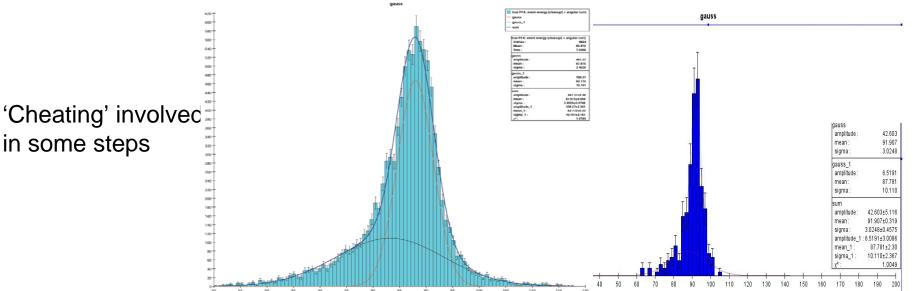
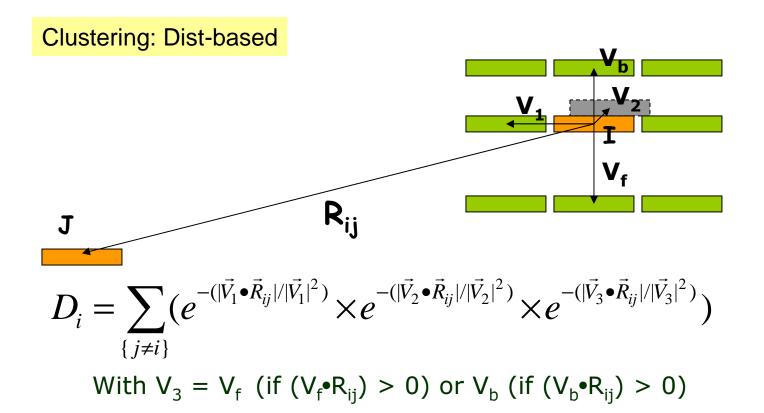
Two Density-based Clustering Algorithms

L. Xia (ANL) V. Zutshi (NIU)

General Comments

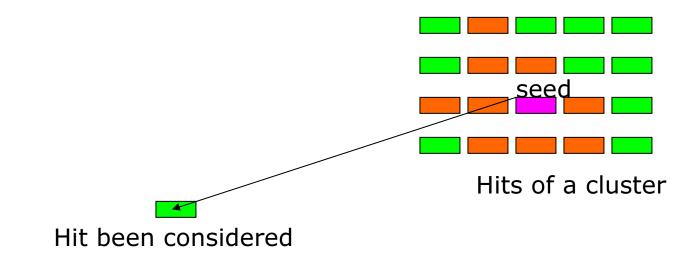
- Both are calorimeter first approaches
 clustering → track match → fragment....
- SiD geometry Si-W ECAL, RPC or Scintillator HCAL





- Hit density reflects the closeness from one hit i to a group of hits {j}
 - {j} = {all calorimeter hits} to decide if hit i should be a cluster seed
 - $\{j\} = \{all hits in a cluster\}$ to decide if hit i should be attached to this cluster
- Consider cell density variation by normalizing distance to local cell separation
 - Density calculation takes care of the detector geometry
 - Clustering algorithm then treat all calorimeter hits in the same way

Clustering: Dist-based



- Find a cluster seed: hit with highest density among remaining hits
- Attach nearby hits to a seed to form a small cluster
- Attach additional hits based on density calculation
 - $i = hit been considered, \{j\} = \{existing hits in this cluster\}$
 - EM hits, $D_i > 0.01$
 - HAD hits, D_i > 0.001
 - Grow the cluster until no hits can be attached to it
- Find next cluster seed, until run out of hits

Clustering: Grad-based

Define the <u>density</u> neighborhood of a cell i

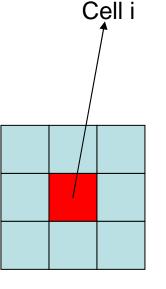
Neighborhood $\rightarrow \pm$ (nly,nz,nphi) window centered on the cell

For each cell i calculate the 'density' D_i over its neighborhood

For each i calculate the 'gradient' $(D_j - D_i)/d_{ij}$ where j is in the <u>clustering</u> neighborhood

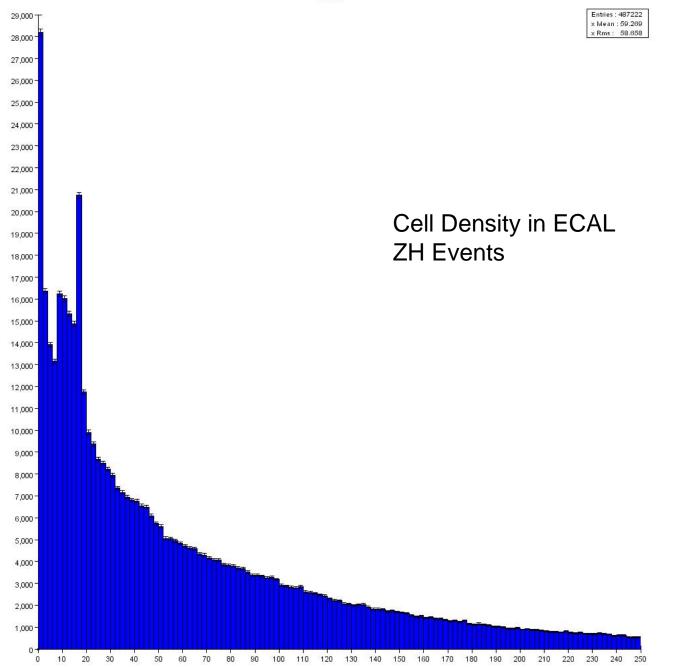
Find max [gradient]_{ij}

The magnitude and sign of max[gradient]_i determines what happens to cell i



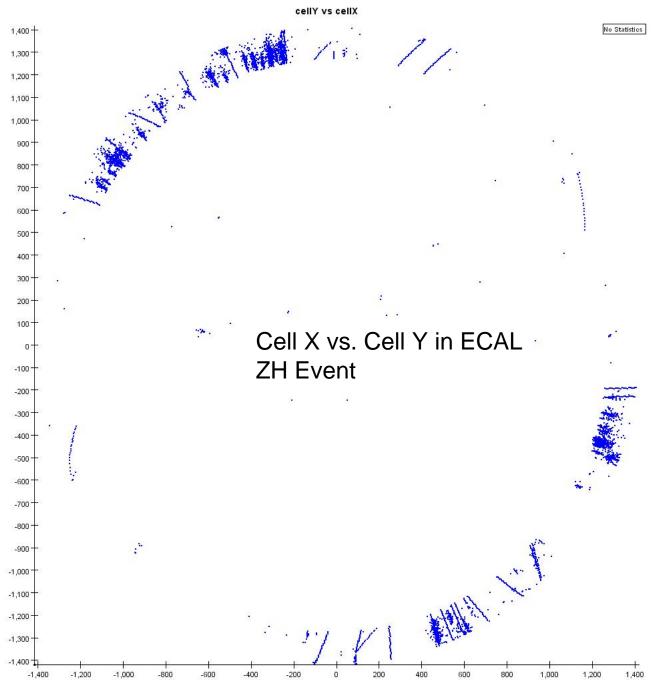
In this presentation simply occupancy or no. of cells above threshold is used

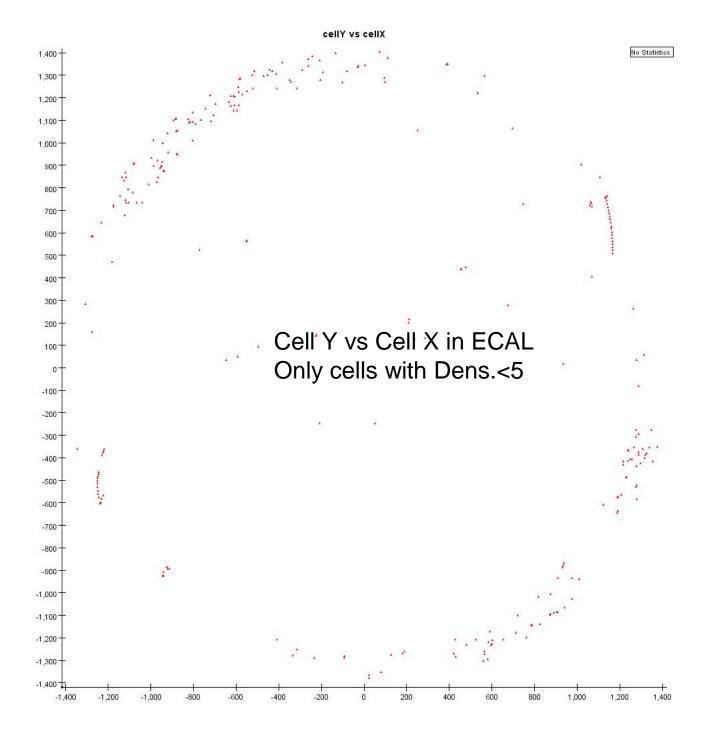
- if max[gradient]_{ij} < 0
 - i becomes the 'root' and starts a branch of its cluster
- if max[gradient]_{ij} > 0
 j is the parent of i and attaches i to its branch
- if max[gradient]_{ij} == 0
 attach i to the nearest j

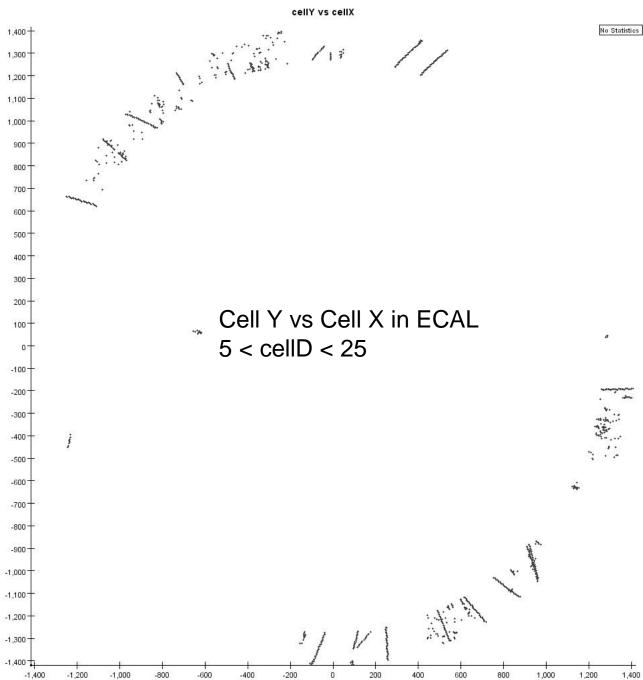


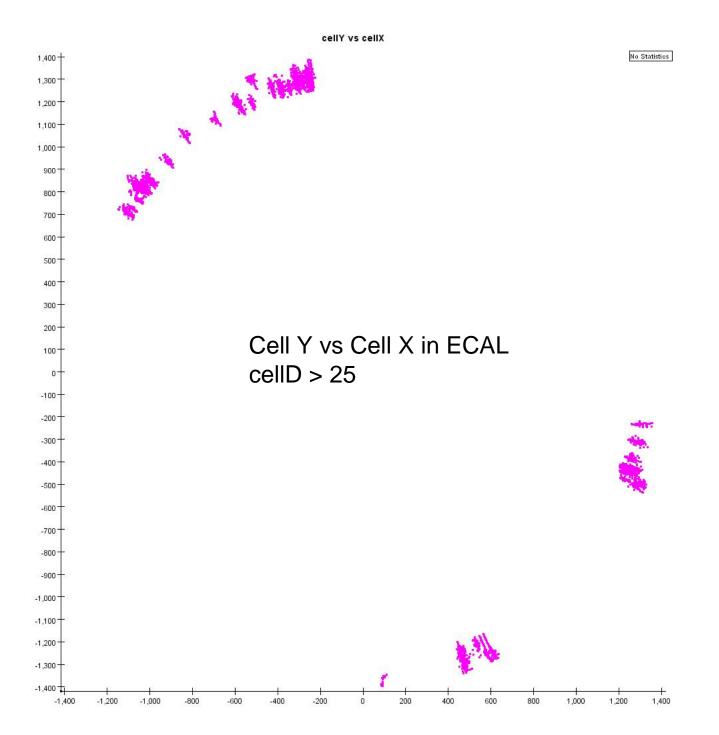
cellD

cellY vs cellX









Dist-based

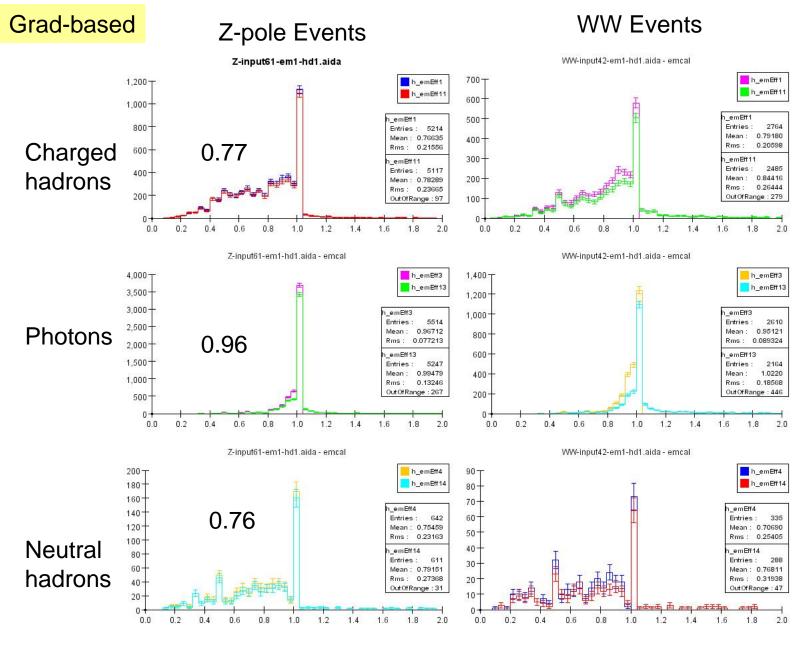
clustering efficiency: single particle

Particle	ECal hit efficiency	HCal hit efficiency	Overall hit efficiency	Overall energy efficiency
Photon (1GeV)	89%	43%	89%	91%
Photon (5GeV)	92%	54%	92%	96%
Photon (10GeV)	92%	61%	92%	97%
Photon (100GeV)	95%	82%	95%	>99%
Pion (2 GeV)	78%	59%	75%	71%
Pion (5 GeV)	81%	70%	79%	80%
Pion (10GeV)	84%	80%	83%	85%
Pion (20GeV)	85%	87%	88%	91%

•Typical electron cluster energy resolution ~ 21%/sqrt(E)

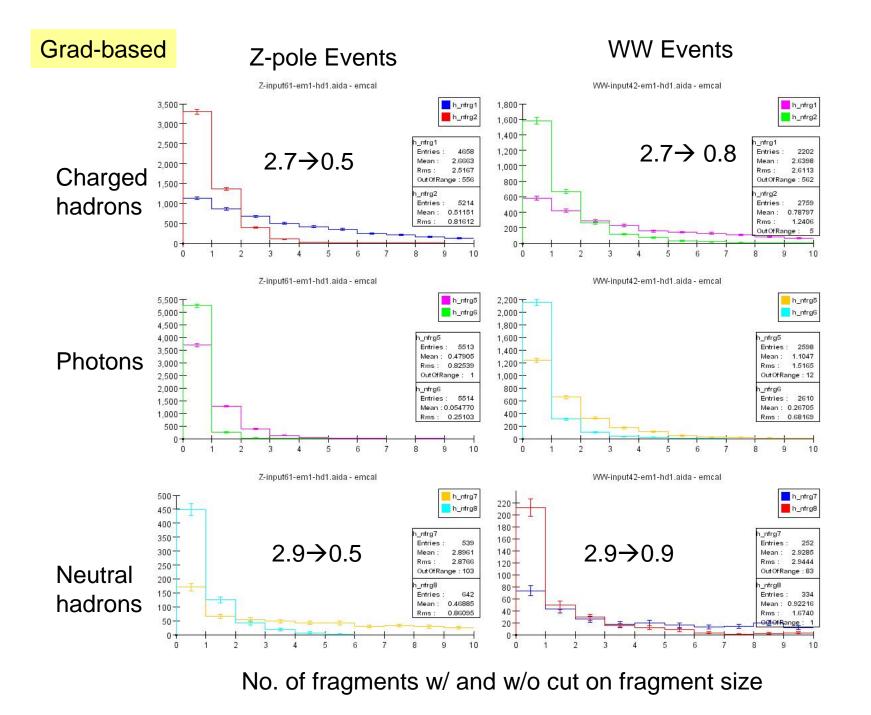
•Typical pion cluster energy resolution ~70%/sqrt(E)

•All numbers are for one main cluster (no fragments are included)

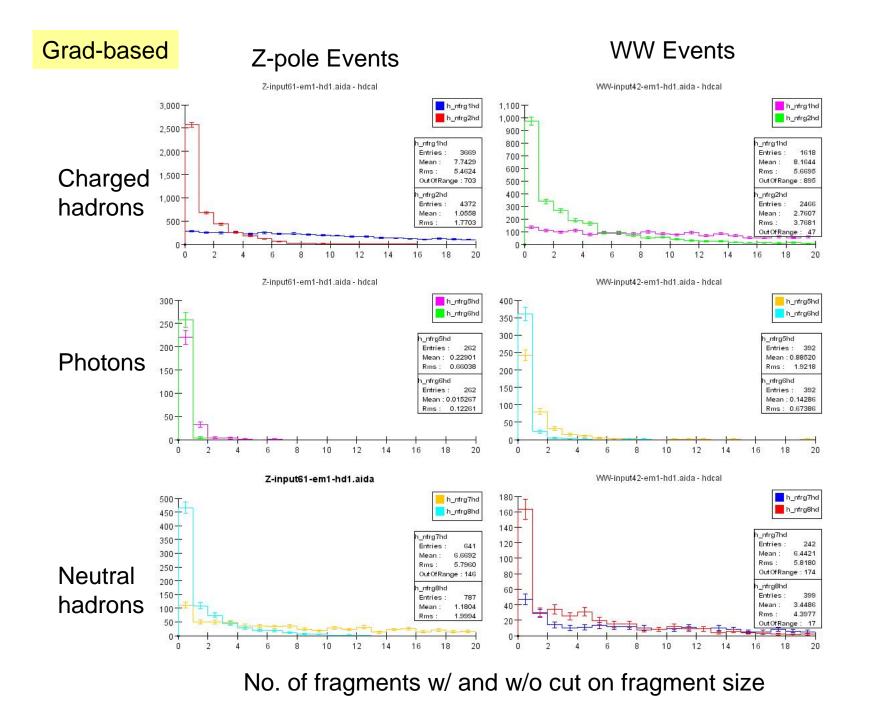


Cluster Efficiency in Z-pole events

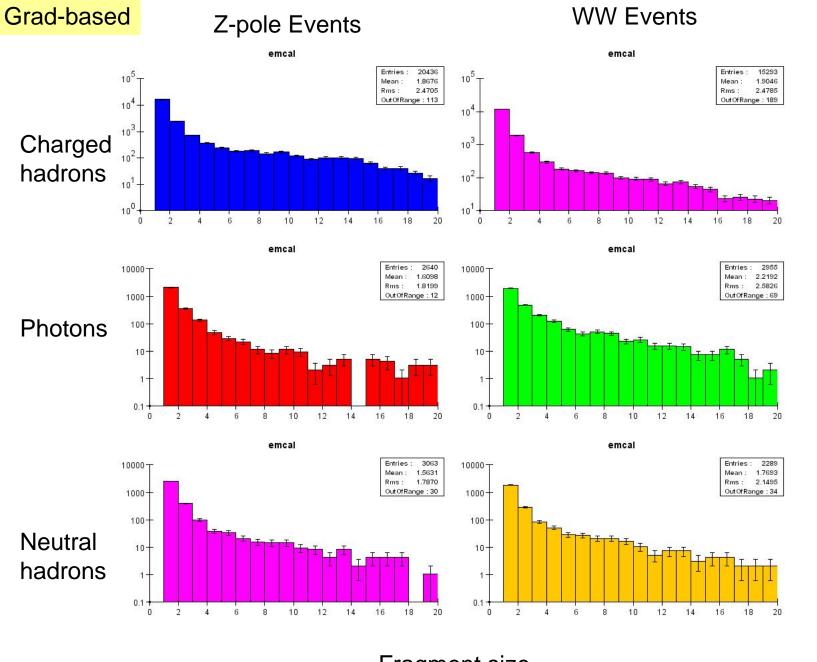
ECAL



ECAL



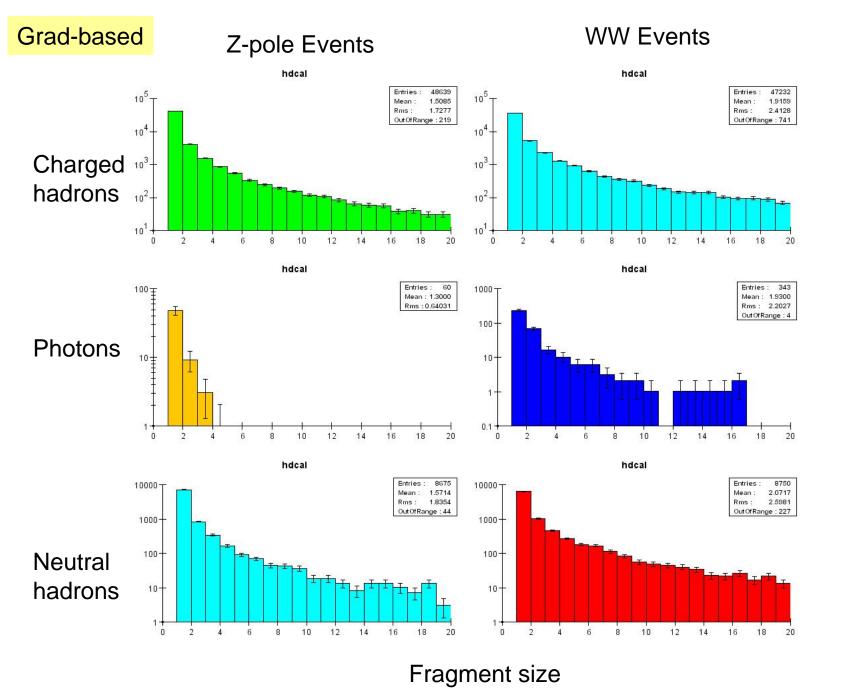




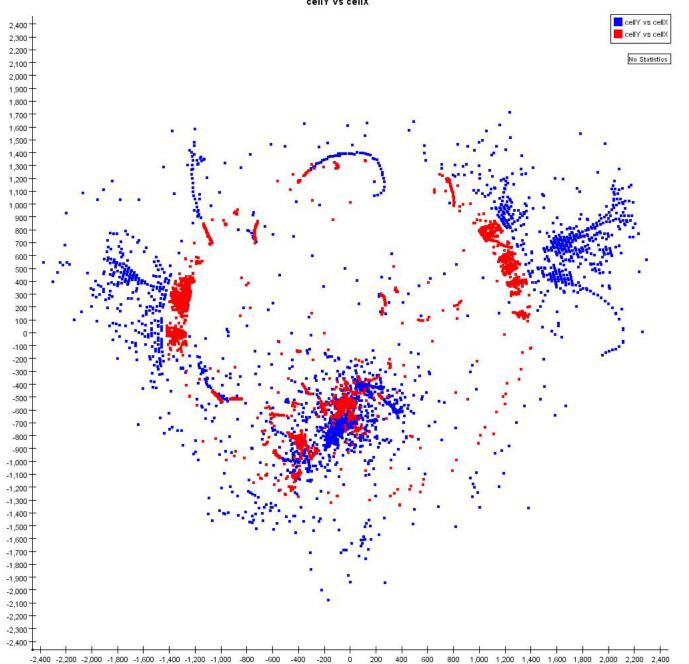
]

ECAL

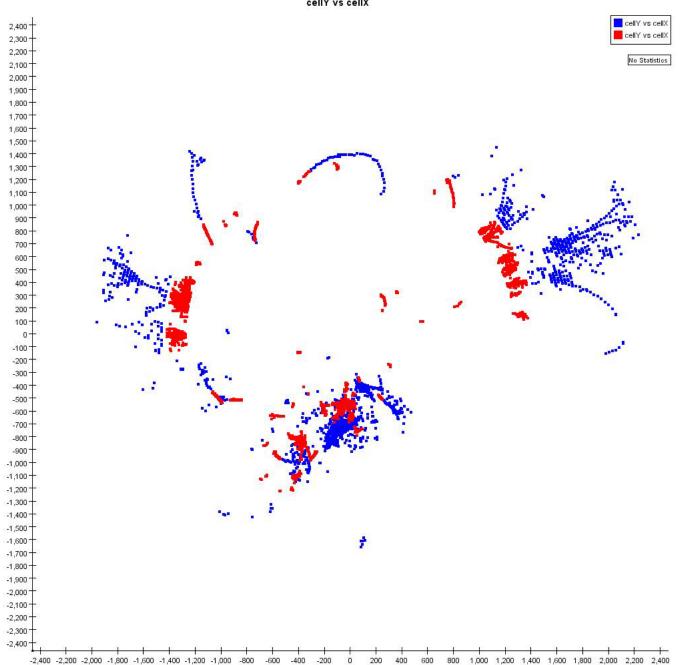
Fragment size



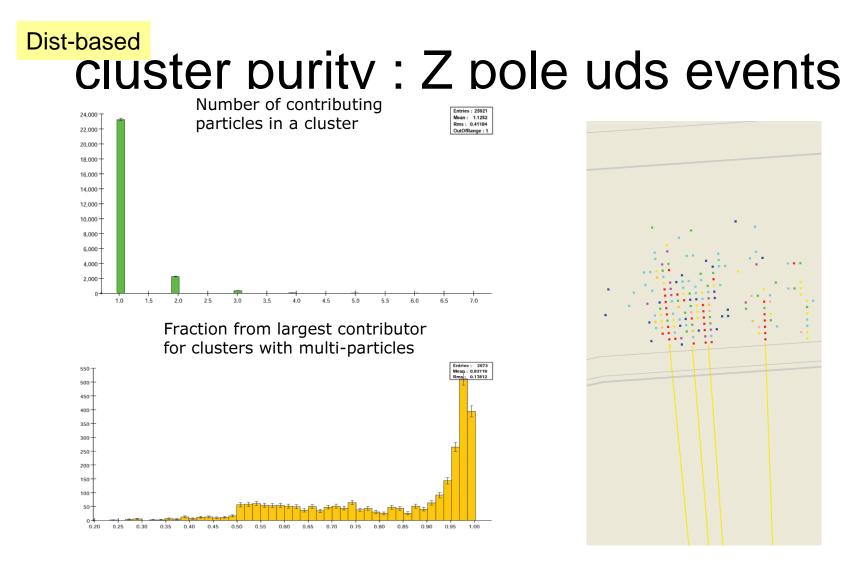
HCAL



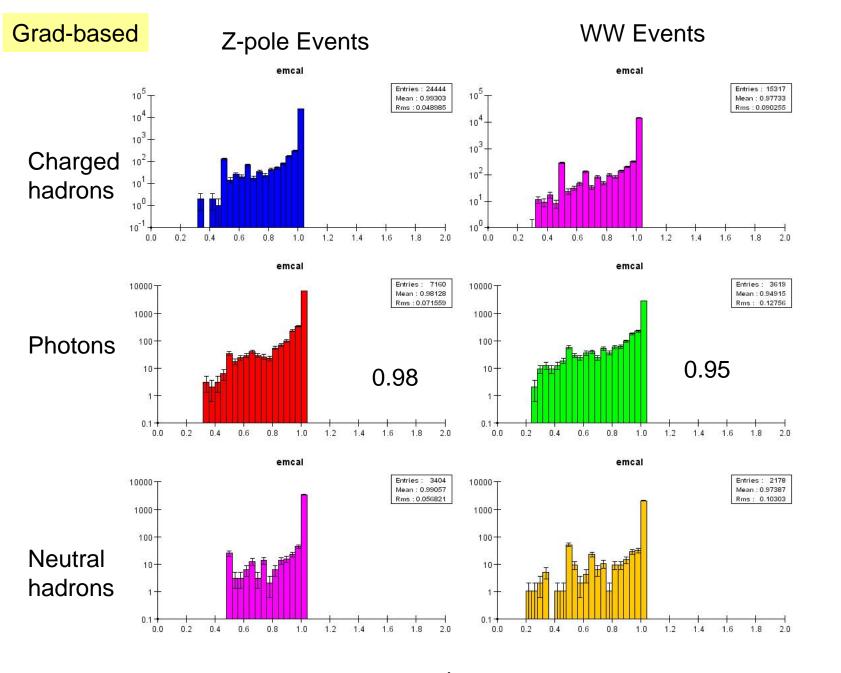
cellY vs cellX



cellY vs cellX

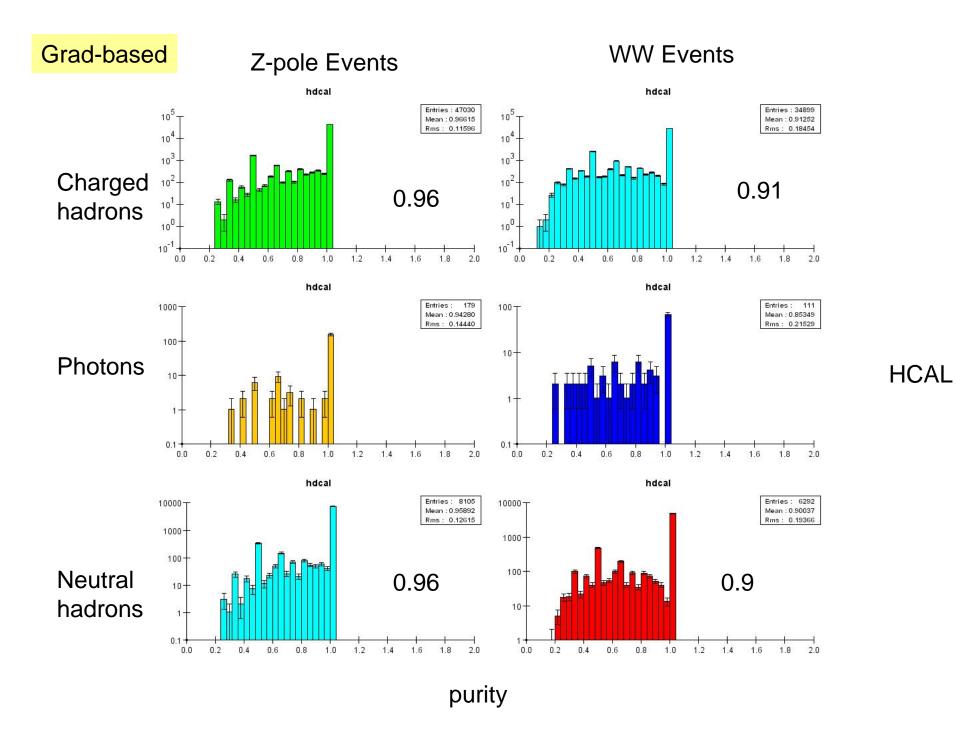


- Most of the clusters (89.7%) are pure (only one particle contributes)
- For the remaining 10.3% clusters
 - 55% are almost pure (more than 90% hits are from one particle)
 - The rest clusters contain merged showers, part of them are 'trouble makers'
- On average, 1.2 merged shower clusters/Z pole event



ECAL

purity

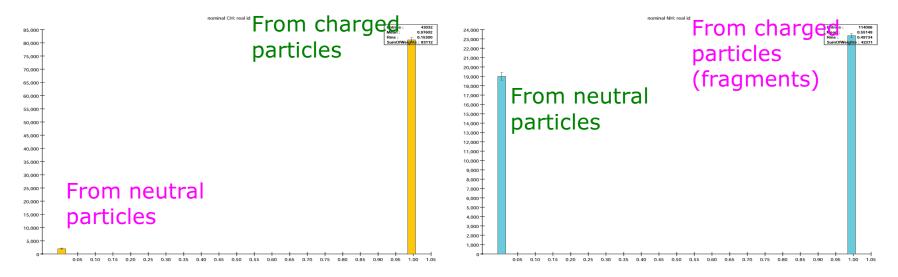


Dist-based

After track-cluster matching

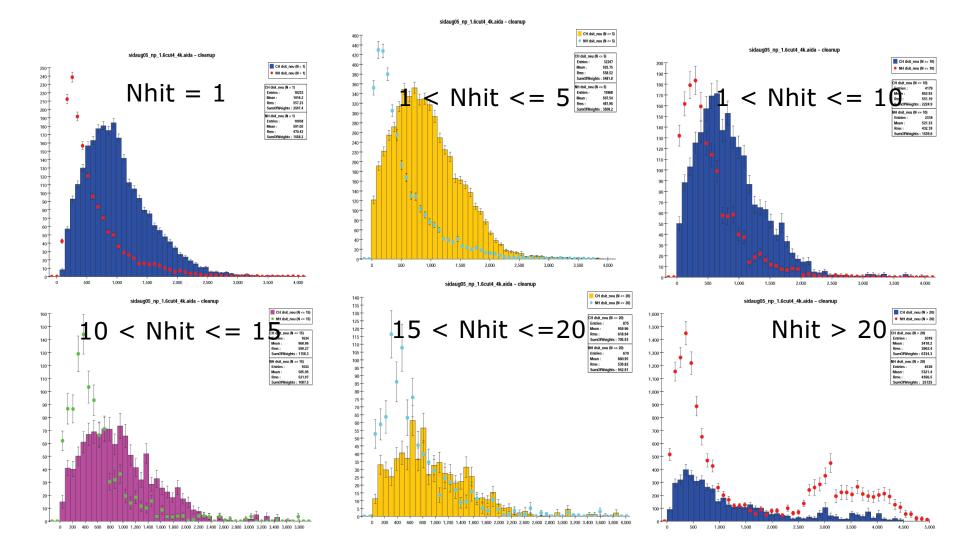
Energy of matched clusters

Energy of clusters not matched to any track: neutral candidate



On average ~3% came from neutral

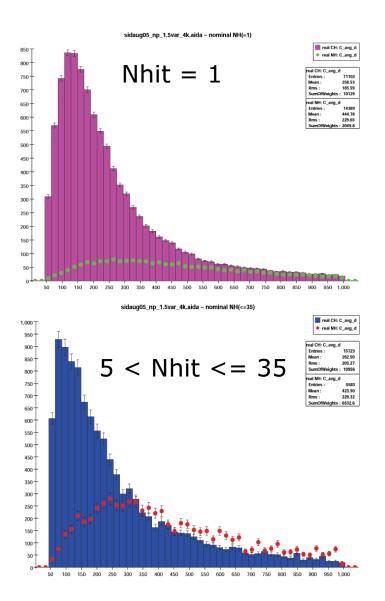
Energy from charged particles is more than real neutral -- need to work on it!

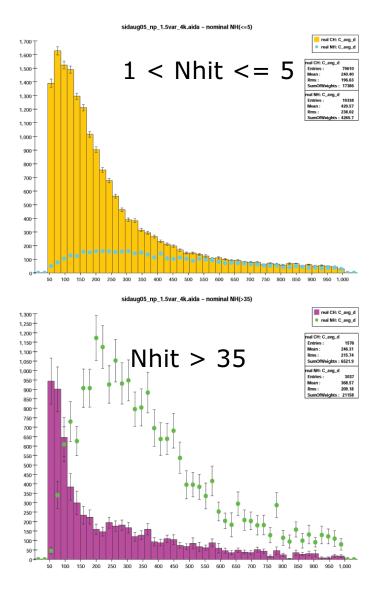


Fragment identification – variable2: distance to any neutrals

Neutral: any 'big' cluster not matched to any track Fragment cluster (histogram) vs. real neutral cluster (dots)

Fragment identification - variable1: distance to any track





Dist-based rayment identification – variable3: ratio of the two distances sidaug05_np_1.6cut4_4k.aida - cleanur sidaug05 np 1.6cut4 4k.aida – cleanup CH avg_d vs dsit_neu (N = 1) CH avg_d vs dsit_neu (N <= 1 600 - 580 - NH avg_d vs dsit_neu (N = 1) NH avg_d vs dsit_neu (N <= 1 CH avg_d vs dsit_neu (N = 1 Nhit = 11 < Nhit <= 5 1 < Nhit <= 0.9097 2599.4 SumOfWeig NH avg_d vs dsit_neu (N = 1) Entries : 10852 Mean : 1,9395 Rms : 1,4894 SumOfWeights : 1545.9 NH avg. d vs duit, new (N -= 5) 1,250 1.200 1.150 1,100 400 1.050 950 850 -100 -750 -100 -100 -550 -450 -450 -350 -150 -150 -250 200 150 100

15 < Nhit <=20

sidaug05 np 1.6cut4 4k.aida - cleanur

18,000 17,000 16,000

15.000

14.000

13,000

11.00

Nhit >

4.0 4.5

5.0 5.5 6.0

2.5 3.0 3.5

6.5

CH avg_d vs dsit_neu (N <= 15)

.75265 .90816 1149.1

99 1.615(1.551)

sidaug05_np_1.6cut4_4k.aida - cleanup

10 < Nhit <=

340 -320 -300 -280 -

260-

240-

220

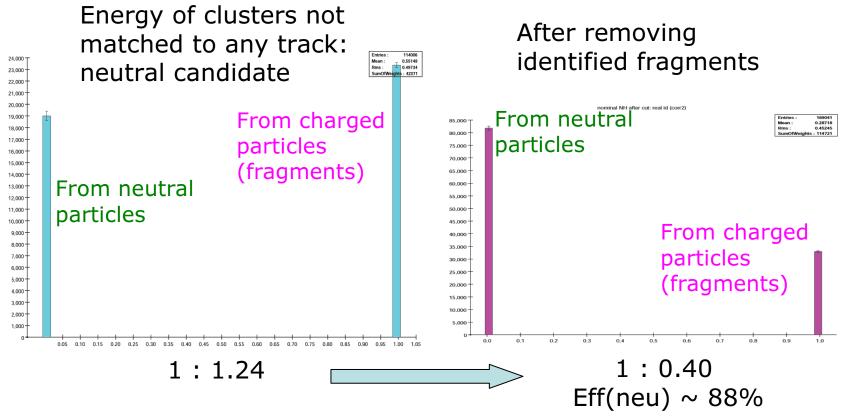
200

180 -160 -140 -120 -100 -80 -

Fragment cluster (histogram) vs. real neutral cluster (dots)

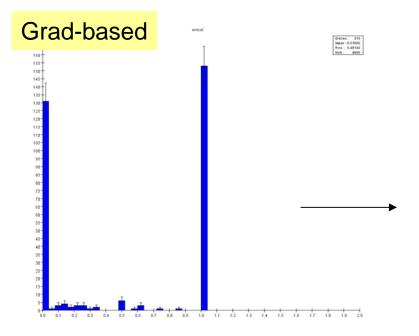
Fragment identification

Dist-based



Use the three variables to identify fragments:

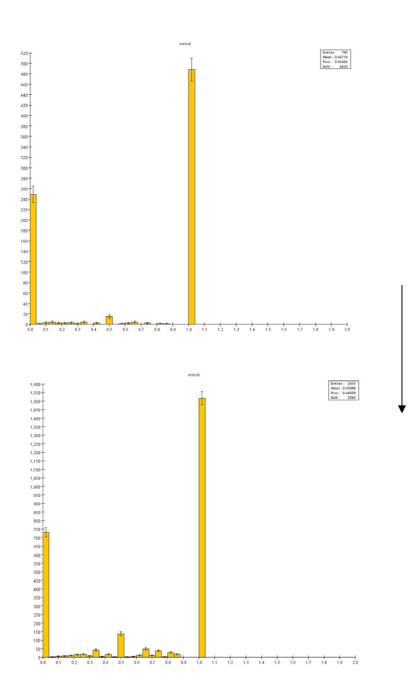
- 1. 72% of the energy from fragments is removed
- 2. Only lose 12% of real neutral energy

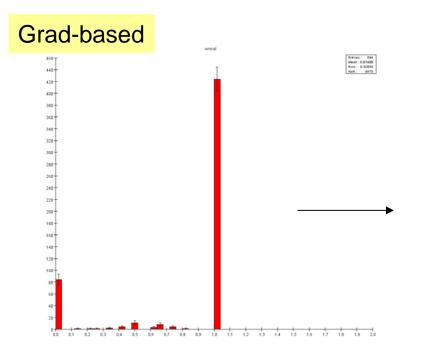


Fraction of cells attached To main cluster which belong to it Attachment is based on angular distance

Z-pole events Charged hadrons

→ Increasing dist

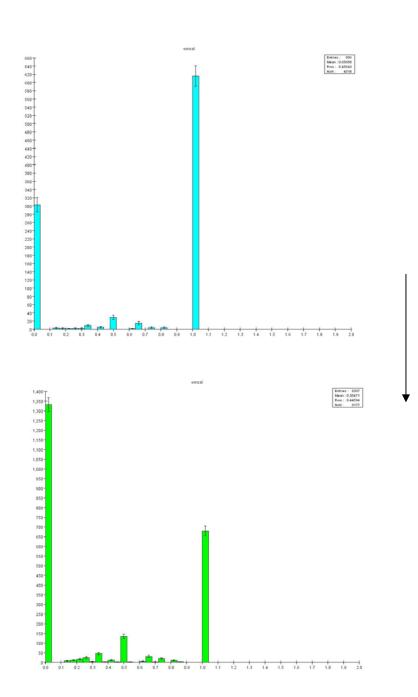


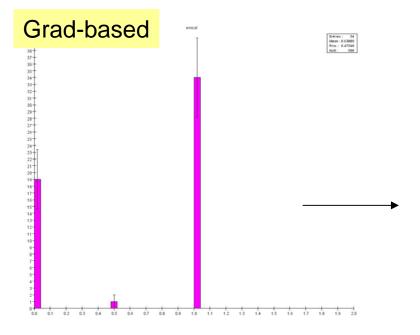


Fraction of cells attached To main cluster which belong to it Attachment is based on angular distance

Z-pole events Photons

→ Increasing dist





Fraction of cells attached To main cluster which belong to it Attachment is based on angular distance

Z-pole events Neutral hadrons

→ Increasing dist

