

# SoLID Tracking Reconstruction

Weizhi Xiong

Duke University

SoLID Collaboration Meeting

January 12-13, 2016

# Previously on Tracking Resolution

- Using Kalman Filter algorithm to reconstruct vertex variables
- Using non-digitized signal track, **no process noise** simulated (Coulomb multiple scattering and energy loss)

# Previously on Tracking Resolution

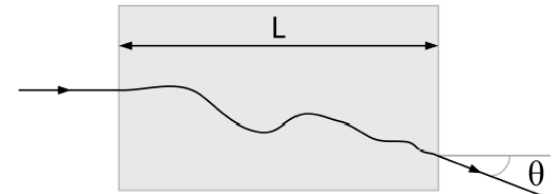
- Using Kalman Filter algorithm to reconstruct vertex variables
- Using non-digitized signal track, **no process noise** simulated (Coulomb multiple scattering and energy loss)
- GEM resolution - 90um, beam spot resolution - 300um.

	$\theta$ (mrad)	$\varphi$ (mrad)	$z$ (cm)	$p$ (%)
SIDIS FA	0.30	0.77	0.54	0.44
SIDIS LA	0.67	0.57	0.47	0.57
PVDIS	0.37	0.77	0.24	0.60

# Process Noise

- Coulomb multiple scattering
  - RMS of the scattering angle can be described by:

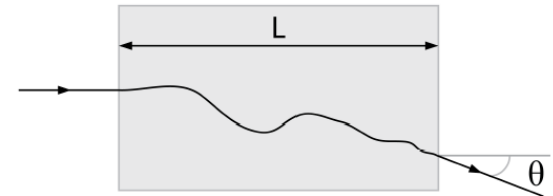
$$\Theta_{CMS} = \frac{13.6 \text{ MeV}}{\beta pc} z \sqrt{t} (1 + 0.038 \ln t) \quad \text{where} \quad t = \frac{l}{X_0}$$



# Process Noise

- Coulomb multiple scattering
  - RMS of the scattering angle can be described by:

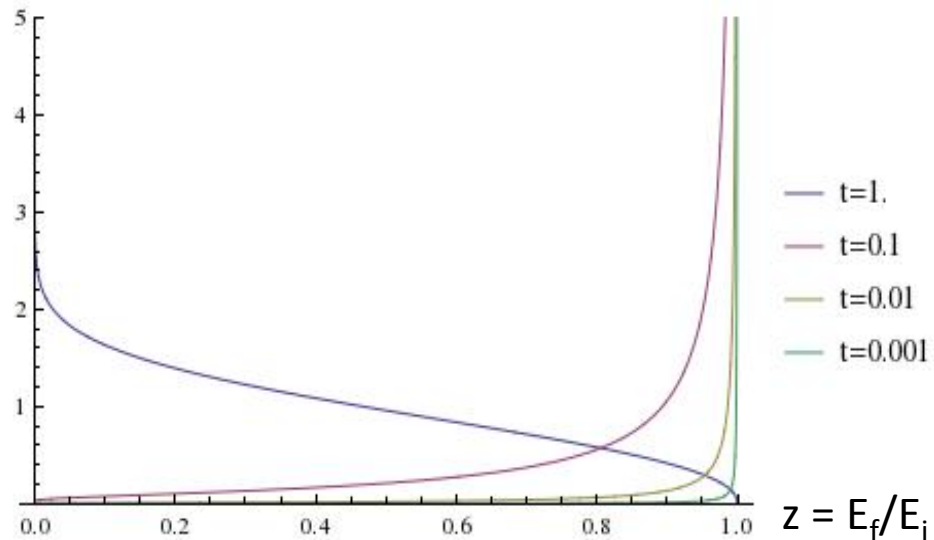
$$\Theta_{CMS} = \frac{13.6 \text{ MeV}}{\beta pc} z \sqrt{t} (1 + 0.038 \ln t) \quad \text{where} \quad t = \frac{l}{X_0}$$



- Bremsstrahlung radiation for electrons
  - Mean value and probability density function given by the Bethe-Heitler model:

$$\left\langle \frac{E_f}{E_i} \right\rangle = e^{-t}$$

$$f\left(z = \frac{E_f}{E_i}\right) = \frac{[-\ln z]^{t/\ln 2 - 1}}{\Gamma(t / \ln 2)}$$

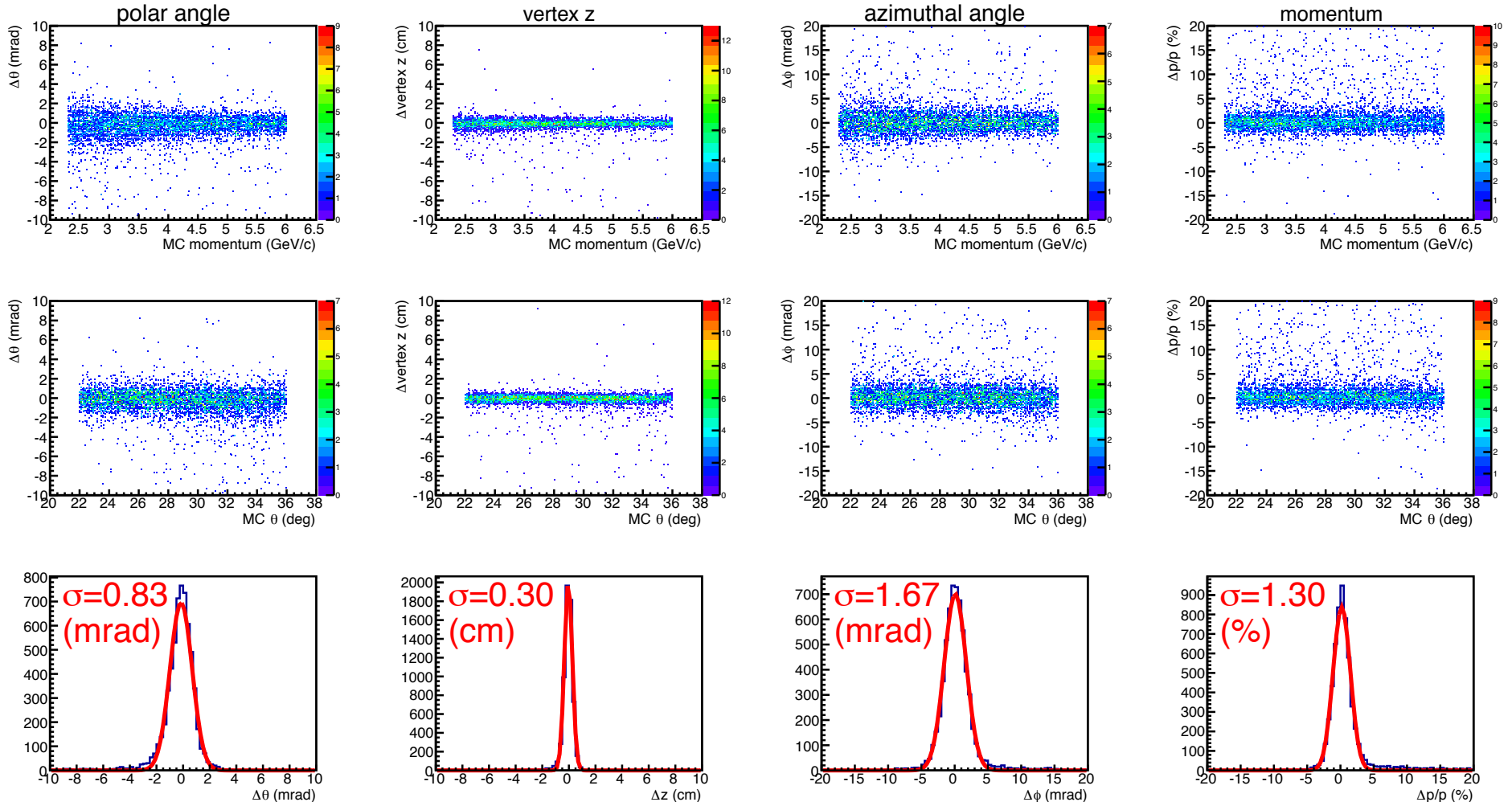


# Material Properties

For everyone		$L/X_0$ (%)
	GEM	0.52
	Air	~1%
SIDIS		$L/X_0$ (%)
	SIDIS Cell	1.43
J/ $\psi$		$L/X_0$ (%)
	J/ $\psi$ Cell	0.20
	LH2	0.21
PVDIS		$L/X_0$ (%)
	PVDIS Cell	0.42
	LD2	0.33

# PVDIS Vertex Resolution

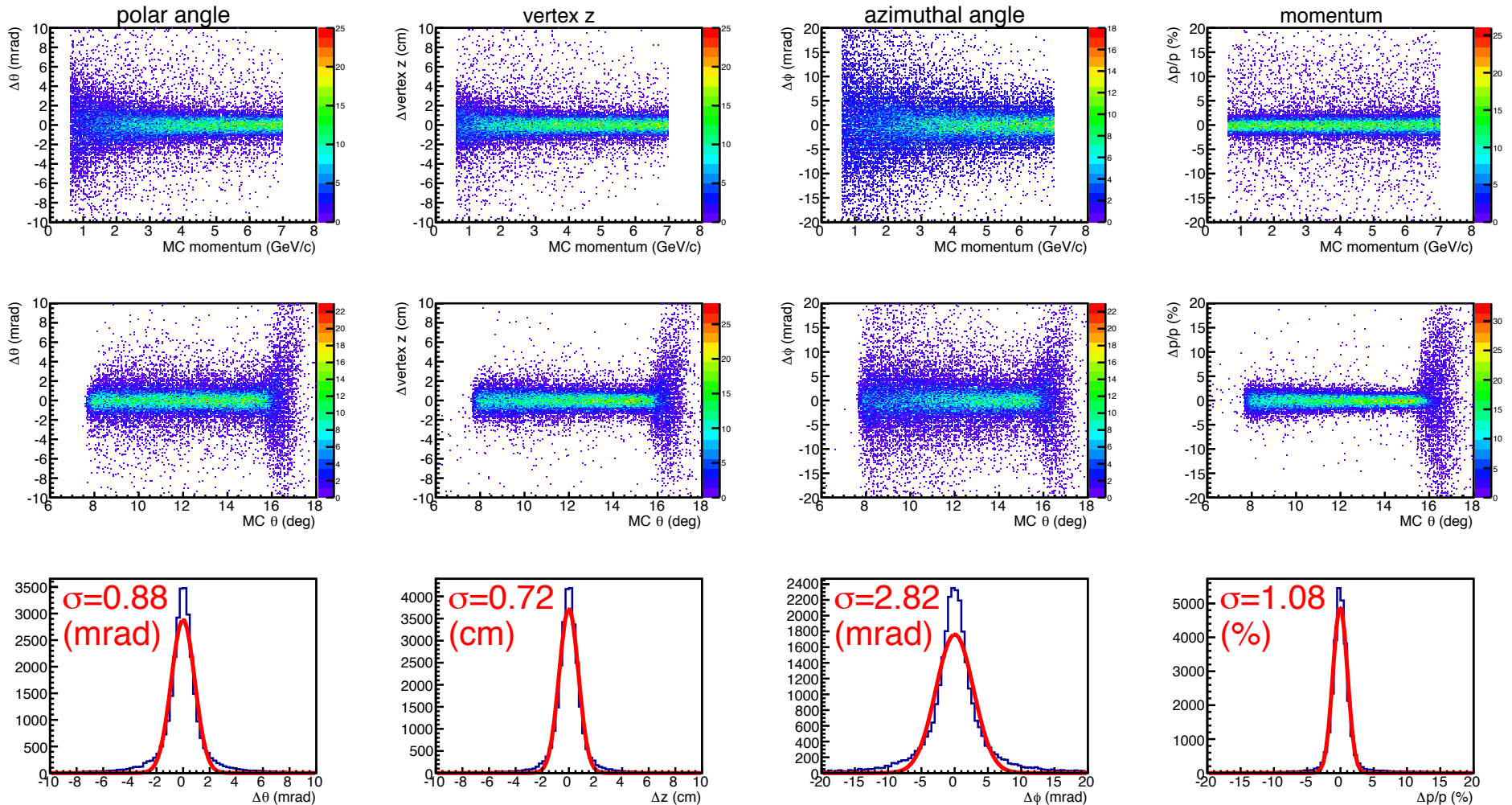
PVDIS electron. GEM resolution 90 $\mu$ m, beam spot resolution 300 $\mu$ m  
Requirement: 1mrad for polar angle and 2% for momentum



# J/ψ FA Vertex Resolution

Pion in J/ψ FA. GEM resolution 90μm, beam spot resolution 300μm

Requirement: 0.6mrad for polar angle, 5mrad for azimuthal angle and 2% for momentum

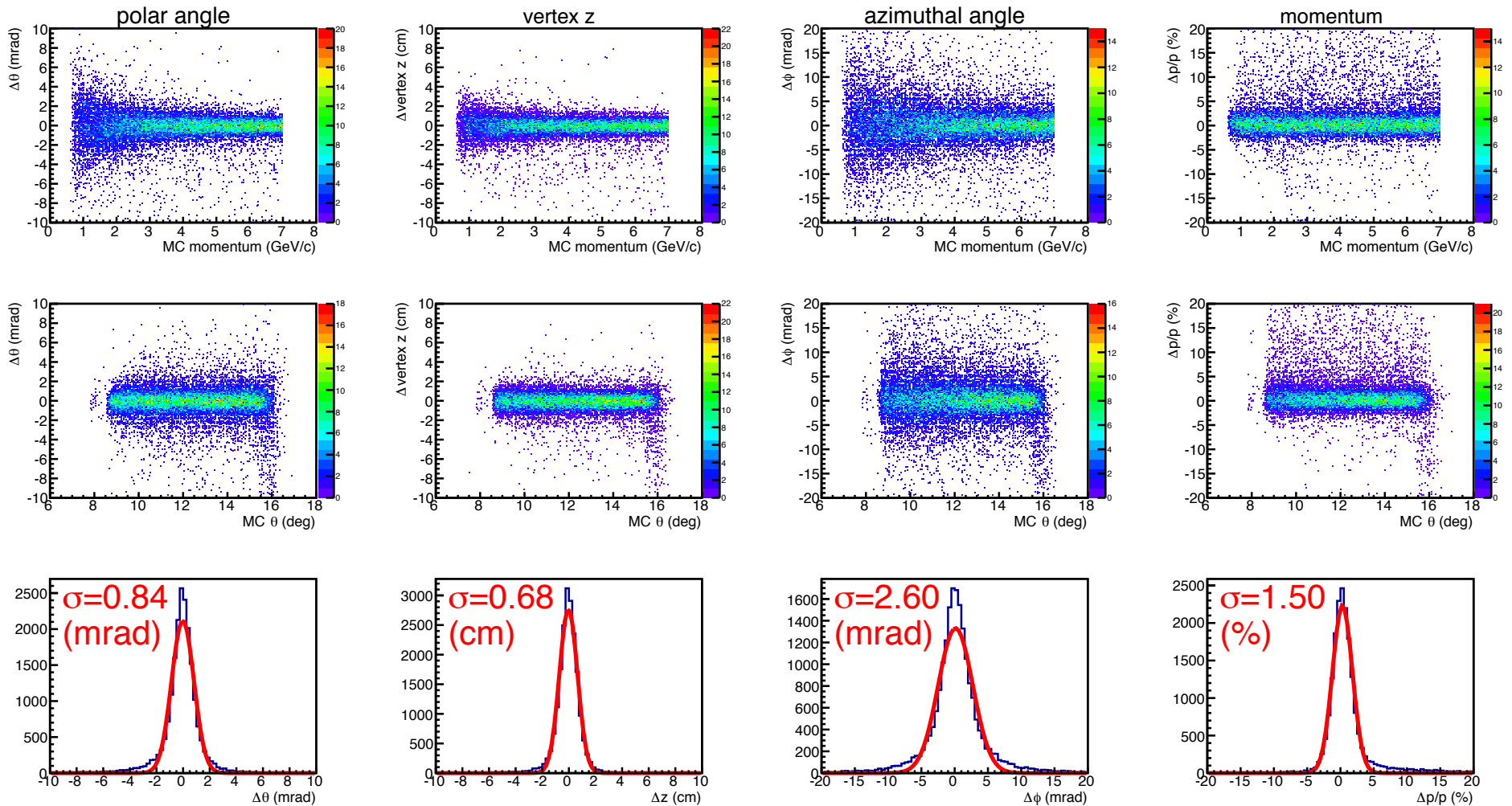




# J/ $\psi$ FA Vertex Resolution

Electron in J/ $\psi$  FA. GEM resolution 90 $\mu$ m, beam spot resolution 300 $\mu$ m

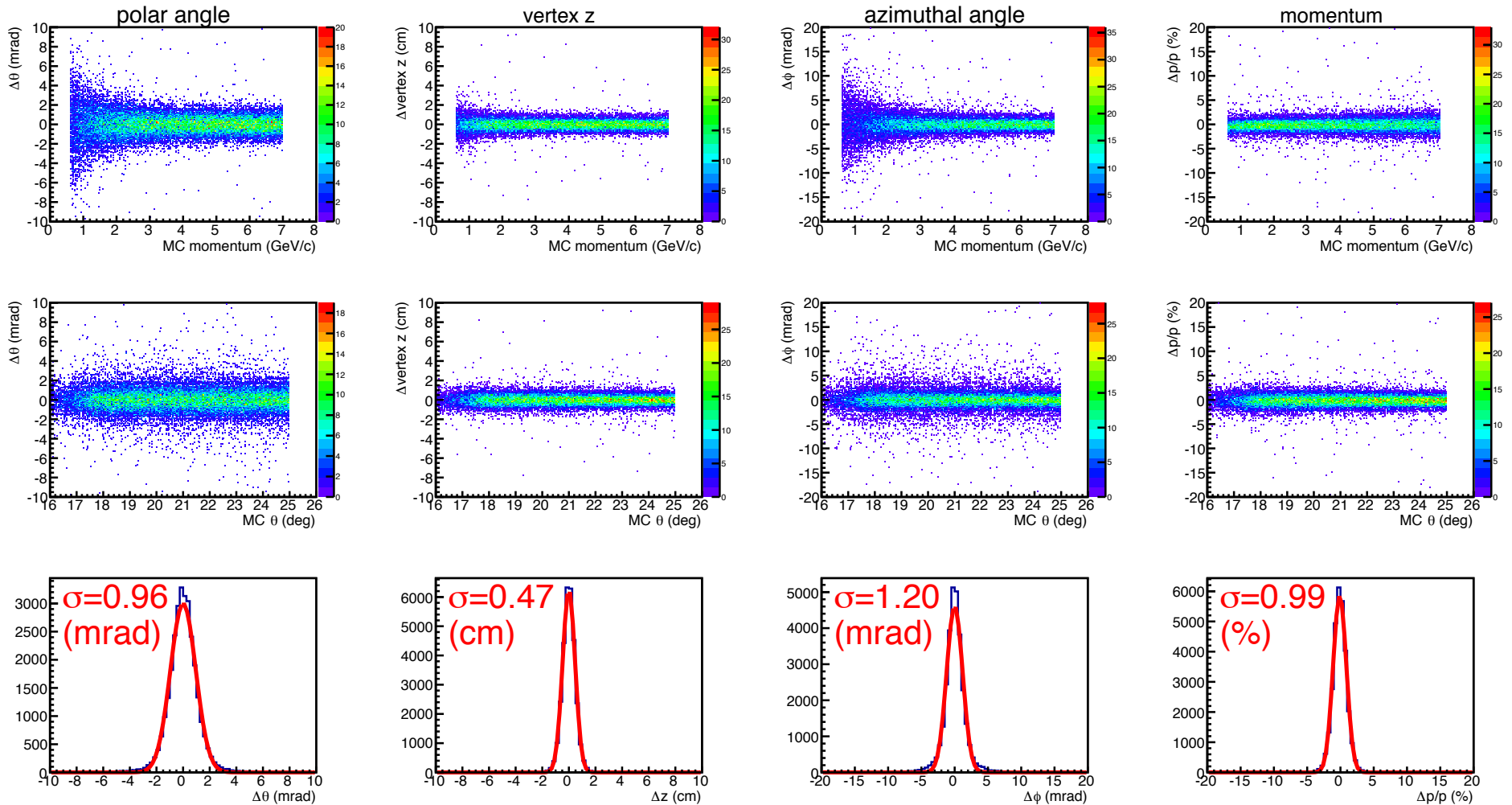
Requirement: 0.6mrad for polar angle, 5mrad for azimuthal angle and 2% for momentum



# J/ψ LA Vertex Resolution

Pion in J/ψ LA. GEM resolution 90μm, beam spot resolution 300μm

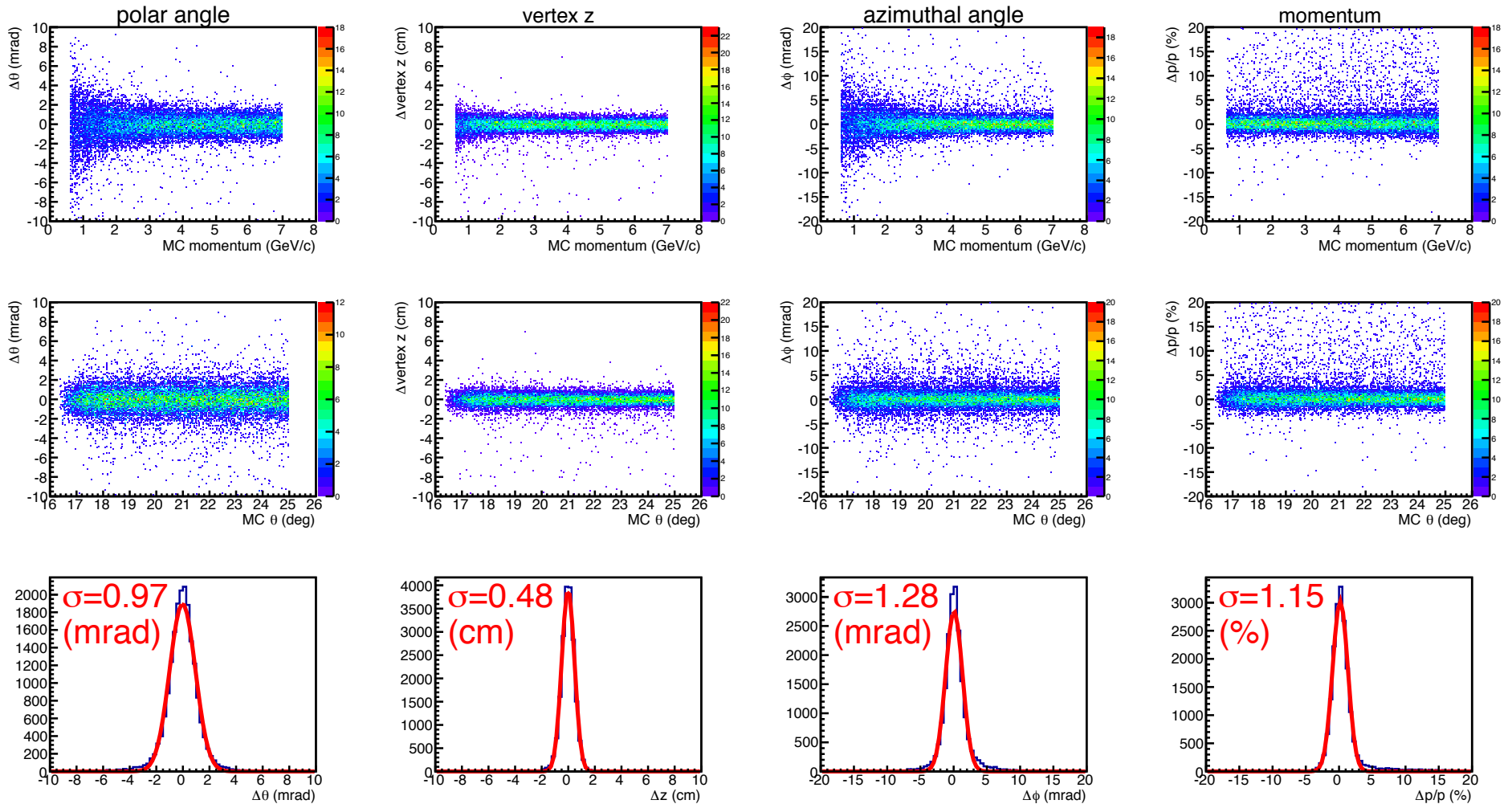
Requirement: 0.6mrad for polar angle, 5mrad for azimuthal angle and 2% for momentum



# J/ψ LA Vertex Resolution

Electron in J/ψ LA. GEM resolution 90μm, beam spot resolution 300μm

Requirement: 0.6mrad for polar angle, 5mrad for azimuthal angle and 2% for momentum

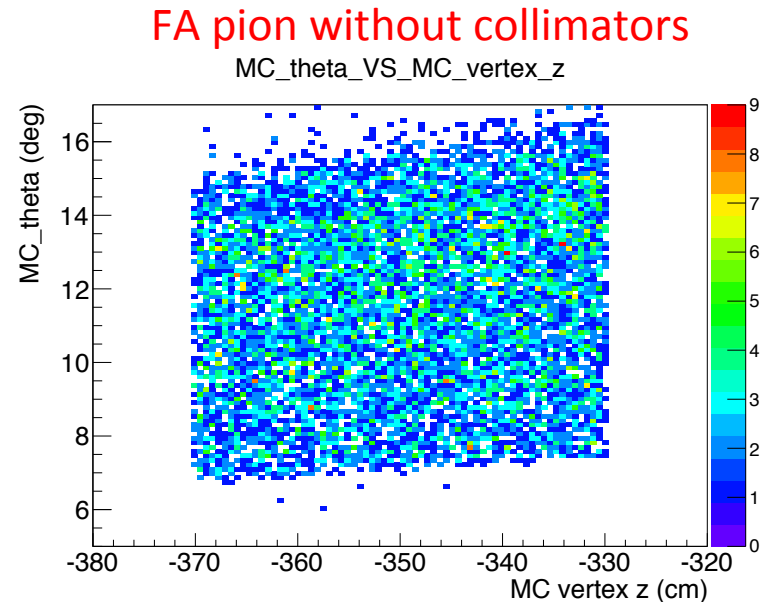


# SIDIS FA Vertex Resolution

- Two collimators used for SIDIS, made of tungsten ( $X_0 = 0.3504\text{cm}$ )
- Effective in blocking electrons, but not as much for hadron

# SIDIS FA Vertex Resolution

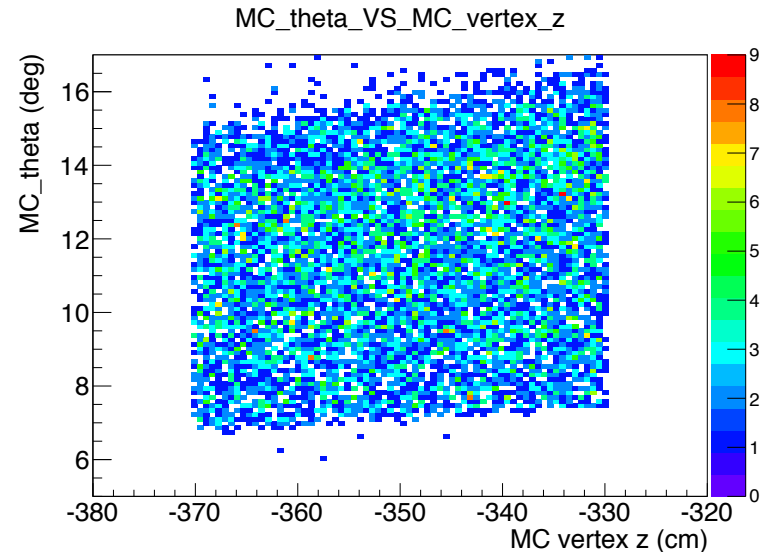
- Two collimators used for SIDIS, made of tungsten ( $X_0 = 0.3504\text{cm}$ )
- Effective in blocking electrons, but not as much for hadron



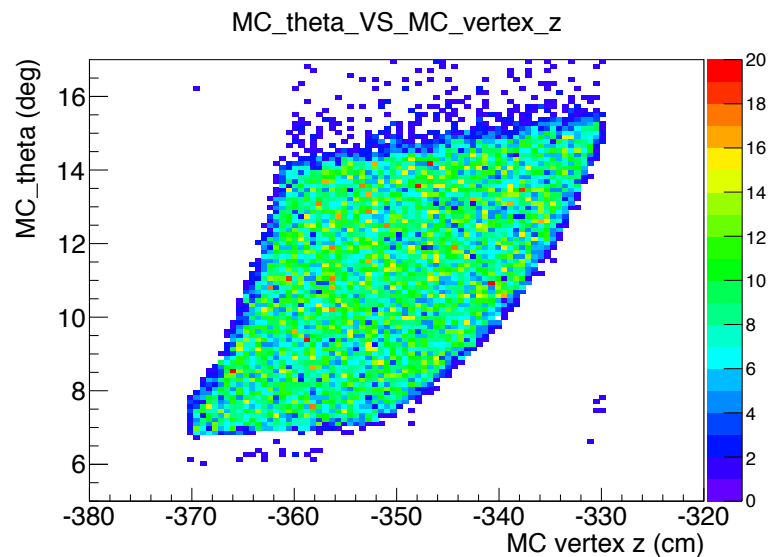
# SIDIS FA Vertex Resolution

- Two collimators used for SIDIS, made of tungsten ( $X_0 = 0.3504\text{cm}$ )
- Effective in blocking electrons, but not as much for hadron

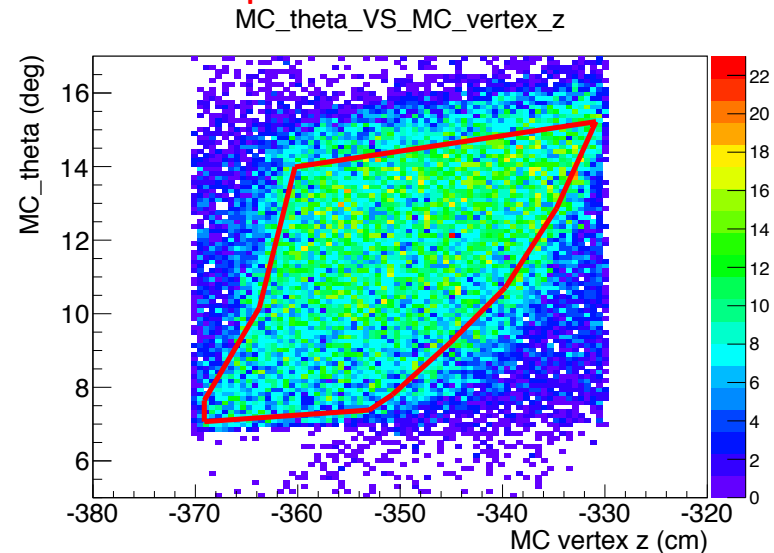
FA pion without collimators



FA electron with collimators



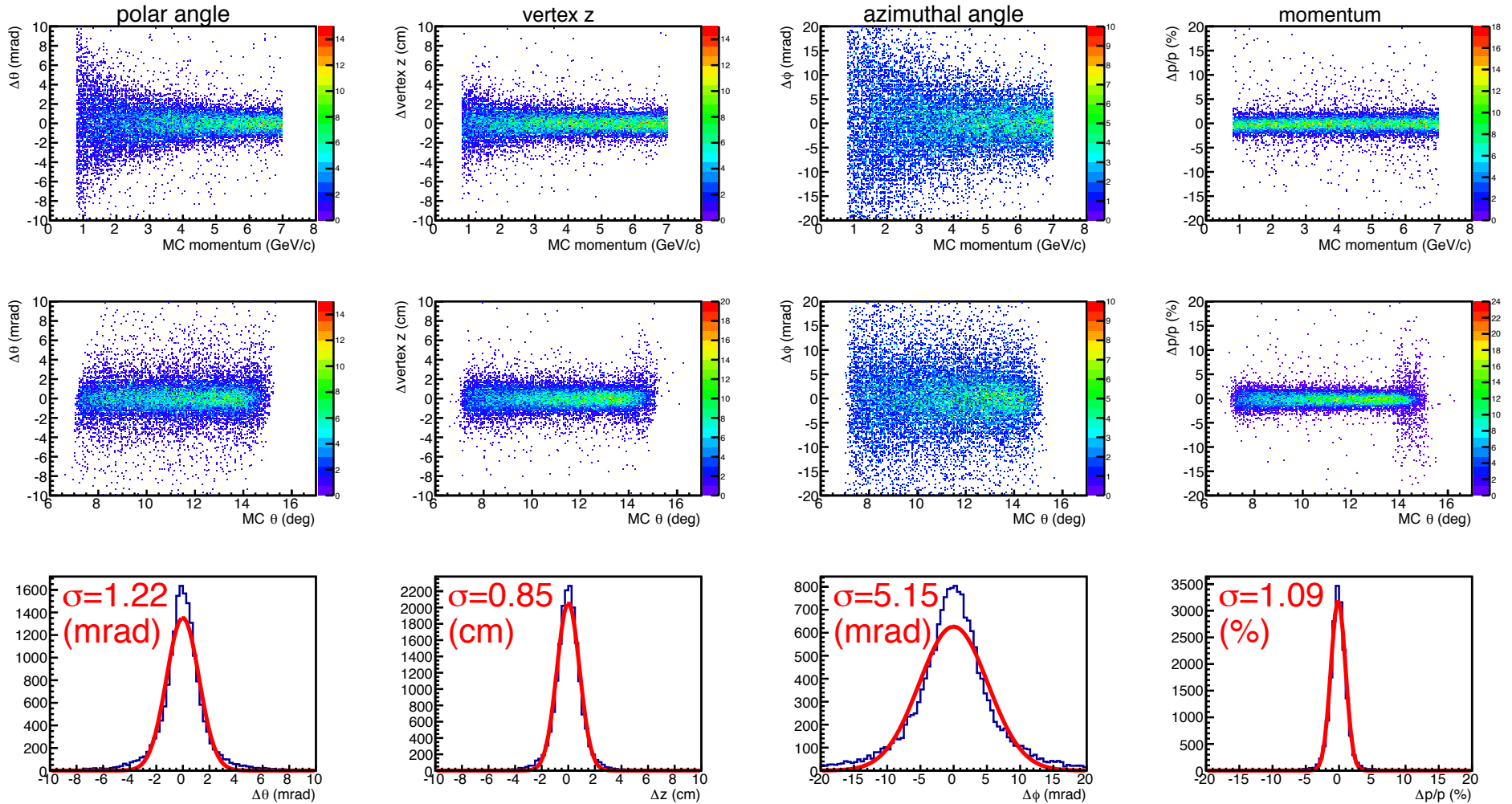
FA pion with collimators



# SIDIS FA Vertex Resolution

Pion in SIDIS FA. GEM resolution 90 $\mu\text{m}$ , beam spot resolution 300 $\mu\text{m}$

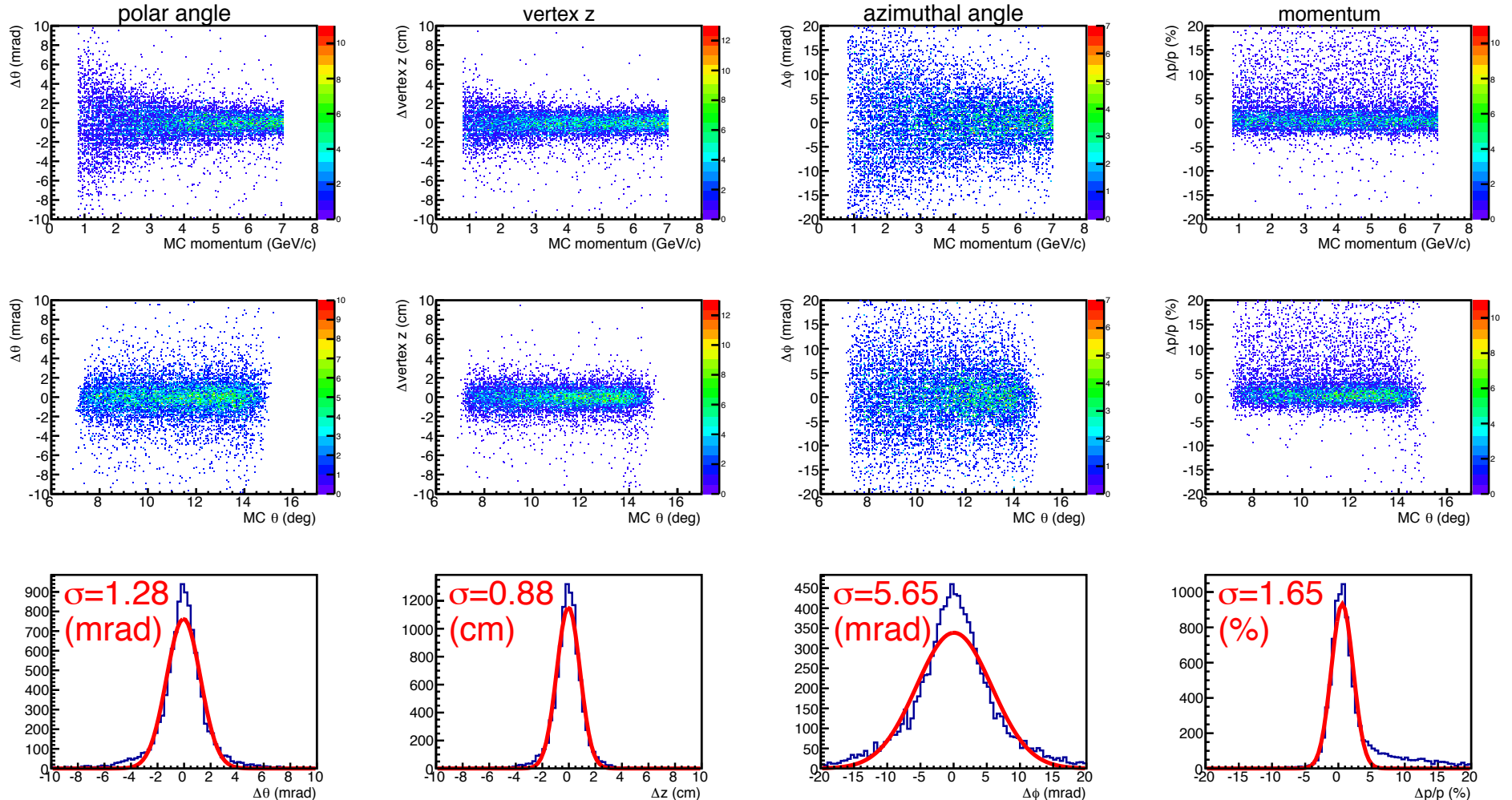
Requirement: 0.6mrad for polar angle, 5mrad for azimuthal angle and 2% for momentum



# SIDIS FA Vertex Resolution

Electron in SIDIS FA. GEM resolution 90 $\mu$ m, beam spot resolution 300 $\mu$ m

Requirement: 0.6mrad for polar angle, 5mrad for azimuthal angle and 2% for momentum

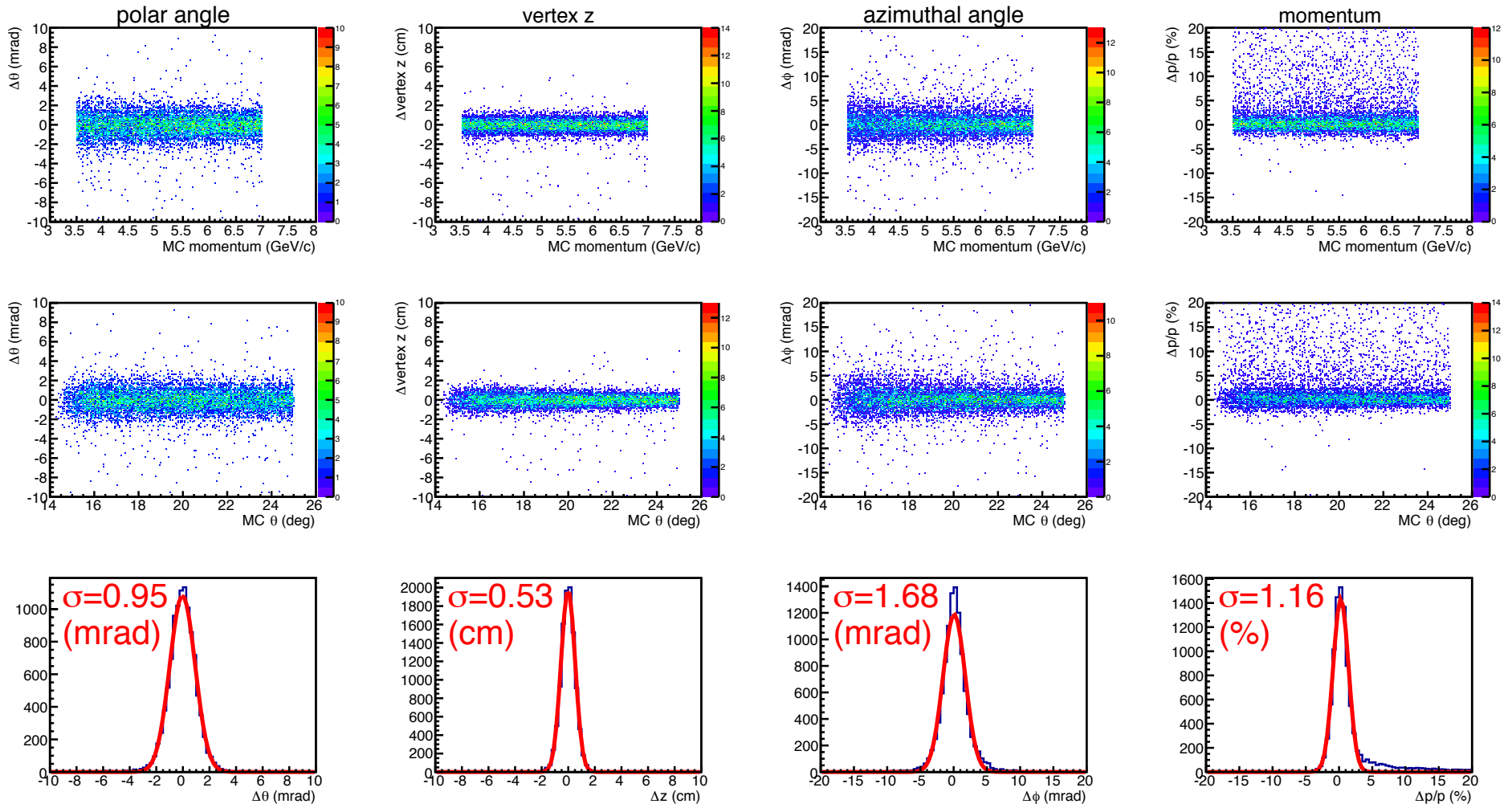




# SIDIS LA Vertex Resolution

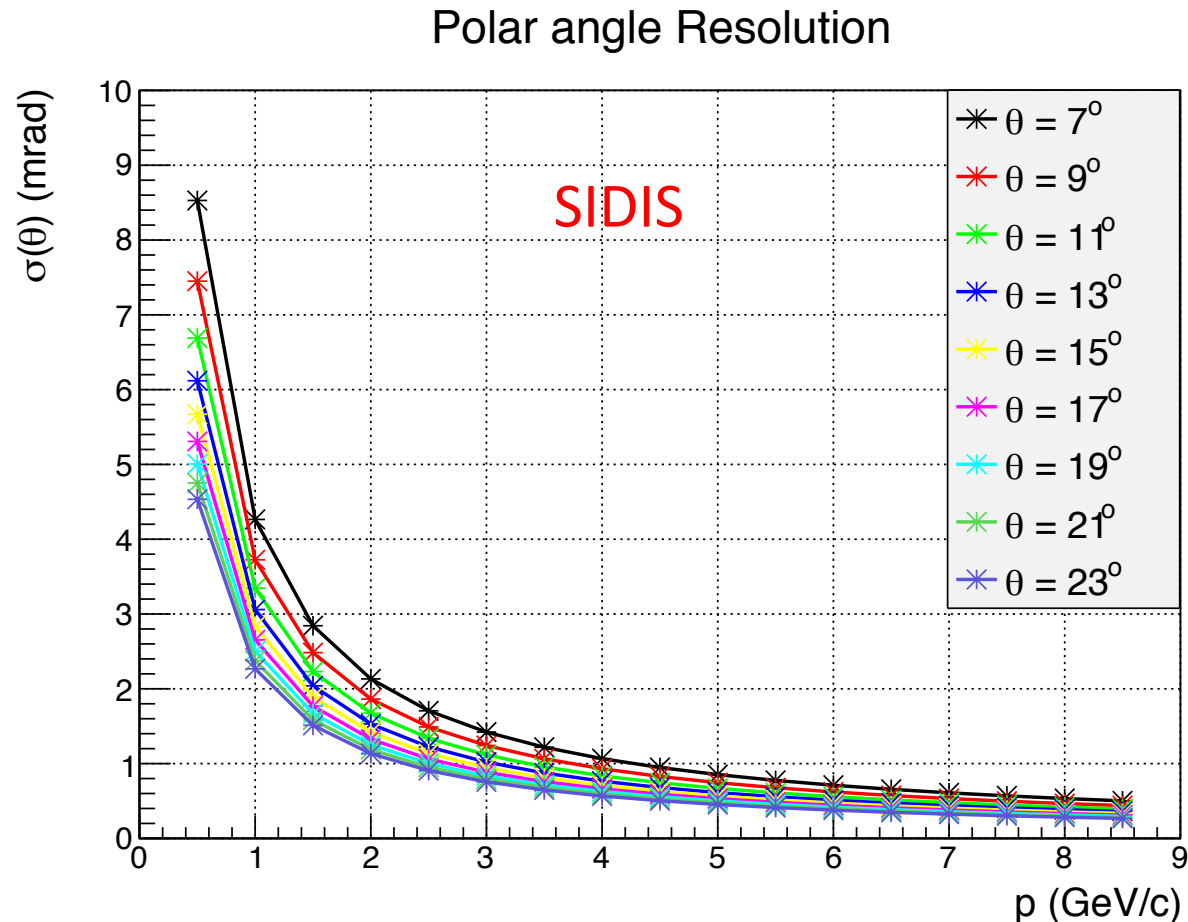
Electron in SIDIS LA. GEM resolution 90 $\mu$ m, beam spot resolution 300 $\mu$ m

Requirement: 0.6mrad for polar angle, 5mrad for azimuthal angle and 2% for momentum



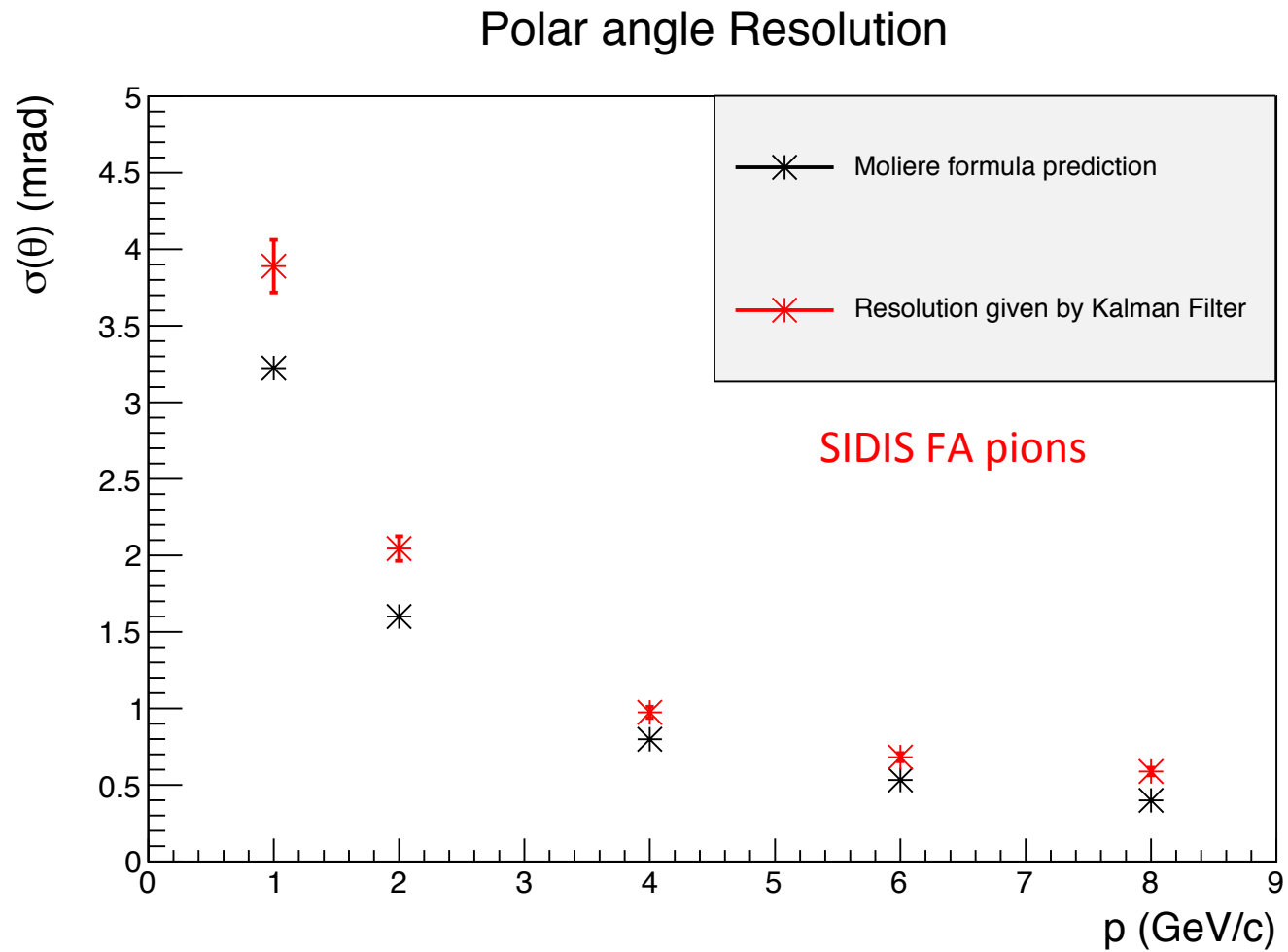
# Polar Angle Resolution Limit

- Best guess for polar angle (for now): the polar angle inside the cell is the same as the polar angle outside
- In this case, polar angle resolution cannot be better than results given by the Moliere formula for particles pass through target cell



# Molière VS Kalman Filter

MC theta fixed at 12 deg. MC momentum fixed at 1, 2, 4, 6, and 8 GeV.



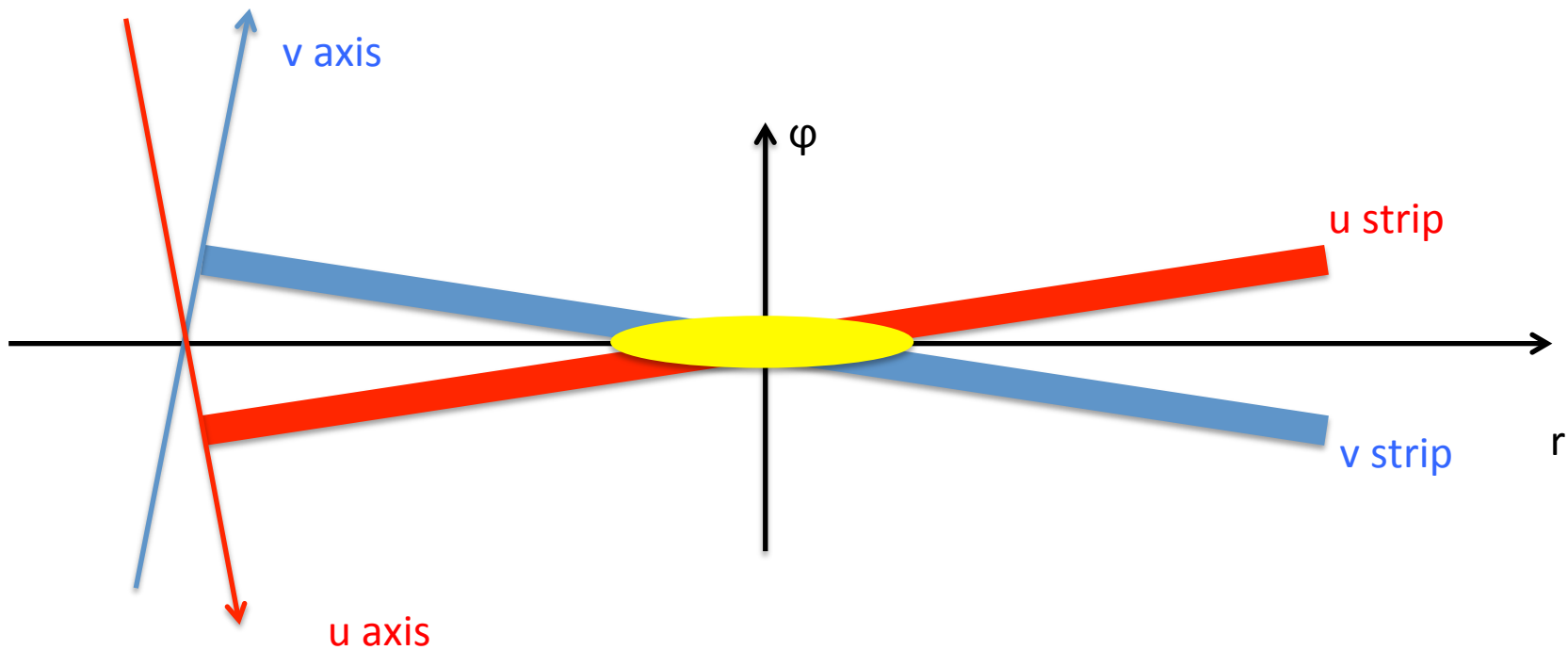
# Conclusion

- Obtained preliminary resolution results for SIDIS,  $J/\psi$  and PVDIS
- Most of the resolutions satisfy requirements, except for polar angle in the SIDIS and  $J/\psi$  configuration
- Improvement quite possible by more careful calculation of the error matrices
- Main challenge is the electron radiation energy loss

**Backup Slides**

# General Info about the Simulation

- Generator is uniform, reconstruct trajectory as long as the particle hits all the GEMs and the calorimeter
- GEM resolution is added according to:



# Introduction on Kalman Filter

- Basic steps for Kalman Filter<sup>[1]</sup>:
  - The optimized state vector  $a_{k-1}$  on detector k-1 is extrapolated to detector k by means of a propagation method

$$a_k^{k-1} = f_{k-1}(a_{k-1})$$

- The covariance matrix of the predicted state vector, and also the covariance matrix of the process noise between detector k-1 and k are computed by error propagation

$$C_k^{k-1} = F_{k-1} C_{k-1} F_{k-1}^T + Q_{k-1}$$

$$F_{k-1} = \frac{\partial f_{k-1}(a_{k-1})}{\partial a_{k-1}}$$

# Introduction on Kalman Filter

- Basic steps for Kalman Filter:
  - The predicted state vector and its covariance matrix are projected into the measurement space by means of a projector  $H$ , thus obtain the predicted measurement vector
  - The weighted mean of the extrapolated and the actual measurement vector  $m_k$  of detector  $k$  is computed, yielding an optimal estimate of the state vector at  $k$

$$C_k = (I - K_k H_k) C_k^{k-1}$$

$$a_k = a_k^{k-1} + K_k (m_k - H_k a_k^{k-1})$$

$$K_k = C_k^{k-1} (H_k)^T (V_k + H_k C_k^{k-1} (H_k)^T)^{-1}$$

In the simplest case, assume 1d state vector  $x$ , and we directly measure this quality, then the Kalman filter formulae can be reduced to a simple form

$$\frac{x_o}{\sigma_o^2} = \frac{x_p}{\sigma_p^2} + \frac{x_m}{\sigma_m^2}$$



# Classical 4<sup>th</sup> Order Runge-Kutta

First of all, take the derivative of the state vector with respect to  $z$ , this is the differential equation that is going to be solved by the Runge-Kutta method:

$$\frac{d\vec{a}}{dz} = \begin{pmatrix} dx/dz \\ dy/dz \\ dt_x/dz \\ dt_y/dz \\ d(q/p)/dz \end{pmatrix} = \begin{pmatrix} t_x \\ t_y \\ t'_x \\ t'_y \\ 0 \end{pmatrix} = f(\vec{a}, z)$$

The solution of the 4<sup>th</sup> order Runge-Kutta method is:

$$\Delta\vec{a}_1 = h \cdot f(\vec{a}(z_0), z_0)$$

$$\Delta\vec{a}_2 = h \cdot f(\vec{a}(z_0) + \frac{1}{2}\Delta\vec{a}_1, z_0 + \frac{1}{2}h)$$

$$\Delta\vec{a}_3 = h \cdot f(\vec{a}(z_0) + \frac{1}{2}\Delta\vec{a}_2, z_0 + \frac{1}{2}h)$$

$$\Delta\vec{a}_4 = h \cdot f(\vec{a}(z_0) + \Delta\vec{a}_3, z_0 + h)$$

$$\vec{a}_f = \vec{a}_0 + \frac{1}{6}\Delta\vec{a}_1 + \frac{1}{3}\Delta\vec{a}_2 + \frac{1}{3}\Delta\vec{a}_3 + \frac{1}{6}\Delta\vec{a}_4 + O(h^5)$$

# Classical 4<sup>th</sup> Order Runge-Kutta

- The propagator matrix of this process is:

$$F = \frac{d\vec{a}_f}{d\vec{a}_0} = I + \frac{1}{6}F_1 + \frac{1}{3}F_2 + \frac{1}{3}F_3 + \frac{1}{6}F_4$$

$$F_i = \frac{d\Delta\vec{a}_i}{d\vec{a}_0}$$

- Using this method, there is no need for a pre-defined geometric track. To initialize the fit,  $x_0$  and  $y_0$  can be taken from the last GEM tracker along the beam direction,  $tx_0$  and  $ty_0$  can be calculated using the last two GEM trackers (where field is low and track is almost straight), finally  $p$  is given by the calorimeter (we always have a EC hit for PVDIS)

# Material Properties

	Radiation Length (cm)	Thickness (cm)
GEM	~302.2	1.55
SIDIS Cell (GE180)	7.038	0.1
J/ $\psi$ Cell (Al)	8.897	0.0178
PVDIS Cell (Al)	8.897	0.038
LH2	890.4	1.9
LD2	769.1	2.5

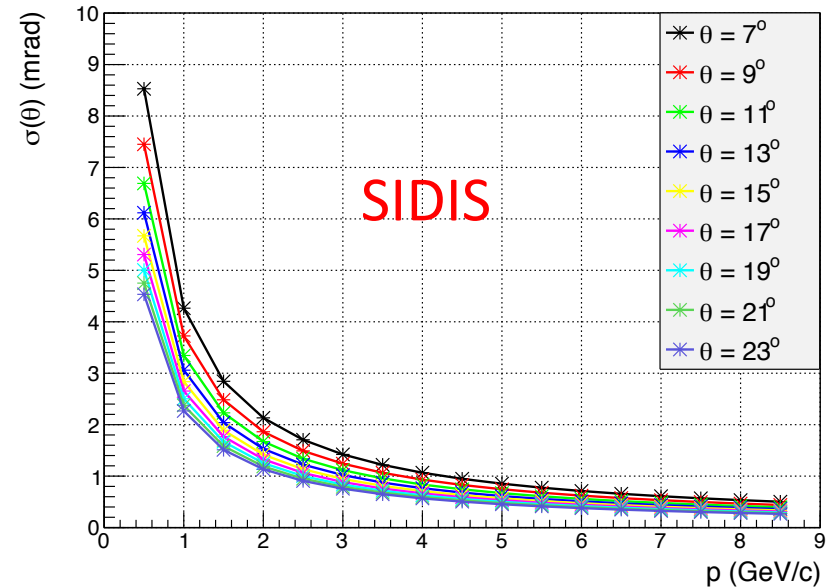
- Some simple calculations for t:

	SIDIS Cell	J/ $\psi$ Cell	PVDIS Cell	GEM
10°	0.0818	0.0115	-	0.0052
20°	0.0415	0.0058	-	0.0055
30°	-	-	0.0085	0.0059

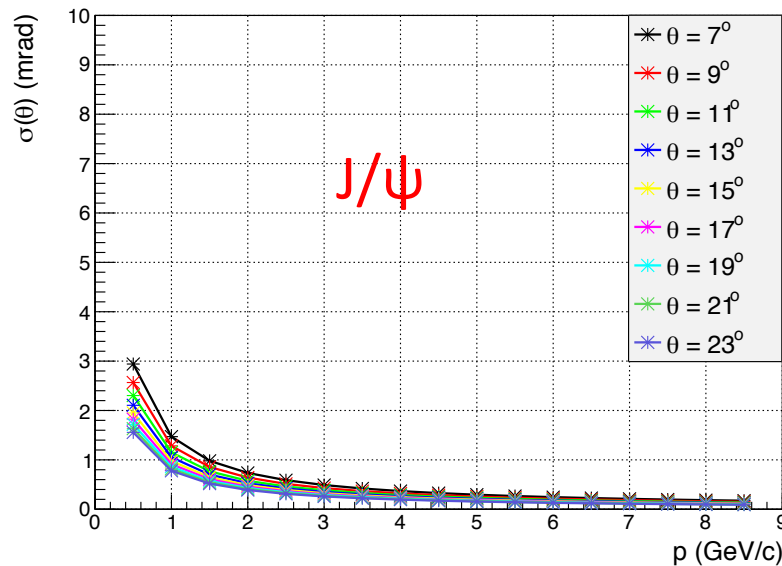
# Polar Angle Resolution Limit

- Best guess for polar angle (for now): the polar angle inside the cell is the same as the polar angle outside
- In this case, polar angle resolution cannot be better than results given by the Moliere formula for particles pass through target cell

Polar angle Resolution



Polar angle Resolution



Polar angle Resolution

