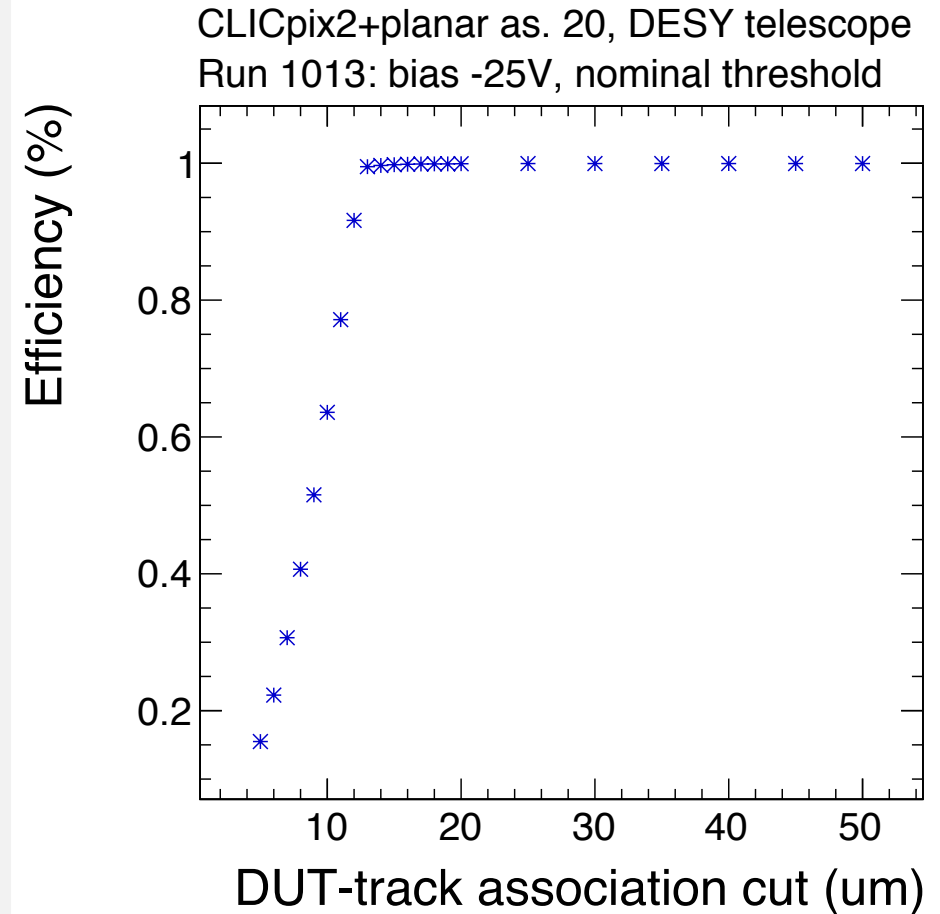
# TRACK-DUT ASSOCIATION CUT

- Now want to determine the best cut value for DUT-track association in space.
- Note: using the tracking cut just determined of 40um.
- Using run 1013: -25V bias, nominal threshold ~800 electrons.



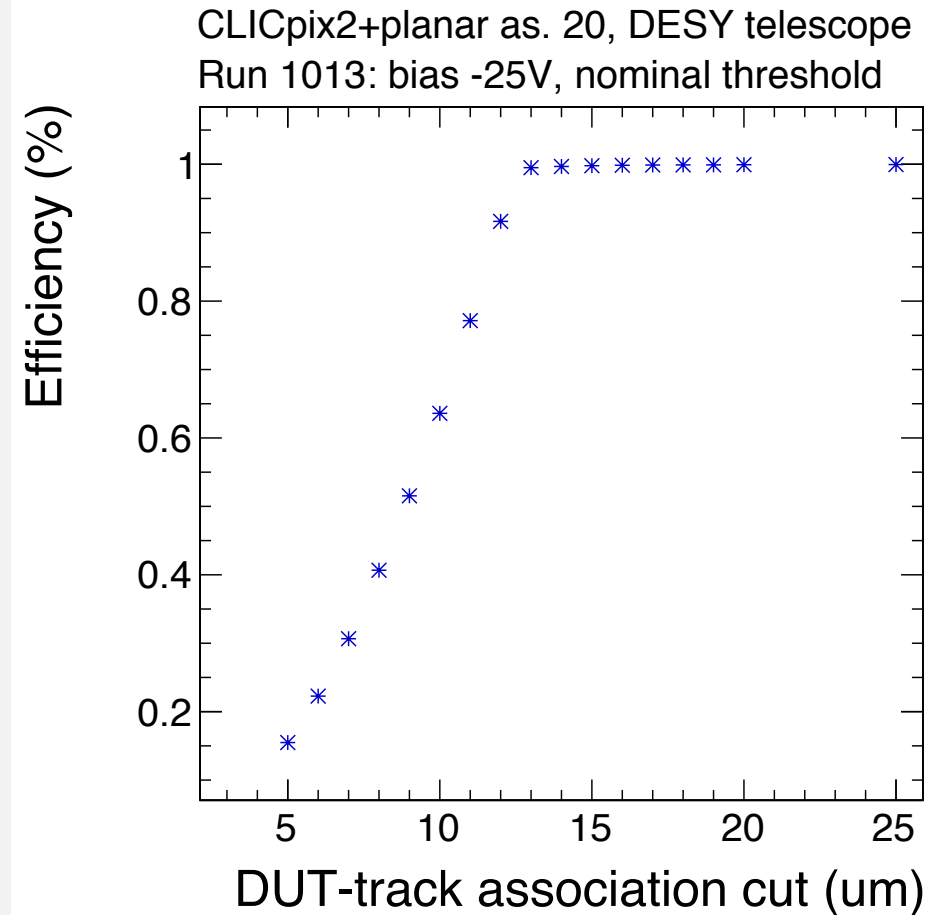
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- Using run 1013: -25V bias, nominal threshold ~800 electrons.
- Scanned association cut value from 5 $\mu$ m to 20 $\mu$ m in steps of 1 $\mu$ m, and 25 $\mu$ m to 50 $\mu$ m in steps of 5 $\mu$ m.
- Right: see a plot of the efficiency at each value.



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- Right: see a plot of the efficiency at each value.
- See high efficiency ( $\sim 99.9\%$ ) for association cuts down to 13 $\mu\text{m}$ , then efficiency decreases rapidly.  
→ choose a DUT-track association cut of 13 $\mu\text{m}$



# QUESTIONS

Sanity-check questions before batch processing runs:

- Is 40um a reasonable track cut?
- Is 13um a reasonable DUT association cut?
- Which runs can I apply these cuts to? (change in bias voltage, threshold, DUT alignment)
- Can I apply the same eta correction to each run? (change in bias voltage, threshold, DUT alignment)