

# Update of cosmic-ray analysis with 3D-line fit

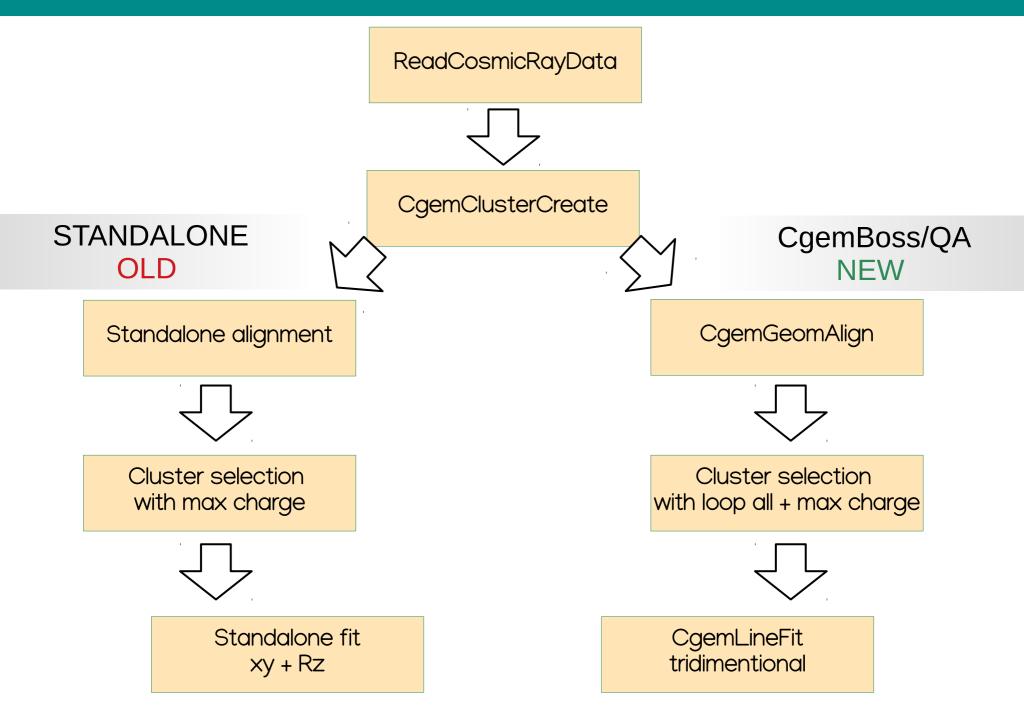


# Quality assurance procedure in CgemBoss for cosmic-ray analysis

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1

### The two procedures



#### Validation with run 17

234154 total events

### **Statistics**

|           | STANDALONE |        | CgemBoss   |        |
|-----------|------------|--------|------------|--------|
| L1 bottom | n fitted 1 | 123607 | n fitted 2 | 153167 |
| L1 top    | n fitted 1 | 124799 | n fitted 2 | 151822 |
| L2 bottom | n fitted 1 | 65932  | n fitted 2 | 150589 |
| L2 top    | n fitted 1 | 69937  | n fitted 2 | 153432 |

#### STANDALONE

Selection (on trackers):

- Three trackers fired
- Total cluster charge

L1, x view, 
$$Q_{cluster} > 20$$
 fC  
L2, x view,  $Q_{cluster} > 15$  fC

L1, v view,  $Q_{CLUSTER} > 10 \text{ fC}$ L2, v view,  $Q_{CLUSTER} > 10 \text{ fC}$ 

- No cut on cluster size
- L1 2D-clusters shall not share the same v cluster

*n fitted*  $1 \rightarrow$  the track is fitted (no limit on chi2)

#### CgemBoss

Selection (on trackers):

- Three trackers fired
- No cut on charge
- No cut on cluster size
- Loop all + max Q

#### *n fitted 2* $\rightarrow$ the track is fitted, with chi2 < 2000

The **Loop all + max Q** method, which loops on all combinations of highest charged clusters to find the usable ones, provides higher statistics

### New QA in CgemBoss

#### **Quality Assurance**

Two packages:

- **TestTrack**: all the hit/ + cluster 1D + cluster 2D + fitted track are saved to a TTree (root file)
- CgemCosmicRavOA reads the TTree and fills all the histograms

/[BESIII]/CgemBossCvs/Reconstruction/CgemLineFit/src/CgemCosmicRayQA.cxx

#### HISTOGRAMS

**HIT** – for each plane, each sheet, each view hit charge (fC), hit time (ns), hit charge (fC) vs stripID, hit time (ns) vs stripID, hit charge (fC) vs time (ns), hit charge (fC) vs length (mm)

CLUSTER 1D – for each plane, each sheet, each view number of cluster1d, cluster1d size, cluster1d charge (fC) vs phi (deg)

#### TRACK and RESIDUALS

number of fitted tracks fitted track chi2 fitted track point of closest approach x (mm) test plane: residual in R \* phi (mm) test plane: residual in z (mm) residual in R \* phi (mm) on each plane/sheet residual in z (mm) on each plane/sheet

CLUSTER 2D – for each plane, each sheet number of cluster2d cluster2d charge (fC) vs phi (deg) cluster2d charge (fC) vs z (mm)

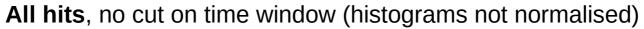
#### SIGNAL

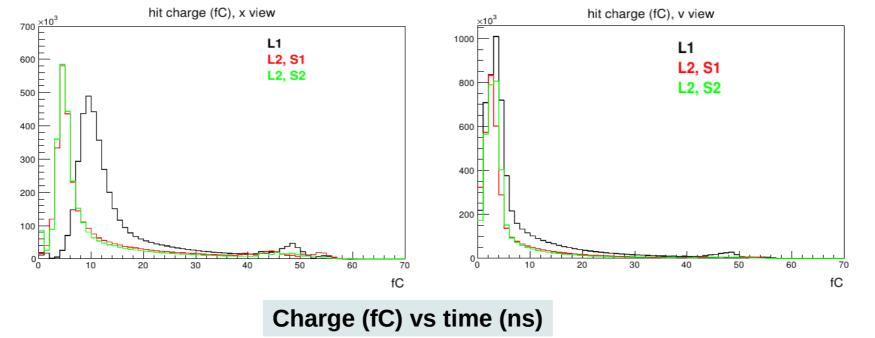
total charge of the signal (fC) charge x of the signal (fC) charge v of the signal (fC) cl.size x of the signal cl.size v of the signal

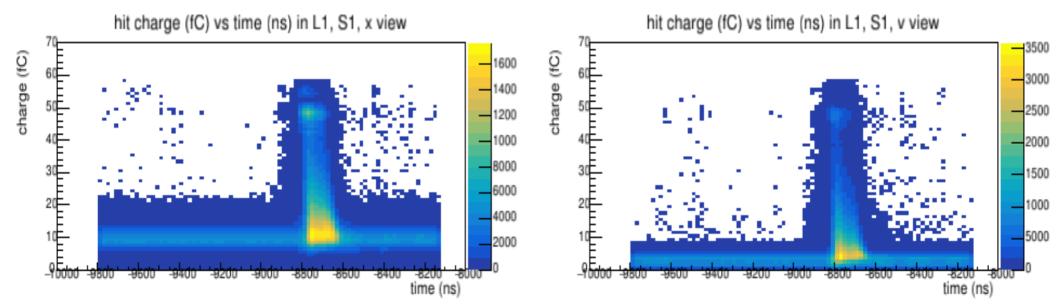
#### BACKGROUND

total charge of the background (fC) charge x of the background (fC) charge v of the background (fC) cl.size x of the background cl.size v of the background

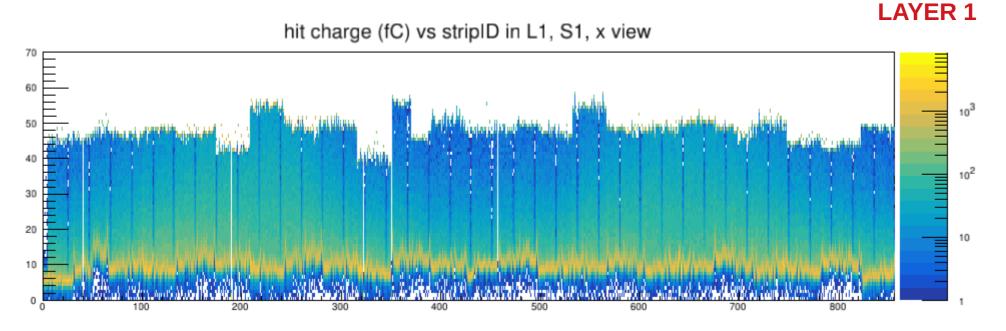
# hit charge (fC)



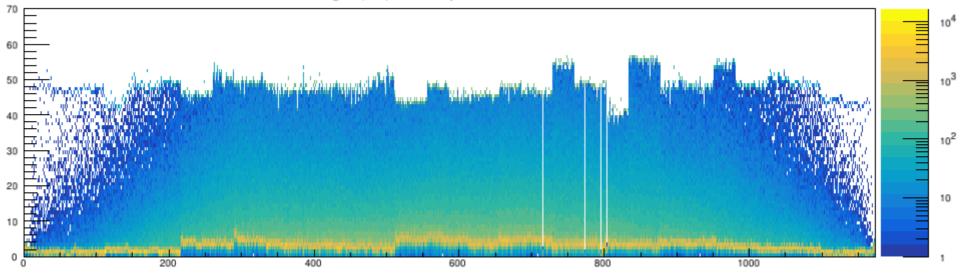




## hit charge (fC) vs strip ID

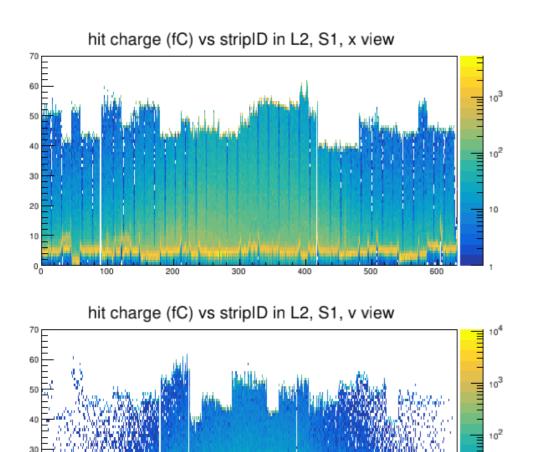


hit charge (fC) vs stripID in L1, S1, v view

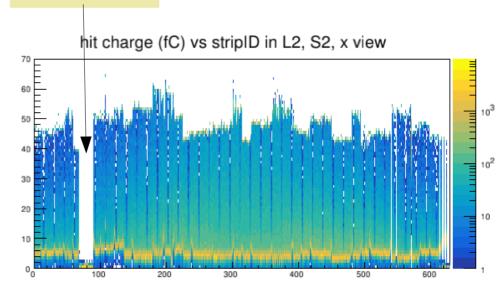


### hit charge (fC) vs strip ID

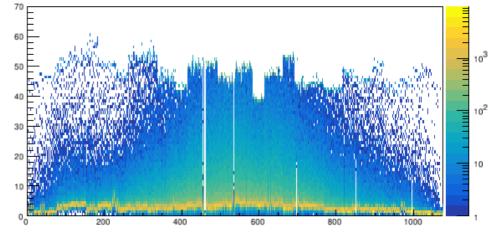
#### LAYER 2



#### broken tail



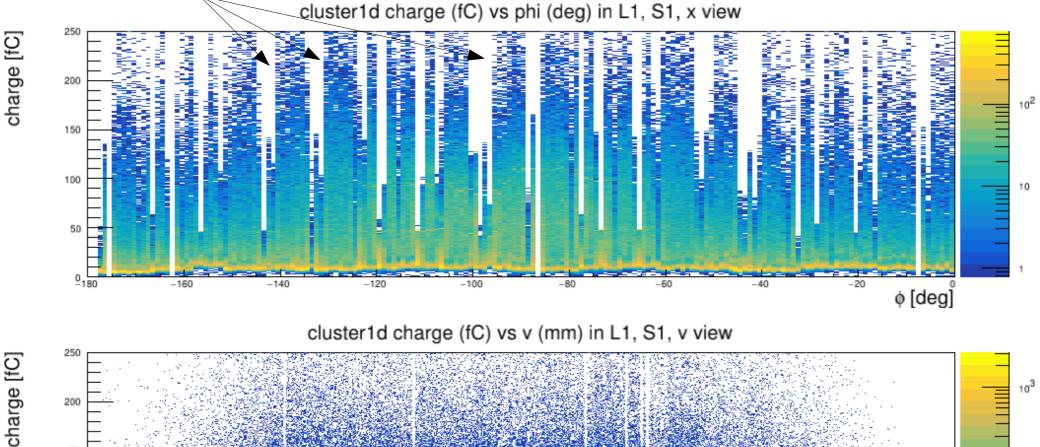
hit charge (fC) vs stripID in L2, S2, v view

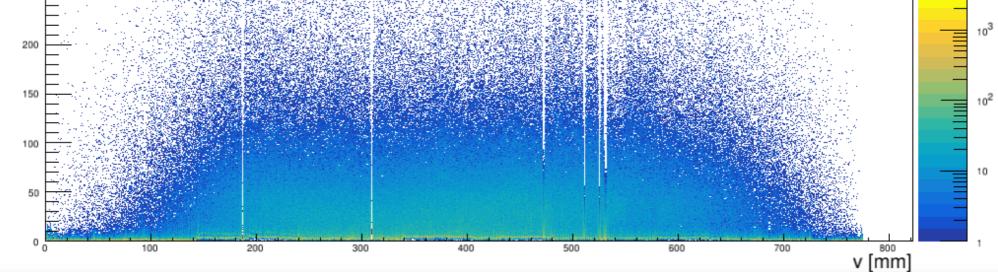


#### 1D cluster charge (fC) $vs \phi/v$

LAYER 1

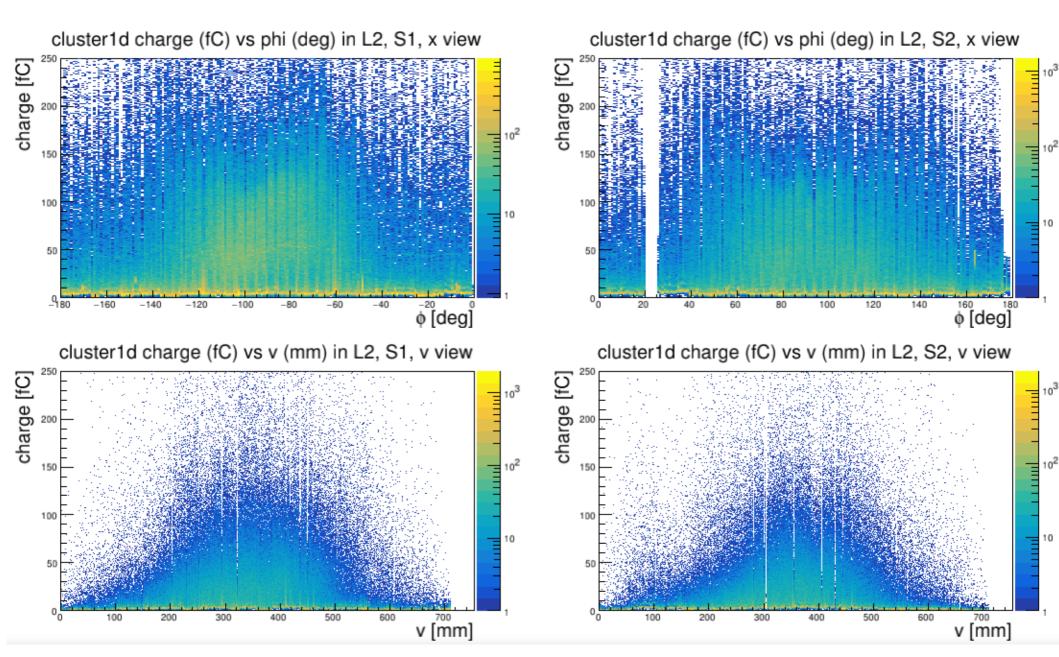
#### Microsectors register less charge



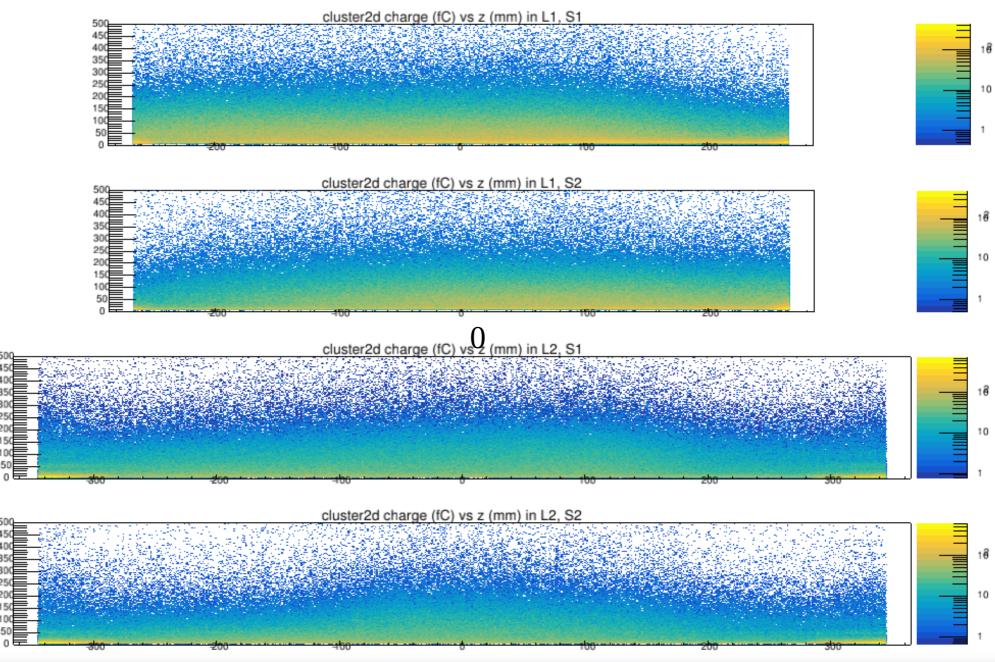


#### 1D cluster charge (fC) $vs \phi/v$

#### LAYER 2

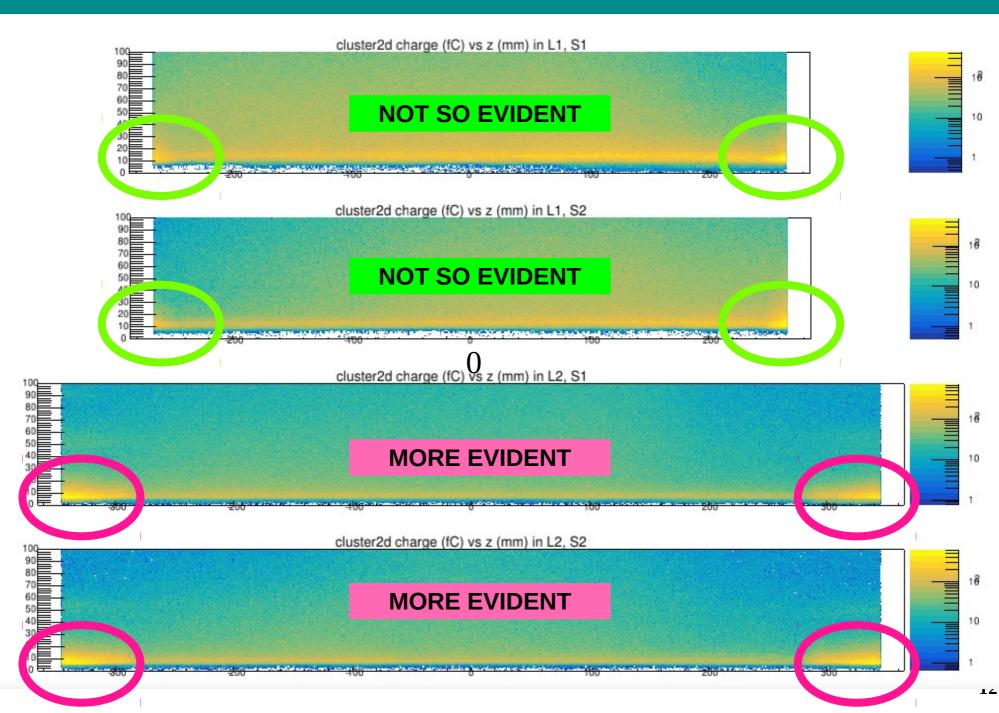


### 2D cluster charge (fC) vs z (mm)



AYER

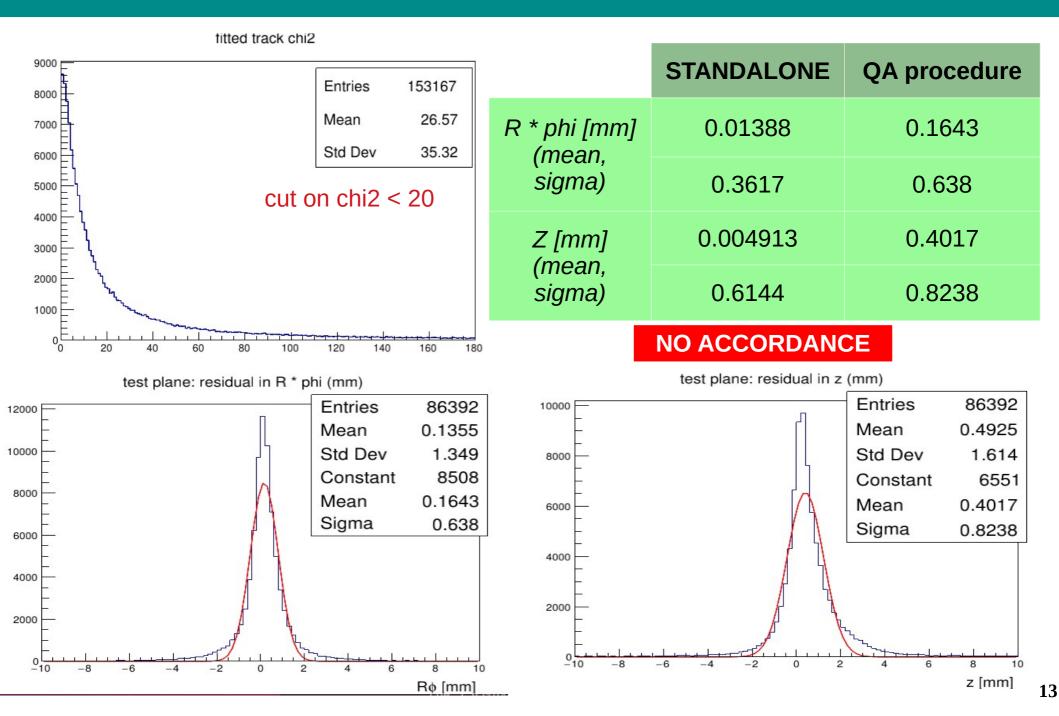
## 2D cluster charge (fC) vs z (mm) - ZOOM



-AYER 2

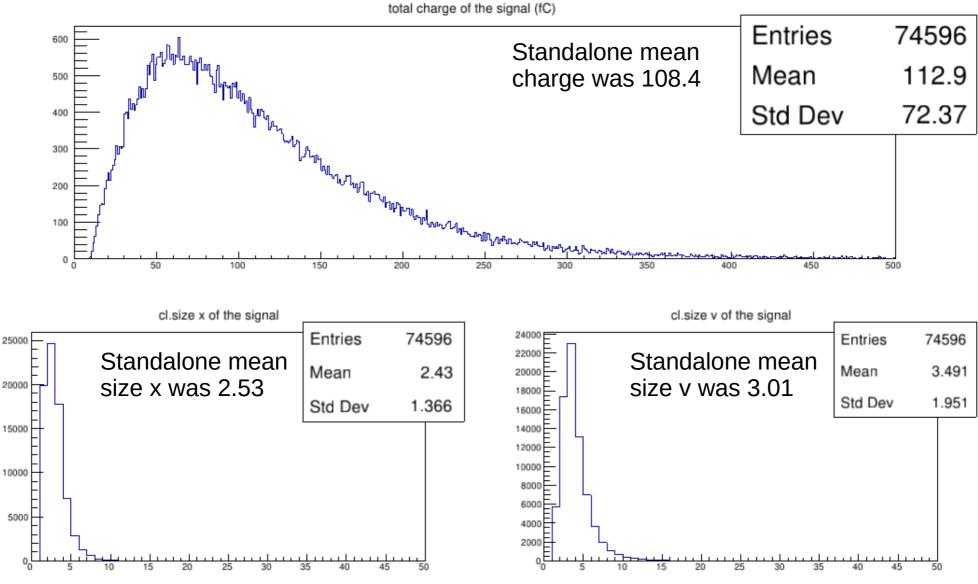
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### Reconstructed track – residual distro – L1 bottom



### Reconstructed track – signal – L1 bottom

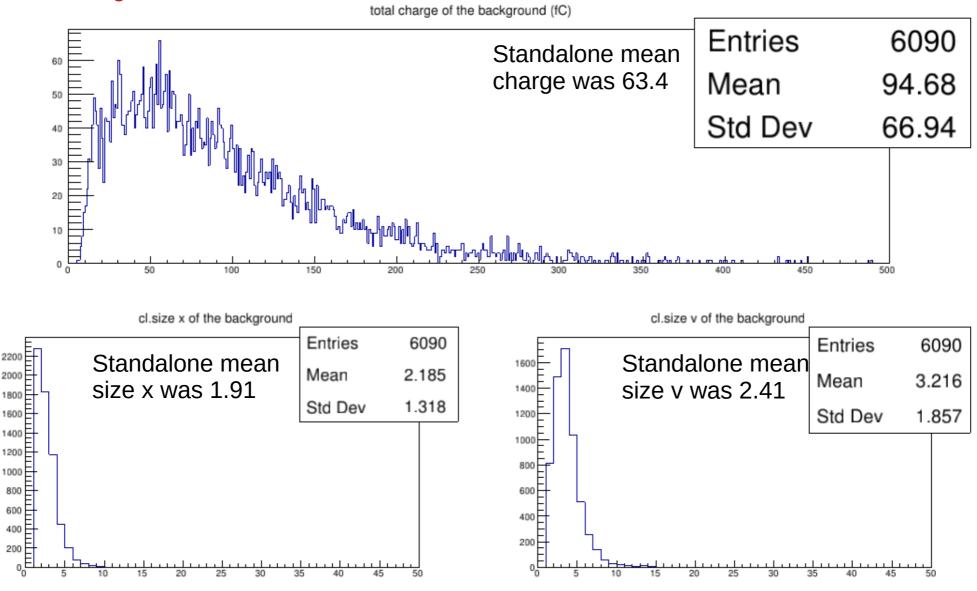
#### Inside 5 sigma



The accordance is not so bad...

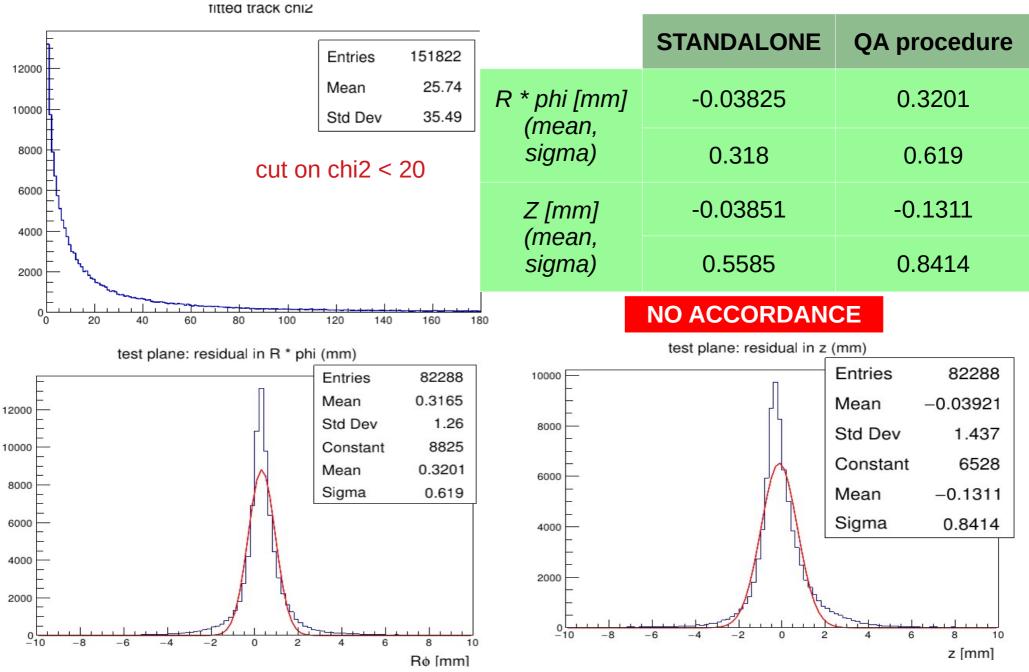
### Reconstructed track – background – L1 bottom

#### outside 10 sigma



The accordance is **not good**...

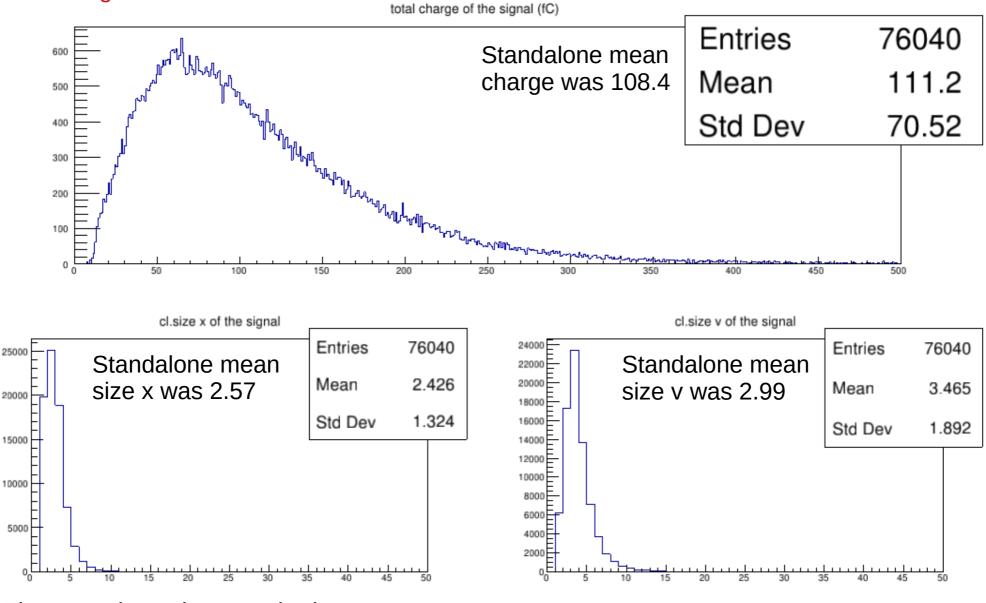
### Reconstructed track – residual distro – L1 top



16

### Reconstructed track – signal – L1 top

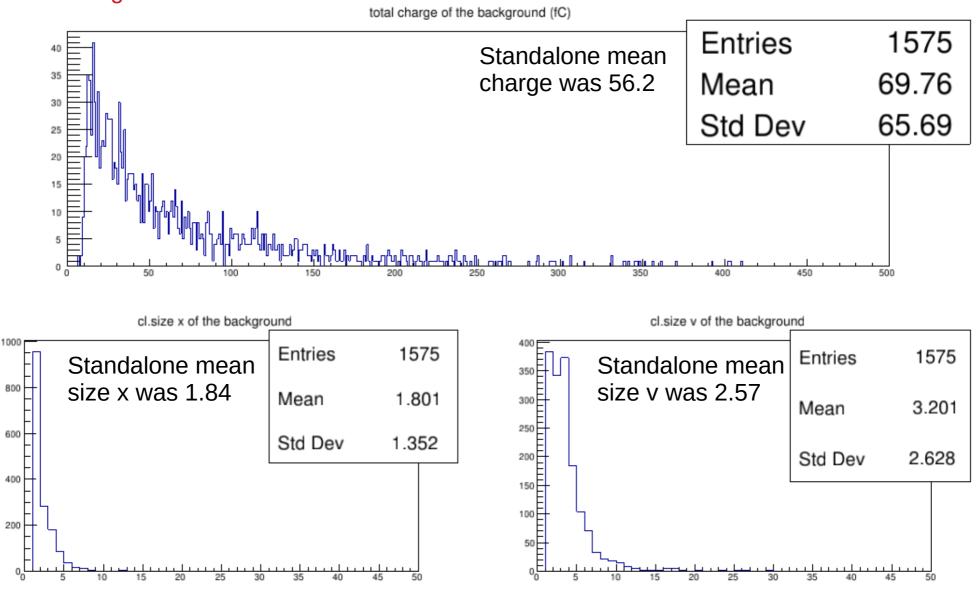
#### Inside 5 sigma



The accordance is not so bad...

### Reconstructed track – background – L1 top

#### outside 10 sigma



The accordance is **not good**...

### Reconstructed track – residual distro – L2 bottom

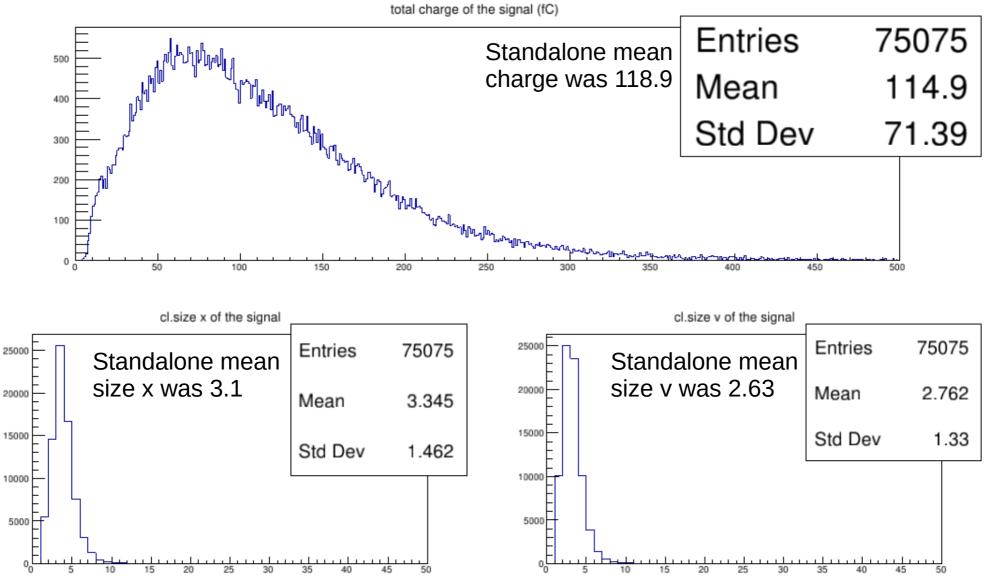
IILLEO LITACK CHIZ 9000 **QA procedure STANDALONE** Entries 150589 8000 26.17 Mean R \* phi [mm] -0.004101-0.050447000 35.22 Std Dev (mean, 6000 sigma) 0.4534 0.827 5000 cut on chi2 < 204000 0.0141 -0.09674Z [mm] 3000 (mean, 2000 sigma) 0.9902 0.8019 1000 **NO ACCORDANCE** 0<sup>L</sup> 0 20 40 60 80 100 120 140 160 180 test plane: residual in R \* phi (mm) test plane: residual in z (mm) Entries 82842 82842 Entries 8000 9000 Mean -0.01309-0.02858Mean 7000 8000 Std Dev 1.582 Std Dev 1.882 6000 7000 Constant 6671 Constant 5560 6000 5000 -0.05044Mean Mean -0.096745000 Sigma 0.827 4000 Sigma 0.9902 4000 3000 3000 2000 2000 1000 1000 -10 0 -2 2 8 -8 0 6 10 -6 -2 0 2 6 8 10 -8 -6-4 -10 z [mm]

Rø [mm]

19

### Reconstructed track – signal – L2 bottom

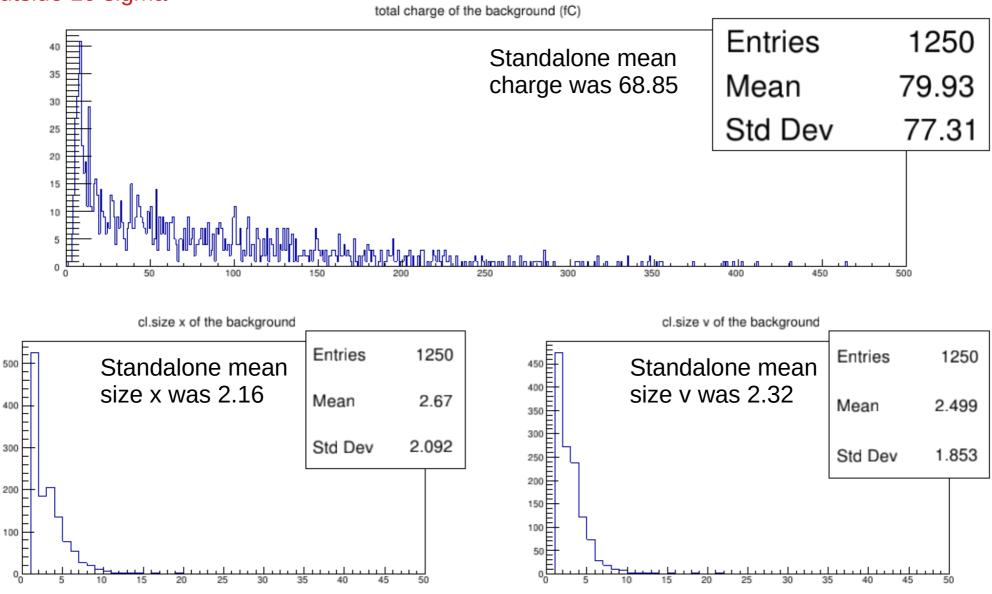
#### Inside 5 sigma



The accordance is not so bad...

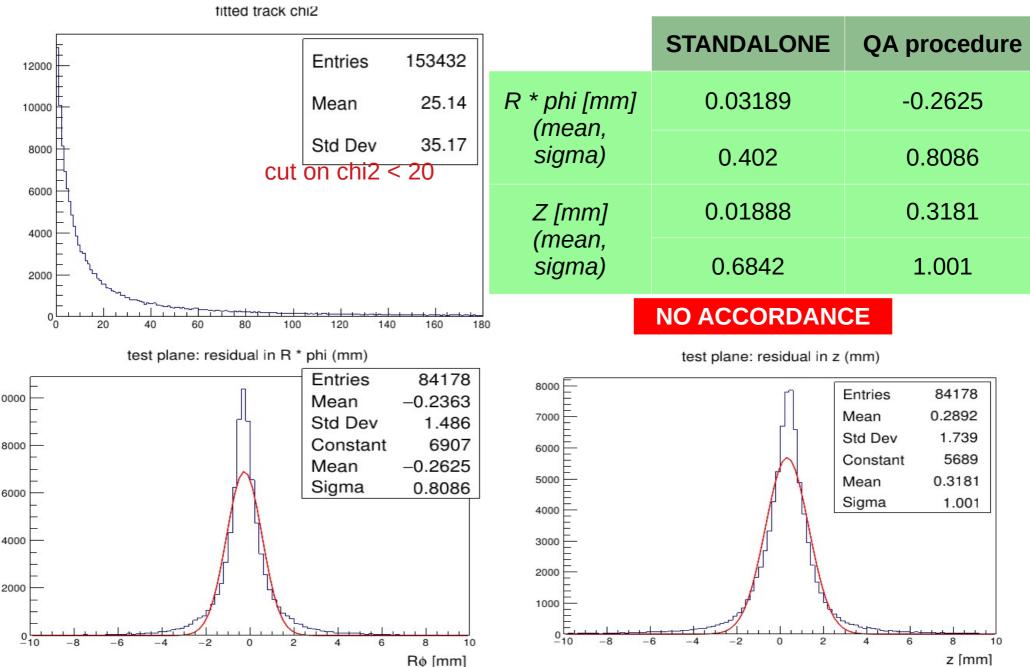
### Reconstructed track – background – L2 bottom

#### outside 10 sigma



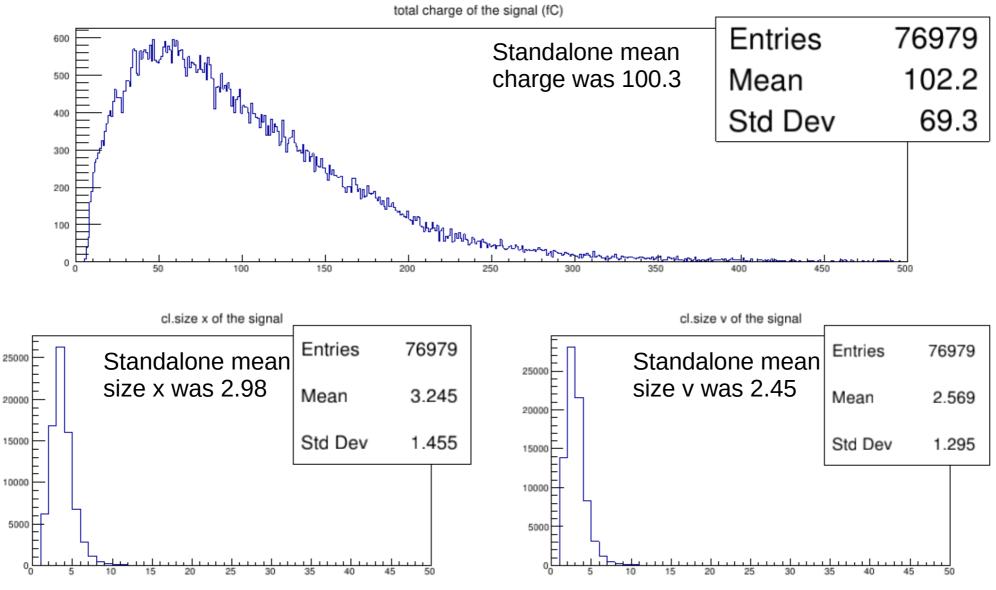
The accordance is not so bad... look a the peak @ 1 in the size

### Reconstructed track – residual distro – L2 top



### Reconstructed track – signal – L2 top

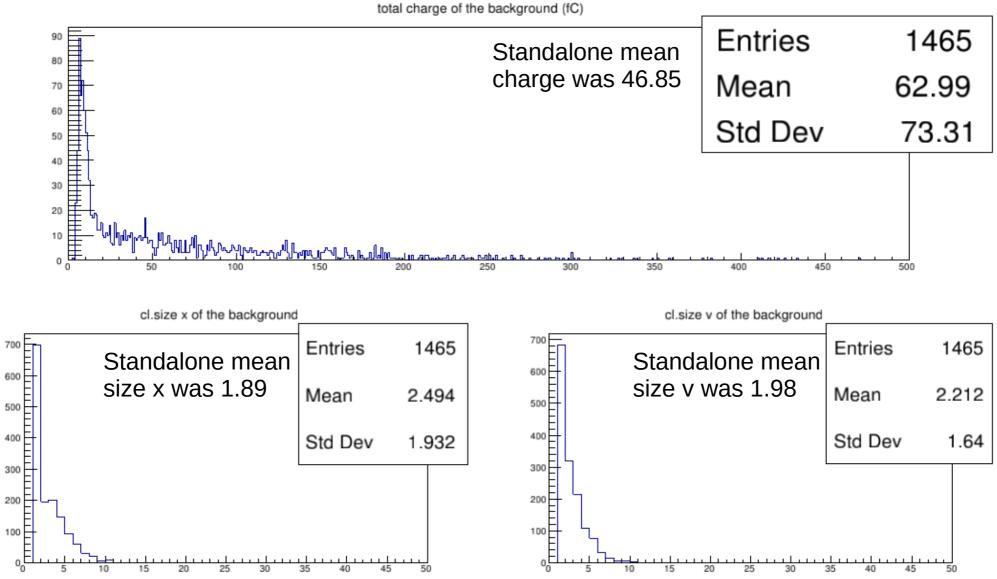
#### Inside 5 sigma



The accordance is not so bad...

### Reconstructed track – background – L2 top

#### outside 10 sigma



The accordance is not so bad... look a the peak @ 1 in the size

### Alignment?

I use the cluster 2D position (pre\_aligned) transformed this way:

HepPoint3D aligned = alignment->point\_invTransform(layer, pre\_aligned)

And used aligned for the residuals

Is this correct?

# Efficiency

| EFFICIENCY | STANDALONE | QA procedure |
|------------|------------|--------------|
| L1 bottom  | 0.86234    | 0.86346      |
| L1 top     | 0.8747     | 0.87784      |
| L2 bottom  | 0.870972   | 0.8726       |
| L2 top     | 0.866423   | 0.866998     |

The values of the efficiency are compatible

### Conclusions

- The **statistics** is enhanced by the loop all + max Q, as expected
- The histograms of the **hits, the cluster 1D and cluster 2D** are compatible with what seen before
- The **efficiency** is the same as from the standalone, around 87%
- The **residual distributions** still have some problem:
  - They are not completely Gaussian, there are tails
  - They are not centered exactly in zero (maybe I did not apply alignment correctly when computing the residual values?)

The code is on CVS (not the last changes, but I will upload them soon)