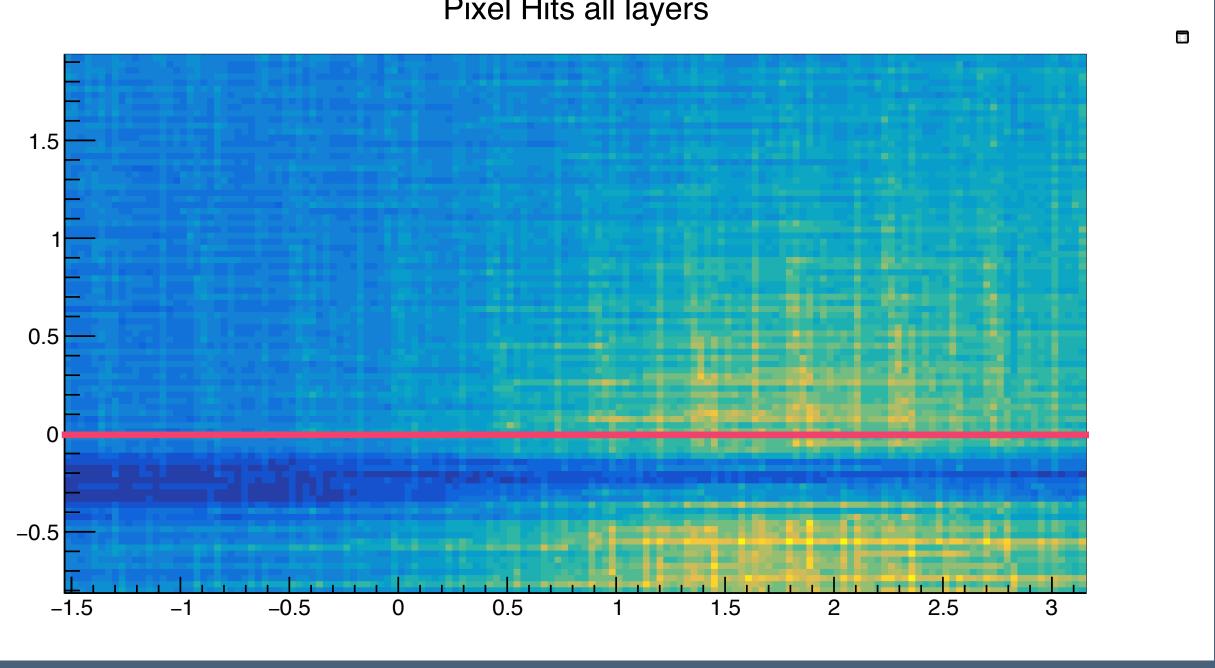
# Chip Alignment

Nick Sundstrom EPICAL2 Meeting Nov 14

#### Reminder from last week

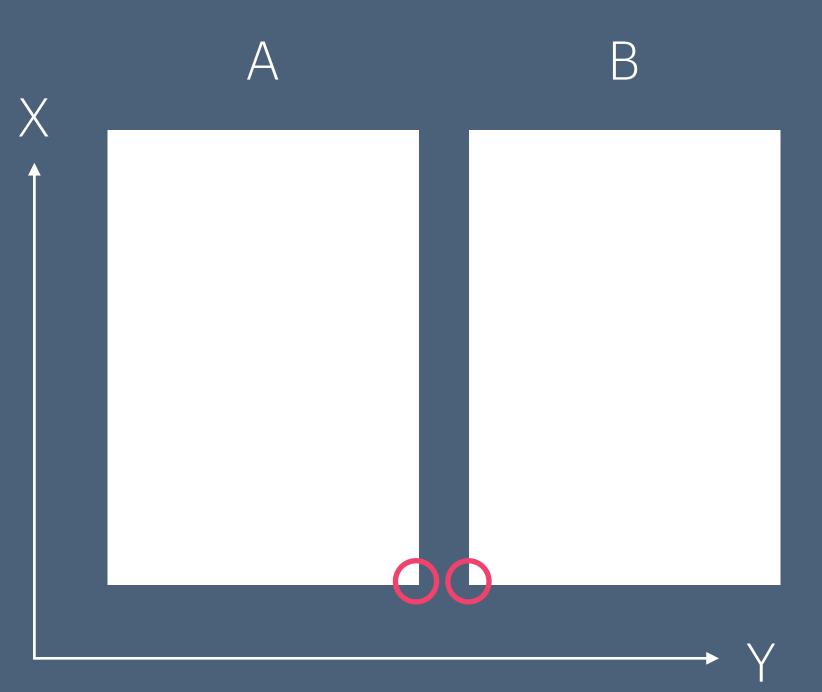
- Expected gap to be centered about O
- Layer alignment explains the gap smearing
  - (Gap larger than ~100micrometer)





## Checking Pixel position

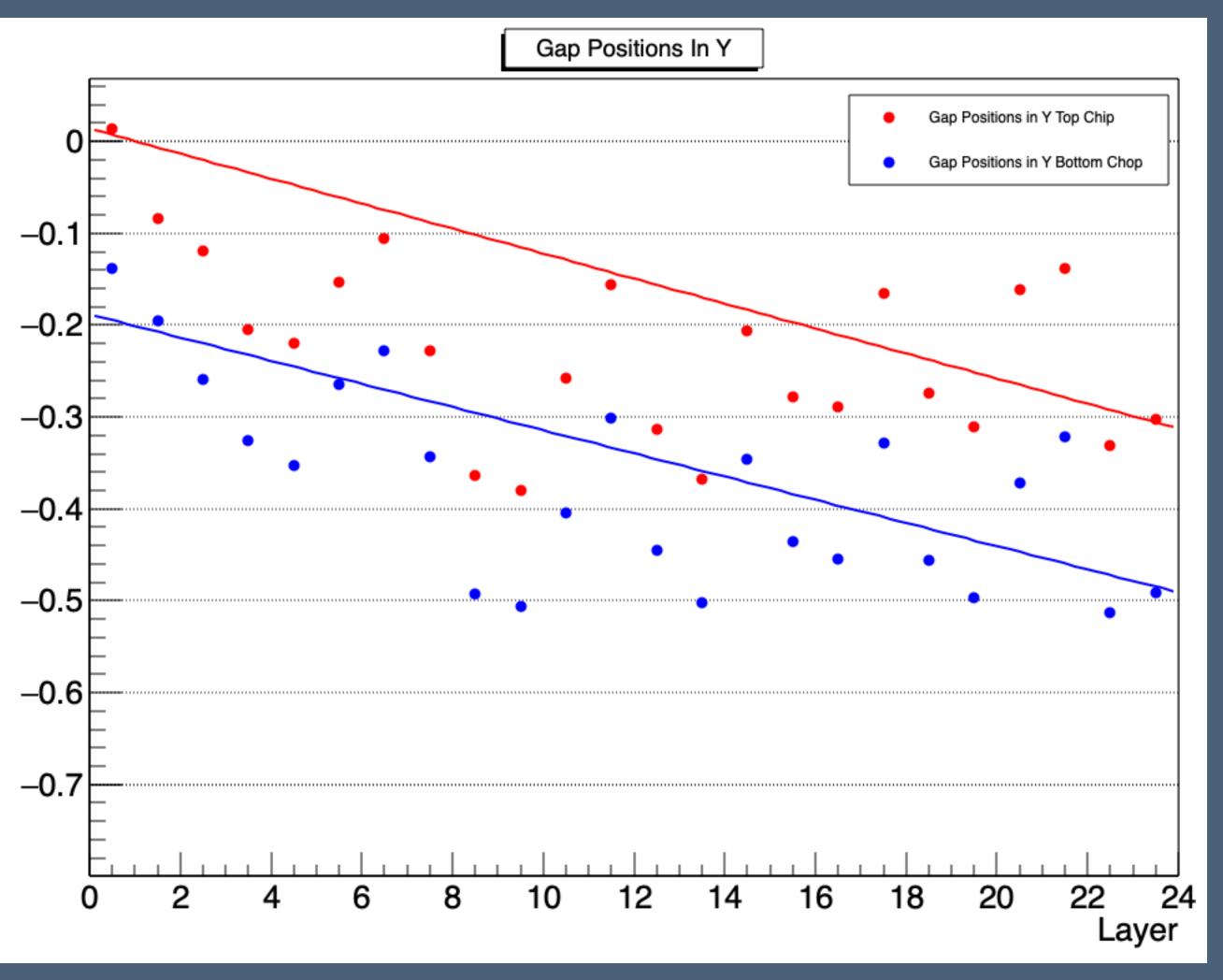
- Checking X and Y Values of pixels circled in red
- Noticed pixel on chip A was at global position Y = .01344 mm
  - This is half of the Pixel Y-size
  - Chip A ends at (X,O)
- Pixel on chip B was located at Y = .13905mm





### Yalignment

- Overall shift in the negative direction
  - (Left)

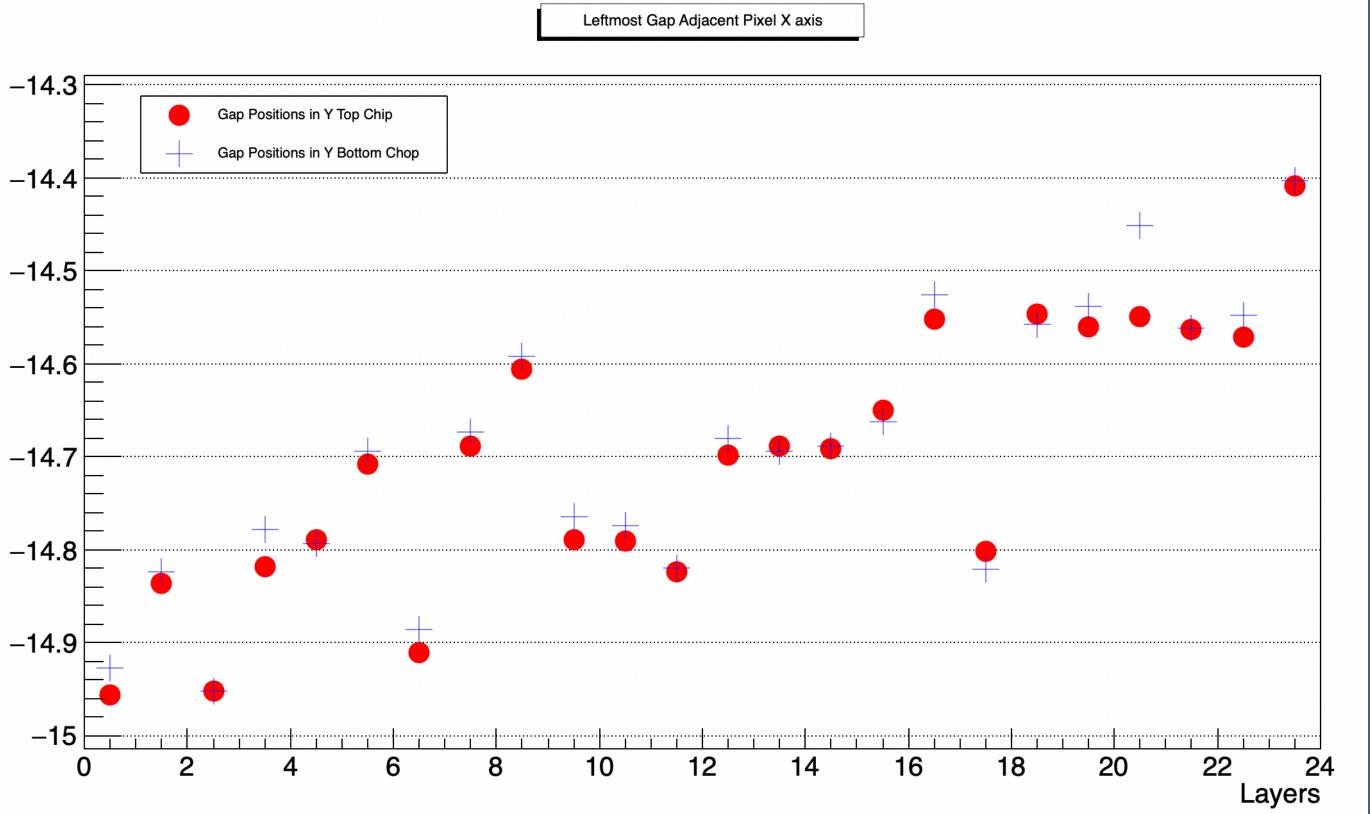


### XAlignment

• Overall shift in the positive direction

x (mm)

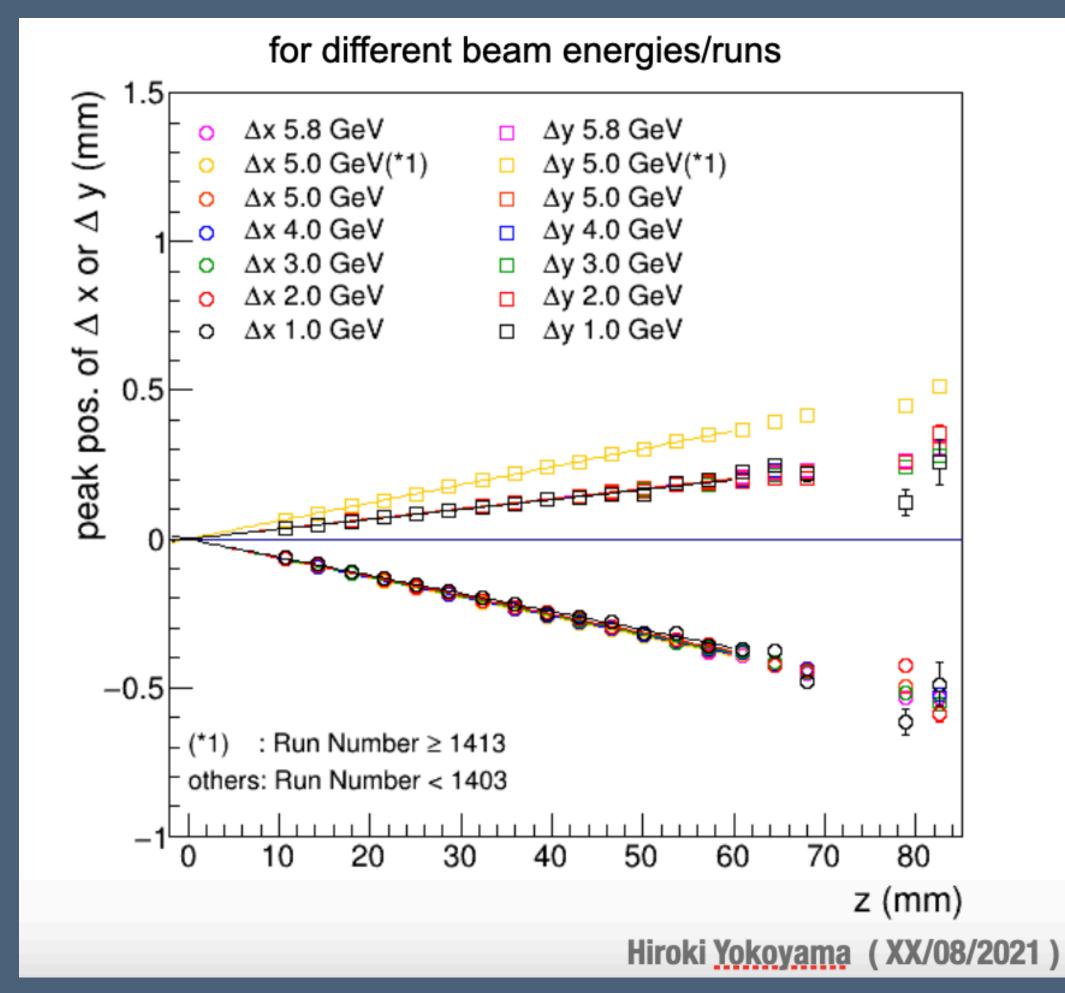
• (Up)



### Hiroki Alignment

• Hiroki slides show the alignment derived from the beam propagation on each layer

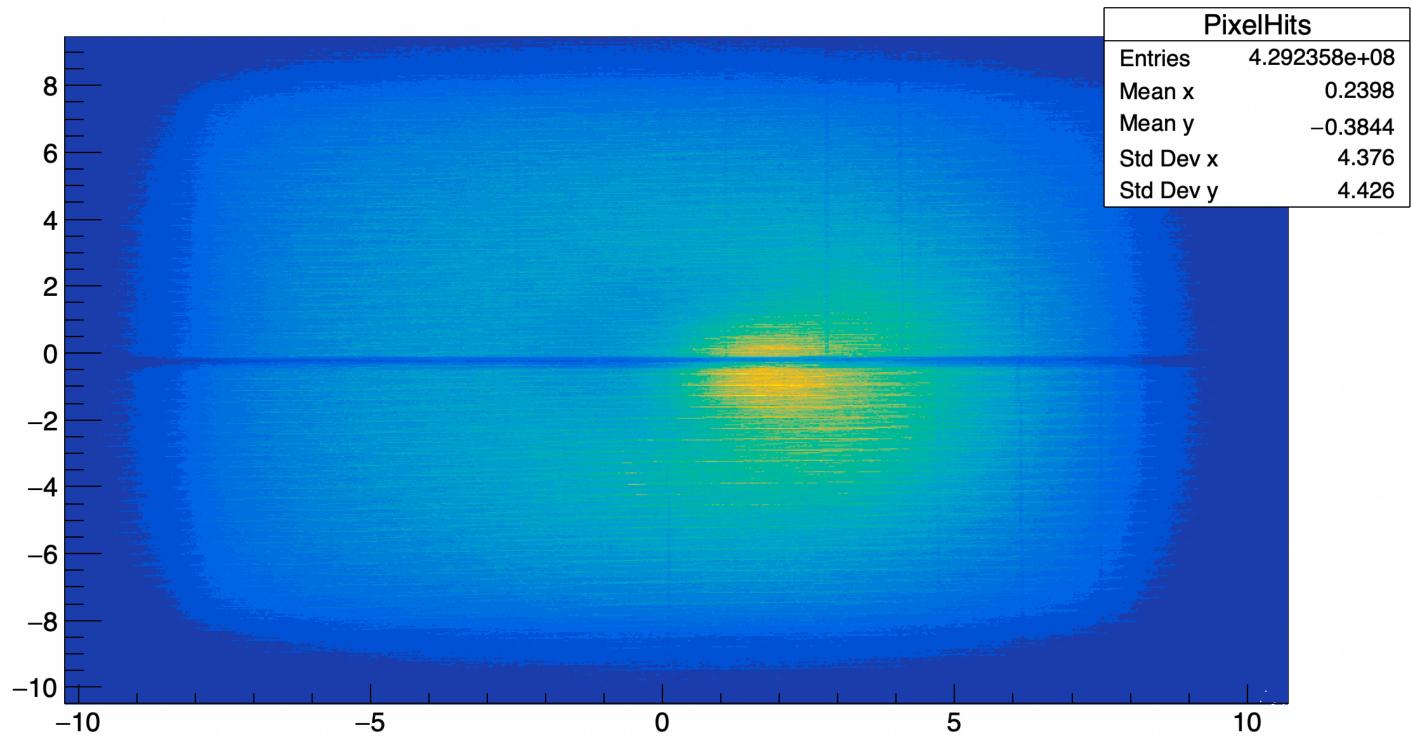
• The X and Y deviation output agrees with what we see in Hiroki slides.





#### Should we account for center in analysis?

- Since the true zero Y value is at the bottom of a chip and not in the center of the gap
  - Should we take this into consideration?
- Certainly means analysis should not extend to the outer .5mm



#### Pixel Hits all layers