



# Performance Study of a New Cluster Splitting Algorithm for the Reconstruction of $\bar{P}$ ANDA EMC Data

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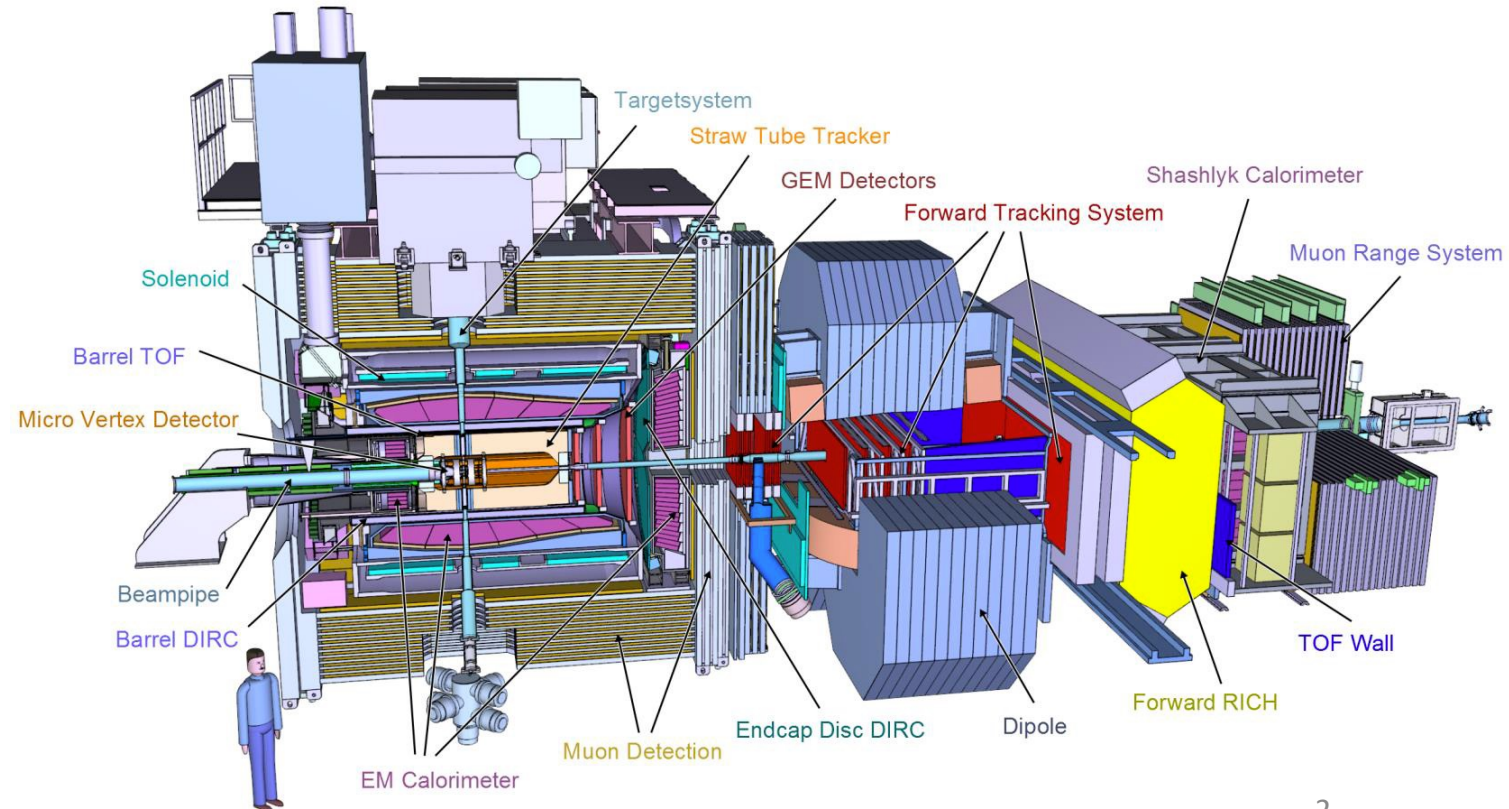
On behalf of the PANDA Collaboration



May 17, 2022

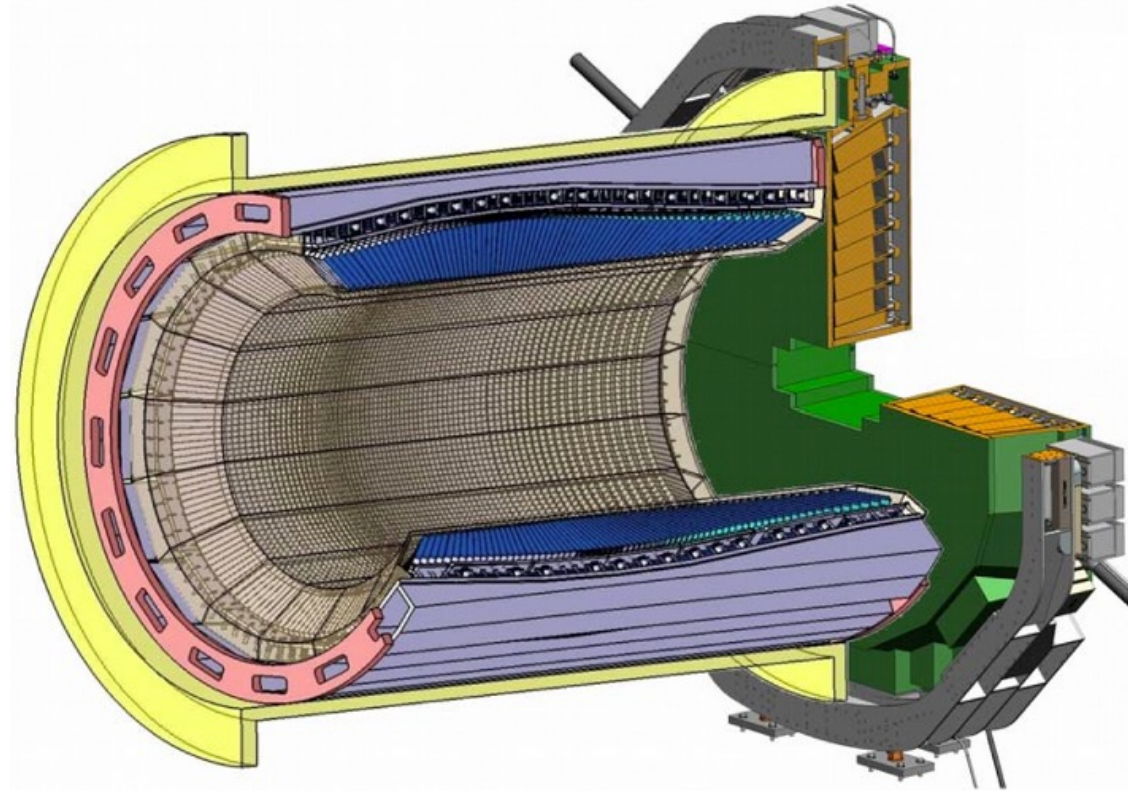
# $\bar{P}$ ANDA Experiment

- AntiProton ANnihilations at DArmstadt
- Cooled antiproton beams between 1.5 GeV/c to 15 GeV/c
- Fixed target experiment : Hydrogen and other
- $\sqrt{s}$  : 2.3 GeV ~ 5.5 GeV
- High luminosity:  $10^{32} \text{ cm}^{-2} \text{ s}^{-2}$
- Hadron spectroscopy
  - Exotic states
  - Baryons
- Nucleon structure
- Mesons in nuclear
- Hypernuclei



# $\bar{P}$ ANDA Electromagnetic Calorimeter

- $\bar{P}$ ANDA physics: Full reconstruction of multi-photon and lepton-pair channels of utmost importance
- Target Spectrometer:
  - Barrel part and two endcaps
  - 16,000 crystals, improved PbWO<sub>4</sub>
  - $X_0 = 0.89$  cm,  $R_M = 2.00$  cm
  - For barrel EMC, 11360 crystals, the average lateral size of crystal is 21.3mm
- Forward Spectrometer:
  - Shashlik type sampling calorimeter
- Good energy and spatial resolution for photons
  - $\leq 1\% \oplus \frac{\leq 2\%}{\sqrt{E/GeV}}$
  - $\leq 0.5^\circ$  (backward),  $\leq 0.3^\circ$  (barrel),  $\leq 0.1^\circ$  (forward)

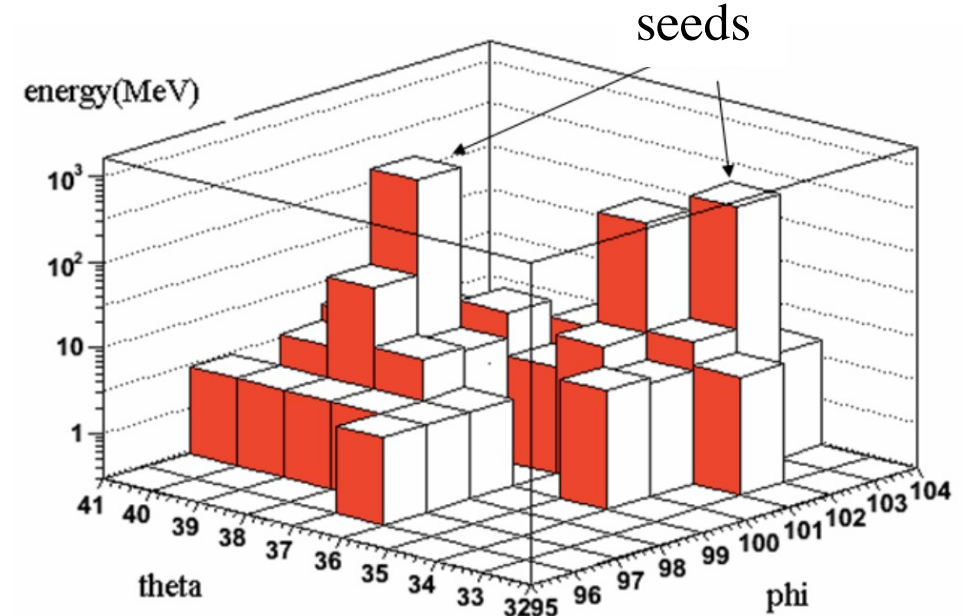


- For barrel EMC :  $22^\circ < \theta < 140^\circ$

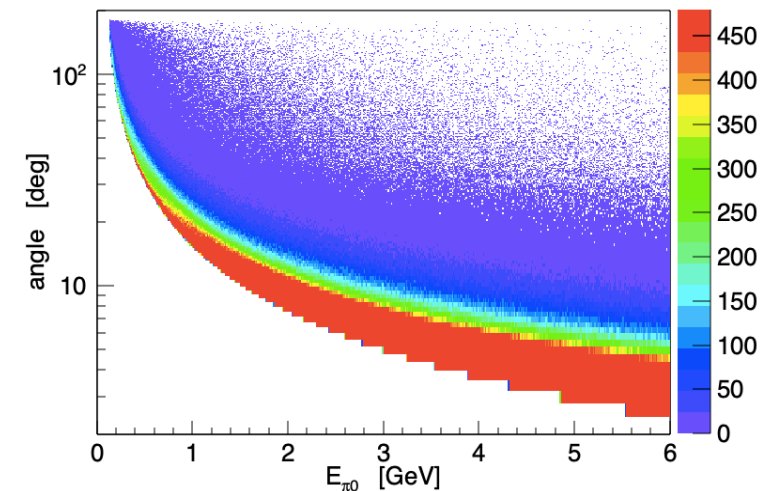


# EMC Reconstruction

- Cluster Finding
  - A contiguous area of crystals with energy deposition
  - Local maximum as the seed of shower
- Cluster Splitting
  - Several local maxima due in general to overlapping showers
  - Especially important for high momentum  $\pi^0$ , significance of  $\pi^0$  depends on mass resolution, and performance of cluster splitting algorithm



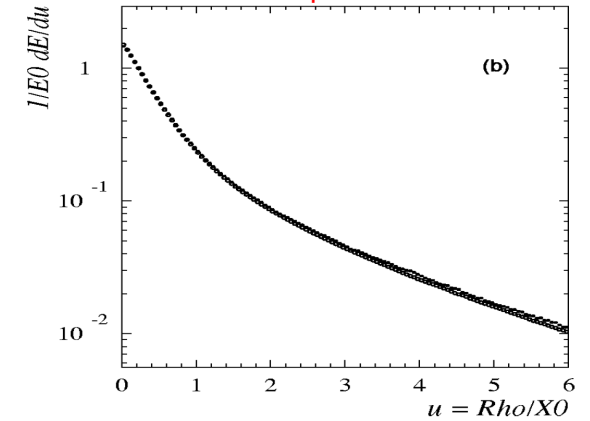
Angles between the  $2\gamma$  from  $\pi^0$



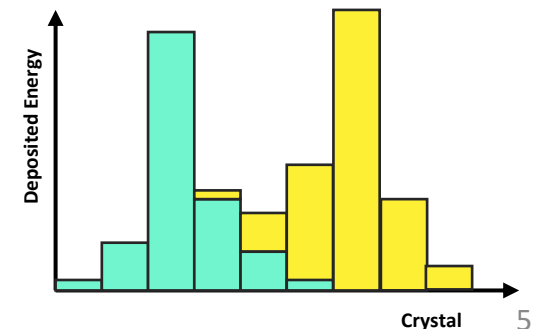
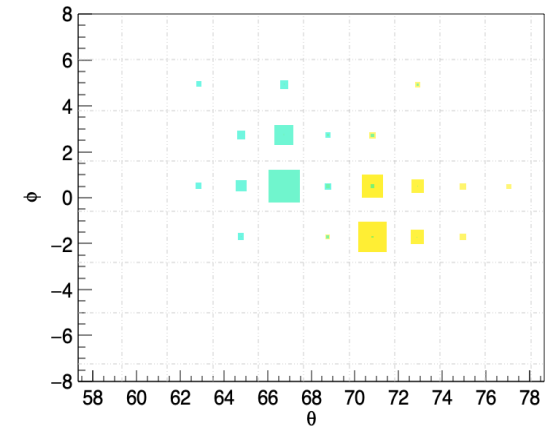
# Cluster Splitting in Reconstruction

- The purpose of the cluster splitting is to precisely assign energy deposits to the correct particles
- Update energy/position iteratively
  - EM lateral development of a shower:
$$E_{\text{target}} = E_{\text{seed}} \cdot \exp(-2.5 r/R_M)$$
  - The fraction of energy for each shower deposited in same crystal is calculated
  - Estimated energies and positions of the single showers involved as input parameters
- In this presentation, the cluster splitting algorithm is improved:
  - Update the lateral development description function based on crystal granularity
  - Correct the seed energy

The lateral development of an EM shower



Cluster splitting for an EMC Cluster



# Cluster Splitting Algorithm\*

- Initialization:

Place the shower center at the center of seed crystal, the total energy of seed crystal as  $E_{seed}$ .

- Iteration:

1. Traverse all crystals to calculate fraction  $f_i$  of shower  $i$ .

$$f_i = \frac{(E_{seed})_i \cdot \exp(-2.5 \cdot r_i / R_m)}{\sum_j (E_{seed})_j \cdot \exp(-2.5 \cdot r_j / R_m)}$$

$i$  or  $j$  : different showers

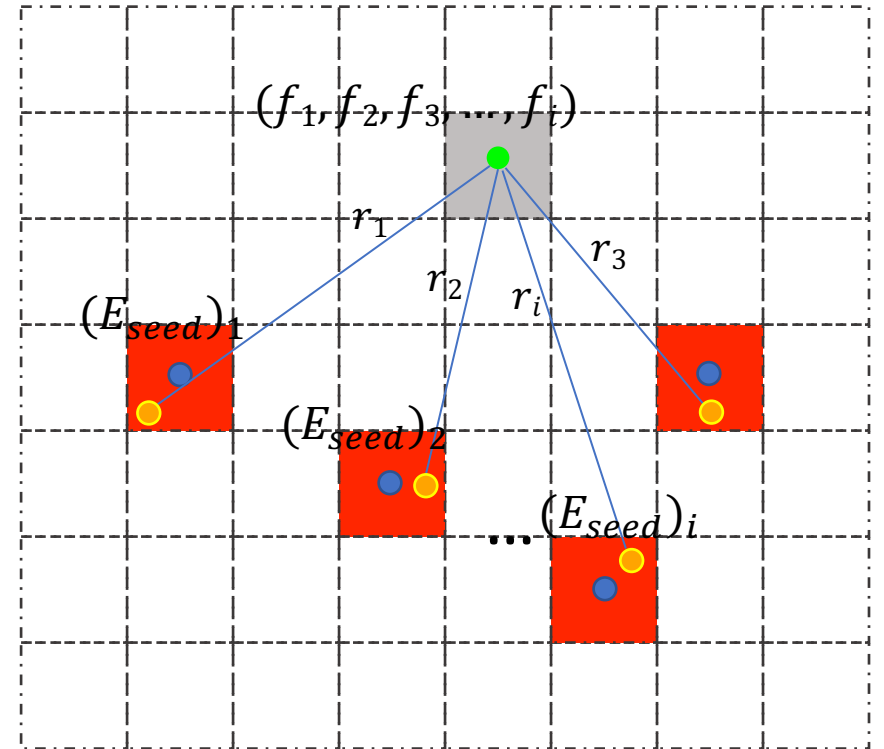
$R_m$ : Moliere radius

$r_i$ : distance from the shower center to the center of target crystal

2. Update the position of the shower center and  $E_{seed}$ .

3. Loop over 1 & 2 until the shower center convergent.

- the target crystal
- the seed crystal
- the shower center



\*Similar as BESIII and BaBar: [1] BESIII Collaboration, Nucl. Inst. Meth. A Volume 614, Issue 3, 11 March 2010. [2] BABAR Collaboration, B. Aubert *et al.*, Nucl. Instr. and Methods A **479**, 1 (2002).

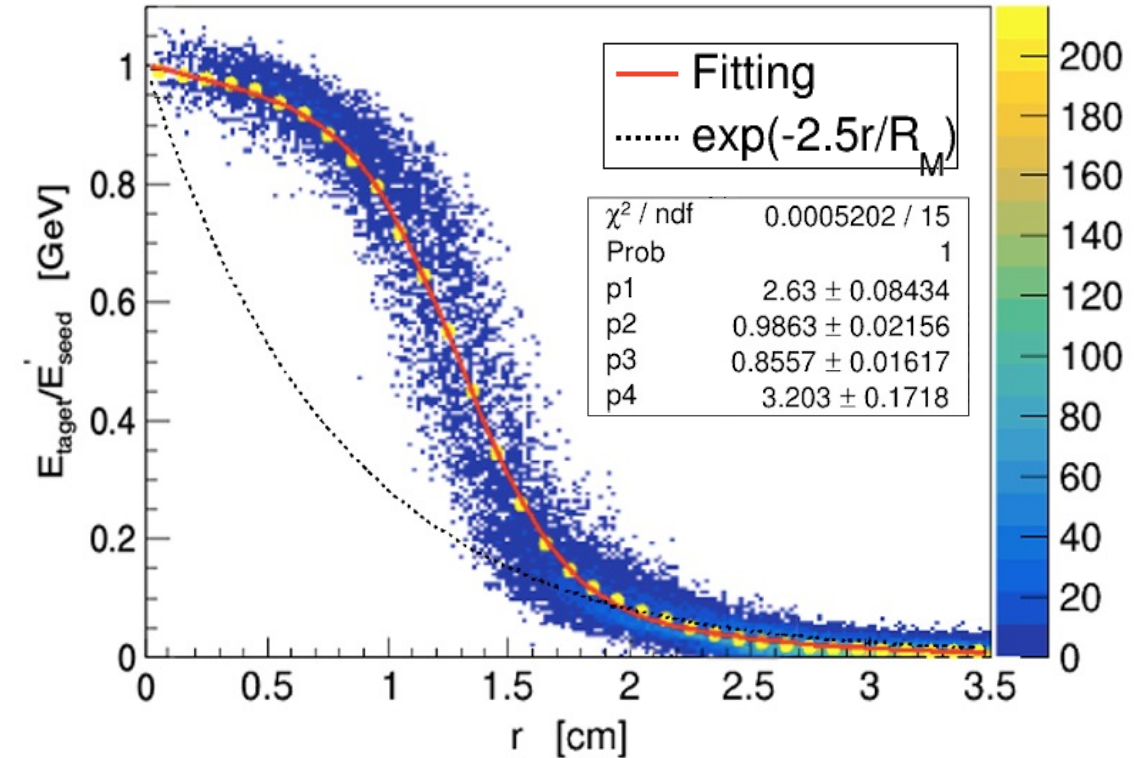
# The Lateral Development of Electromagnetic Shower

- The conventional lateral development formula  $\frac{E_{\text{target}}}{E_{\text{seed}}} = \exp(-2.5 r/R_M)$  has no consideration of **crystal granularity**.

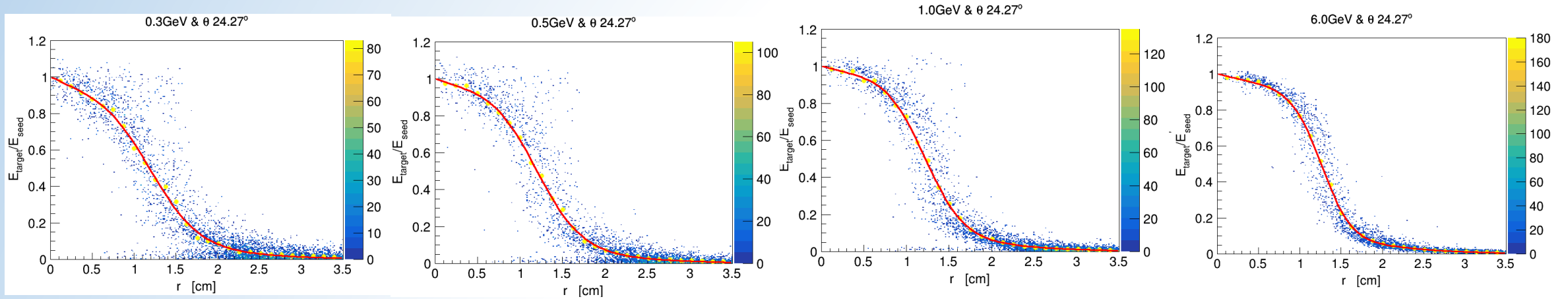
- The function form used for fitting:

$$f(r) = \frac{E_{\text{target}}}{E_{\text{seed}}} = \exp\left[-\frac{p_1}{R_M} \cdot \xi(r)\right],$$
$$\xi(r) = r - p_2 \cdot r \cdot \exp\left[-\left(\frac{r}{p_3 \cdot R_M}\right)^{p_4}\right]$$

here  $p_1$ ,  $p_2$ ,  $p_3$  and  $p_4$  are parameters.



# Parametrization



Fit the distribution of different detector polar angle and energy regions for barrel EMC to get the parameters.

$$\frac{E_{\text{target}}}{E_{\text{seed}}} = \exp\left\{-\frac{p_1}{R_M} \cdot \xi(r, p_2, p_3, p_4)\right\} \quad \xi(r) = r - p_2 \cdot r \cdot \exp\left[-\left(\frac{r}{p_3 R_M}\right)^{p_4}\right] \quad (R_M = 2.00 \text{ cm})$$

$$p_1(E_\gamma, \theta) = -0.384 * \exp(3.88 * E_\gamma) + 5.44 * 10^{-5} * (\theta - 97.7)^2 + 2.6$$

$$p_2(E_\gamma, \theta) = -0.352 * \exp(4.21 * E_\gamma) + (-3.94) * 10^{-6} * (\theta - 69)^2 + 0.932$$

$$p_3(E_\gamma, \theta) = 0.151 * \exp(4.52 * E_\gamma) + (-2.14) * 10^{-5} * (\theta - 91)^2 + 0.841$$

$$p_4(E_\gamma, \theta) = -3.51 * \exp(1.15 * E_\gamma) + 2.26 * 10^{-4} * (\theta - 80.3)^2 + 4.96$$

Energy dependency

Angle dependency

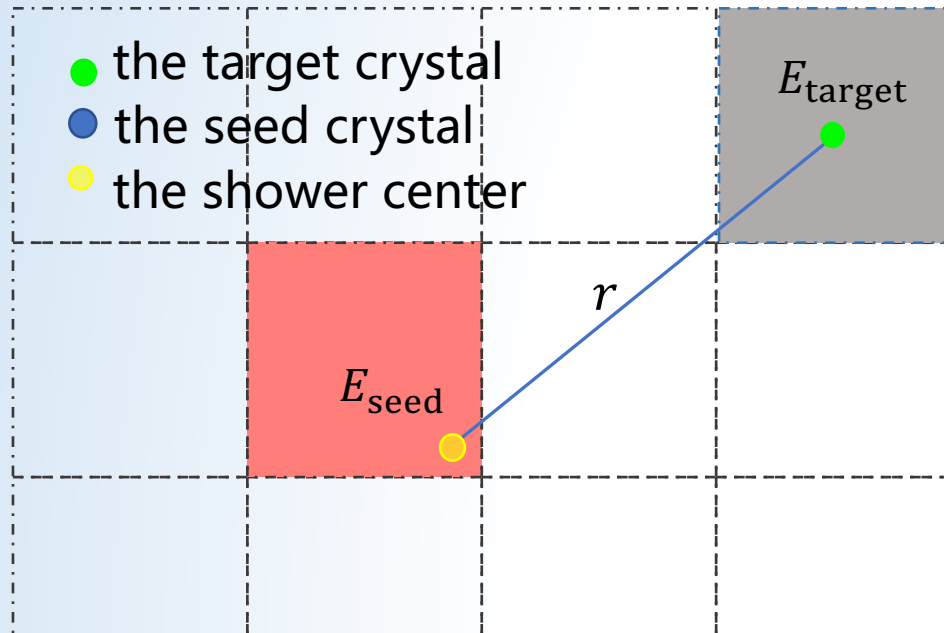


# Seed Energy Correction

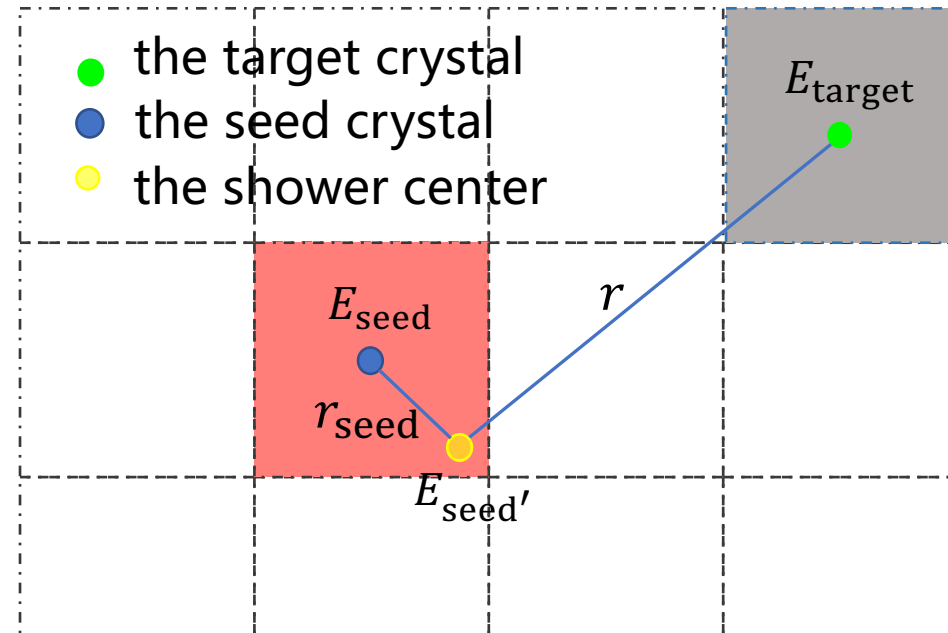
- In the default algorithm, the seed energy is used to calculate the  $E_{\text{target}} = E_{\text{seed}} \cdot f(r)$
- If the shower center not coincide with the crystal center,  $E_{\text{seed}}$  needs to be corrected

$r$  or  $r_{\text{seed}}$ :  
the distance from  
the center of the  
shower to the  
geometric center  
of the crystal.

Default algorithm



New algorithm



# Seed Energy Correction

- In the new method,  $E_{\text{target}}$  can be calculated by the lateral development  $f(r)$ :

$$E_{\text{target}} = E_{\text{seed}'} \cdot f(r)$$

- $E_{\text{seed}'}$  can be related to the  $E_{\text{seed}}$  :

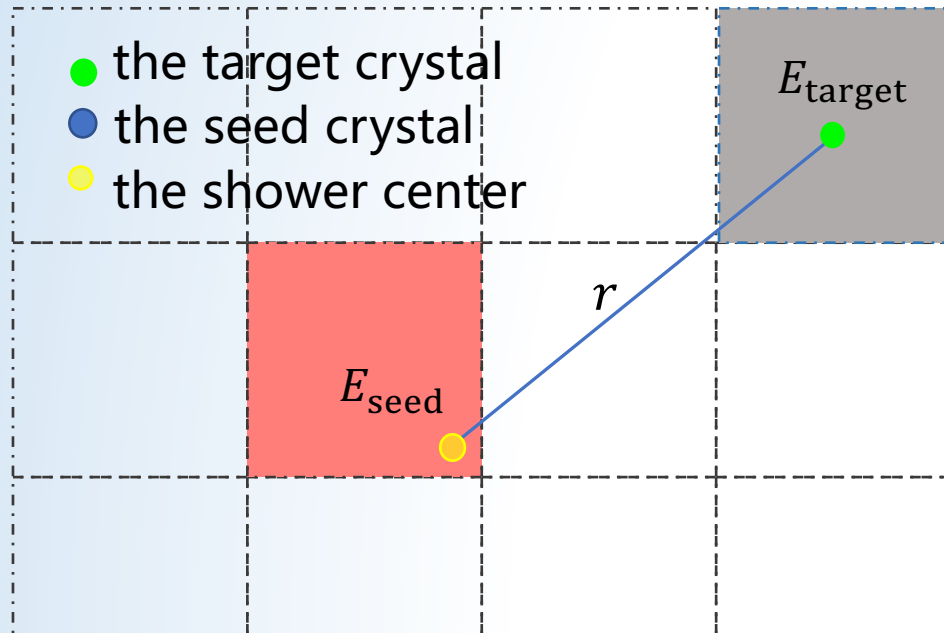
$$E_{\text{seed}'} = \frac{E_{\text{seed}}}{f(r_{\text{seed}})}$$

- $E_{\text{target}}$  can be calculated as  $\left(\frac{1}{f(r_{\text{seed}})} \text{ as the correction factor}\right)$  :

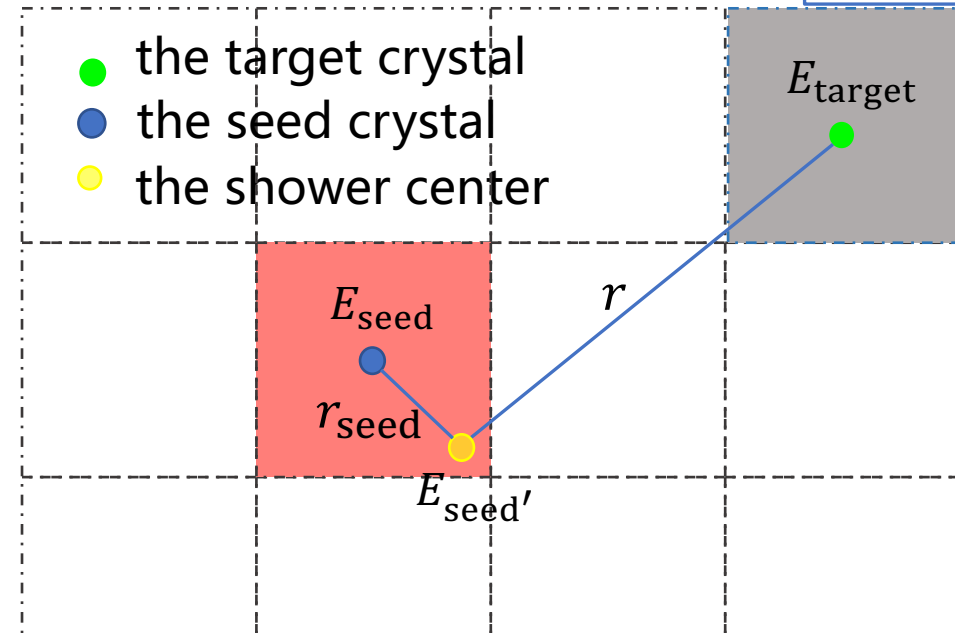
$$E_{\text{target}} = \frac{E_{\text{seed}}}{f(r_{\text{seed}})} \cdot f(r)$$

$r$  or  $r_{\text{seed}}$ :  
the distance from  
the center of the  
shower to the  
geometric center  
of the crystal.

Default algorithm



New algorithm



# Parametrization after Seed Correction

Seed energy correction

$$\frac{E_{\text{target}}}{E_{\text{seed}}} = \exp\left\{-\frac{p_1}{R_M} [\xi(r, p_2, p_3, p_4) - \xi(r_{\text{seed}}, p_2, p_3, p_4)]\right\} \xi(r) = r - p_2 \cdot r \cdot \exp\left[-\left(\frac{r}{p_3 \cdot R_M}\right)^{p_4}\right] \quad (R_M = 2.00 \text{ cm})$$

$$p_1(E_\gamma, \theta) = -0.384 * \exp(3.88 * E_\gamma) + 5.44 * 10^{-5} * (\theta - 97.7)^2 + 2.6$$

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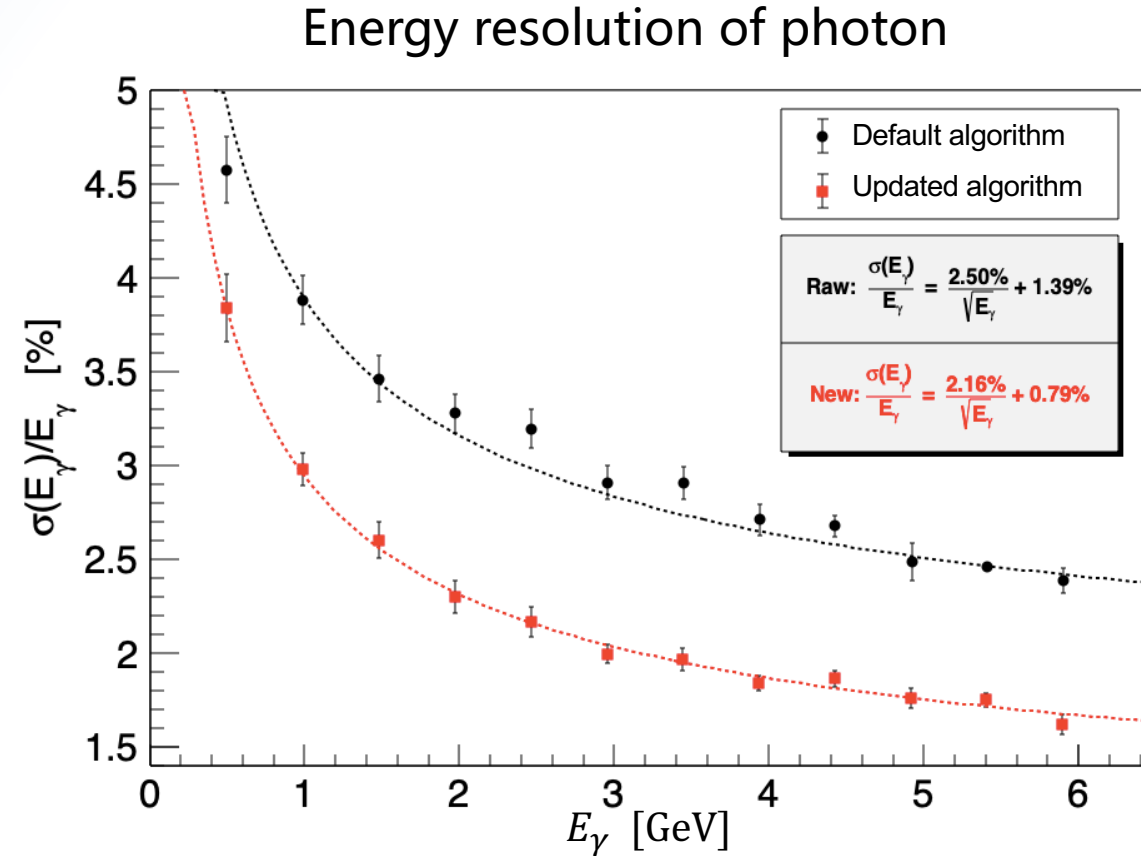
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Energy dependency

Angle dependency

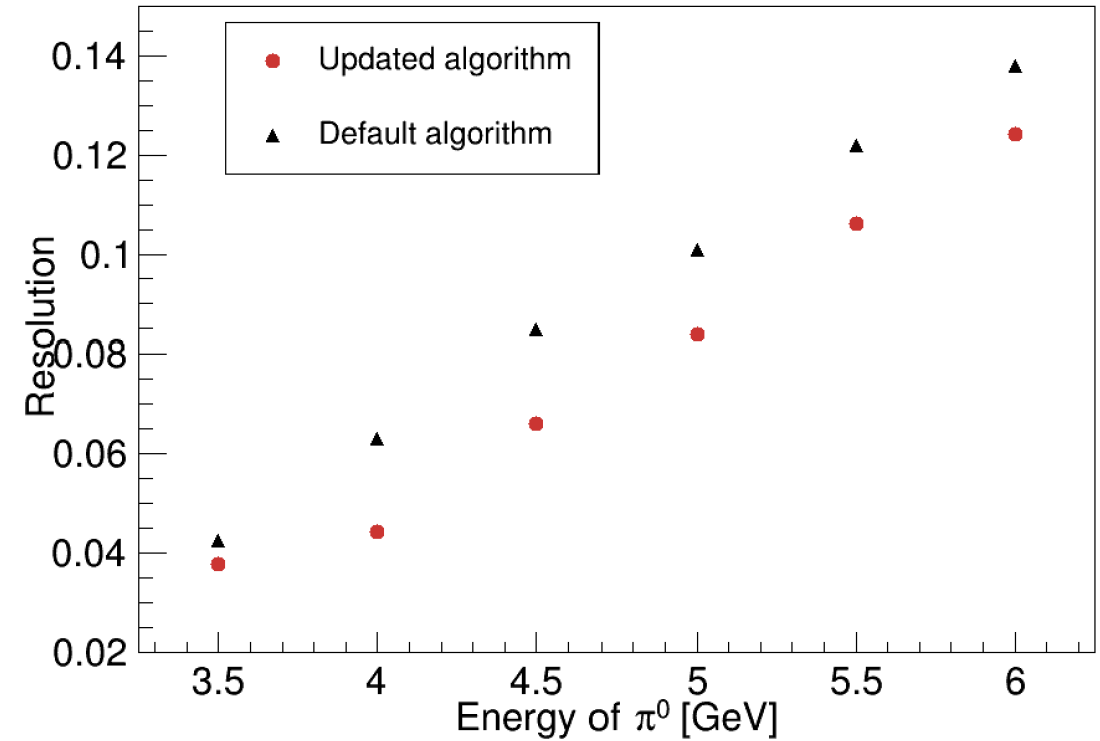
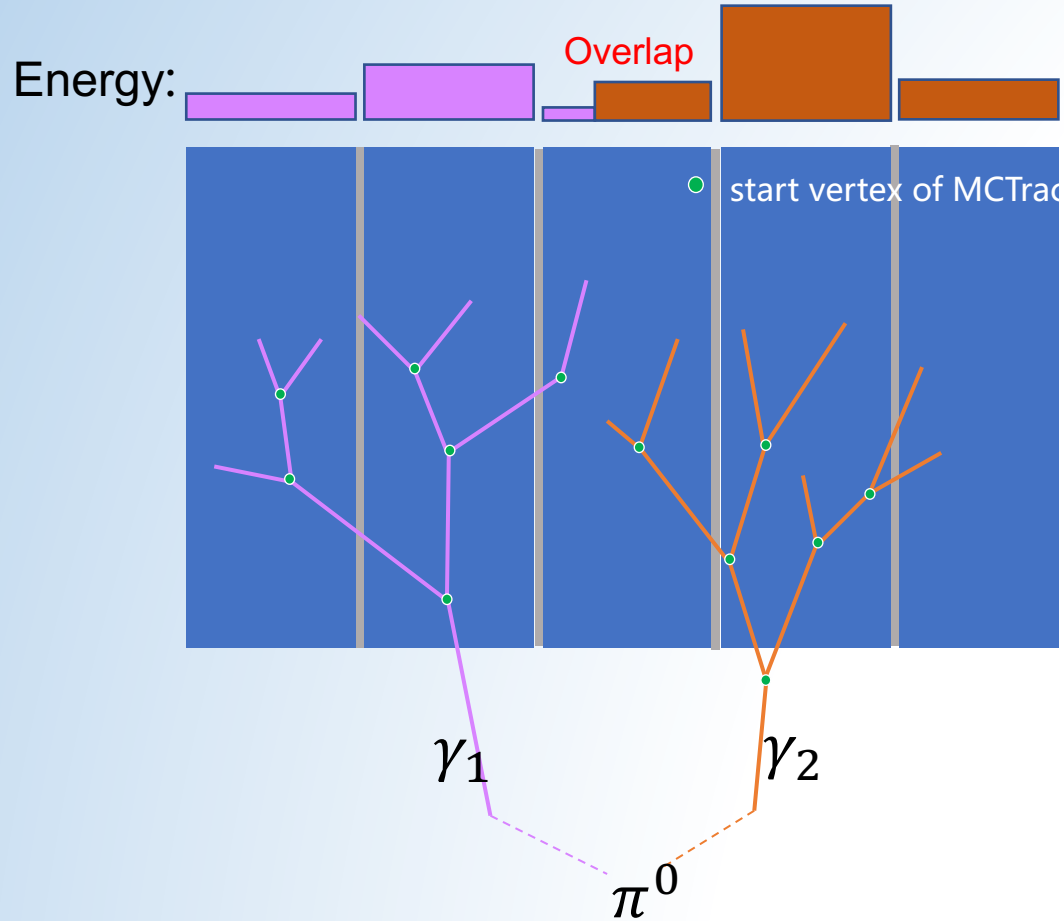
# Photon Resolution from Small-cross-angle Photon Samples

- The angle between two photons  $< 6.75(\text{deg})$
- Energy:  $0.5 \sim 6 \text{ GeV}$





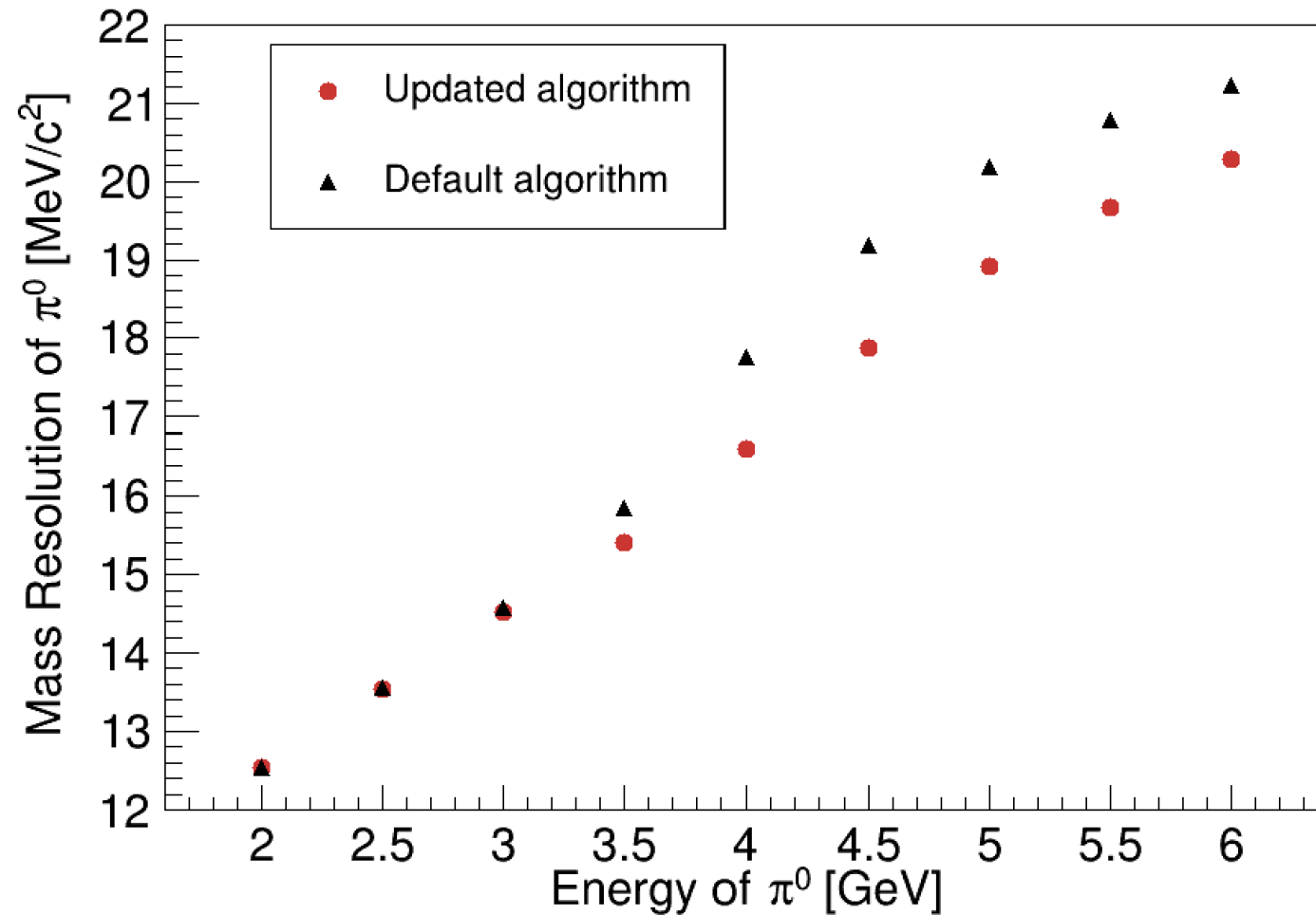
# Photon Resolution from $\pi^0$ Samples



- MC Truth information stores energy deposited in calorimeter of two photons from  $\pi^0$

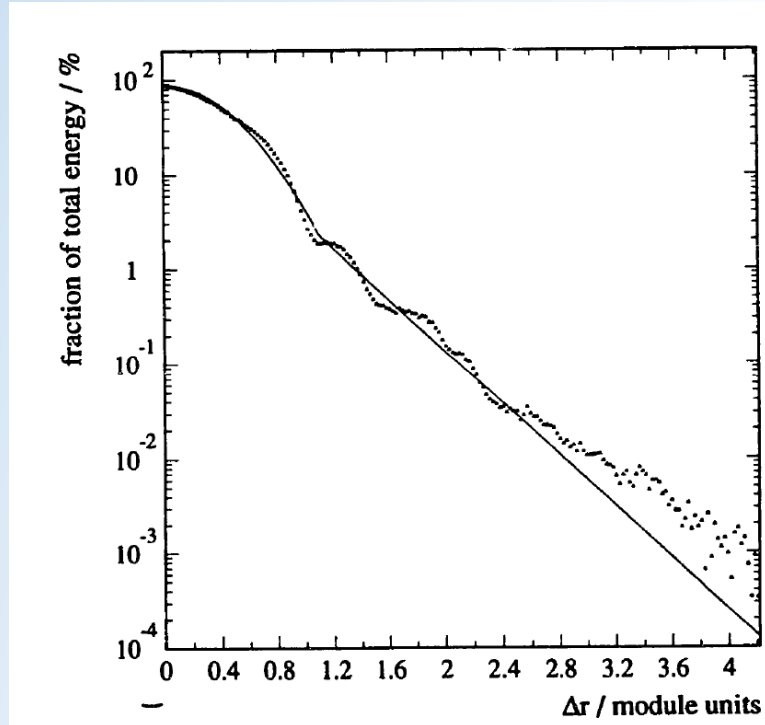
- The deviation to MC Truth is smaller for new algorithm

# $\pi^0$ Mass Resolution

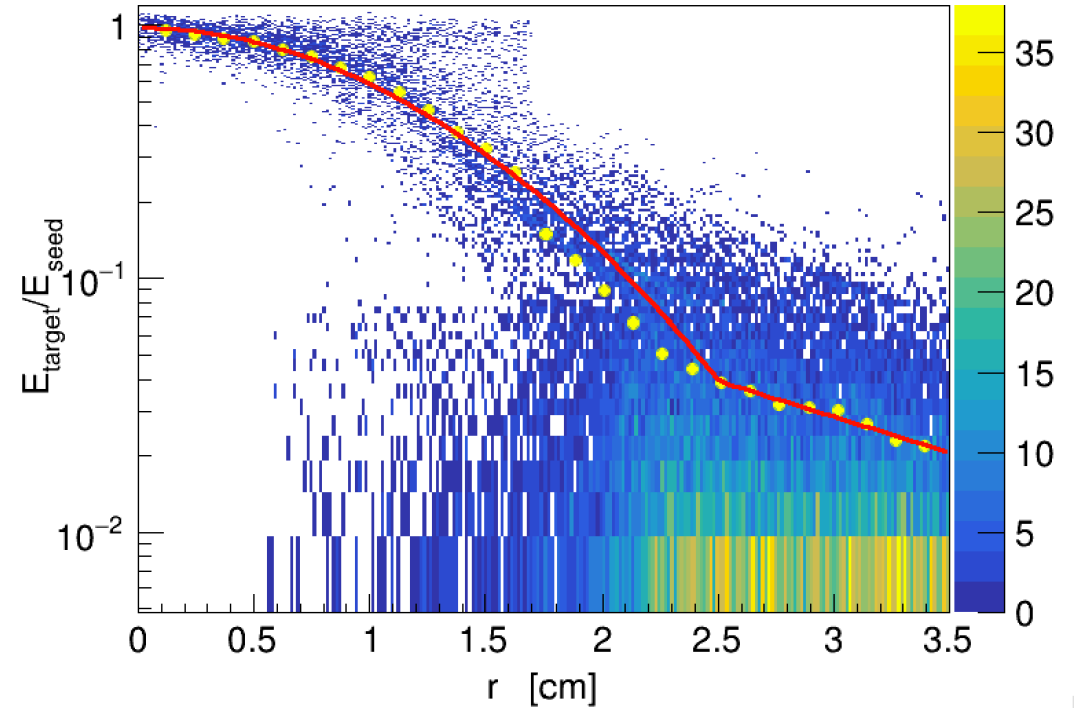


- The mass resolution of the  $\pi^0$  has been improved in the high energy range compared with the conventional method.

# Validation by experimental data



\*The lateral development of electromagnetic showers extracted from experimental data.



$$* f(r, E) = A \cdot \text{Max}\left(\exp\left(-\frac{r^2}{p_1}\right), d \cdot \exp\left(\frac{-r}{s}\right)\right)$$

$$d = p_2 + p_3 \cdot E^{p_4}$$

$$s = p_5 - p_6 \cdot \ln(E)$$

- [1] F. Berger et al., Nucl. Instrum. Methods A321 (1992) 152.  
 [2] ALICE Technology design report of the photon spectrometer.

# Summary

- The cluster splitting algorithm is improved with a new lateral development description function
  - Lateral development based on the crystal granularity
  - Dependency of energy and angle of incident particle
  - Seed energy is corrected
- Good performance of new splitting algorithm
  - Energy resolution of small-cross-angle photons and daughter photons of  $\pi^0$
  - Mass resolution of  $\pi^0$
- Obvious improvements of mass resolution and significance of high momentum  $\pi^0$

Thank you!



# Backup Slides