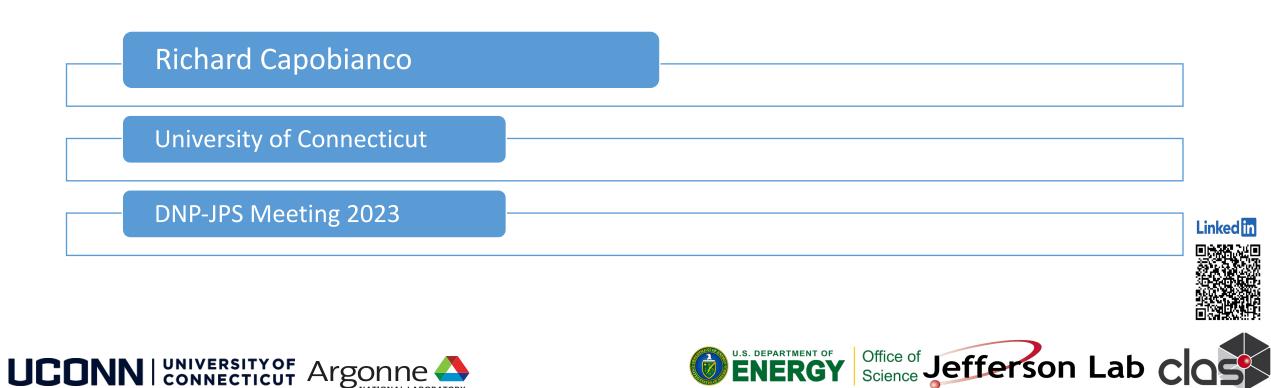
Measurements of the Cos ϕ and Cos2 ϕ Moments of the Unpolarized SIDIS π^+ Cross-section at CLAS12



Motivation

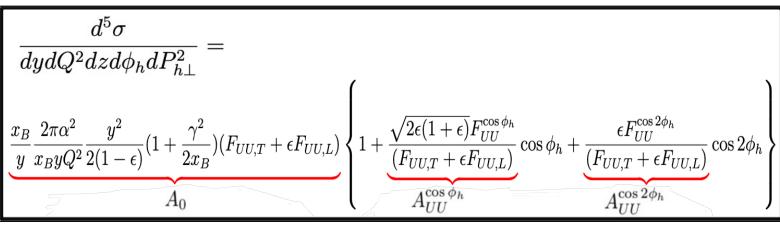
- Semi-Inclusive Deep Inelastic Scattering (SIDIS) experiments allow us to address questions about the 3D structure of nucleons
- Azimuthal modulations in unpolarized SIDIS cross-section for charged pion electroproduction can give access to the Cahn and Boer-Mulders effects
 - **Boer-Mulders Effect:** Sensitive to the correlation between the quark's transverse momentum and intrinsic transverse spin in an unpolarized nucleon
 - Cahn Effect: Sensitive to the transverse motion of quarks inside the nucleon
- A non-zero Boer-Mulders requires quark orbital angular momentum contributions to the proton spin (aspect of the proton missing spin puzzle)



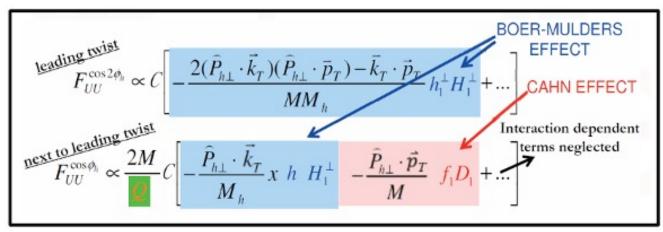


SIDIS Cross-Section and Boer-Mulders

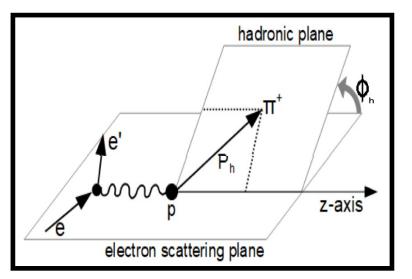
The lepton-hadron Unpolarized SIDIS Cross-Section:



The Boer-Mulders and Cahn effects are present in the Structure Functions:





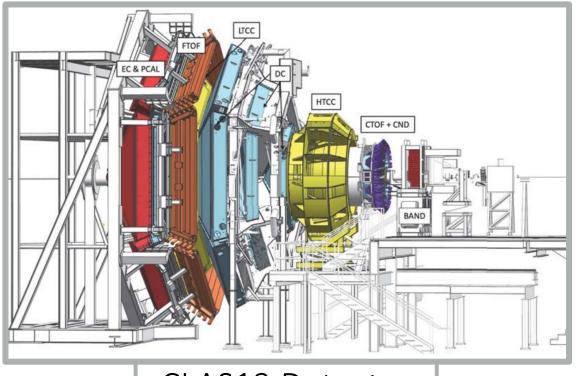


Reaction Studied: $ep \rightarrow e\pi^+(X)$





Data Collection



CLAS12 Detector

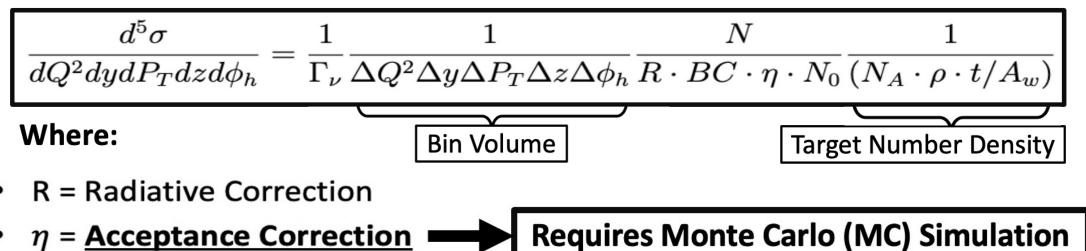
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- CLAS12 detector in Hall B at Jefferson Lab
 - \circ $\,$ Upgrade from the CLAS detector $\,$
 - Enabled the higher energy and statistics for our experiments, not previously accessible
- Data from the Fall 2018 RG-A experiment
 - Used a 10.6 GeV polarized electron beam and unpolarized liquid hydrogen target
- Data presented uses forward tracking only



Analysis Procedure

Experimental extraction of cross-section



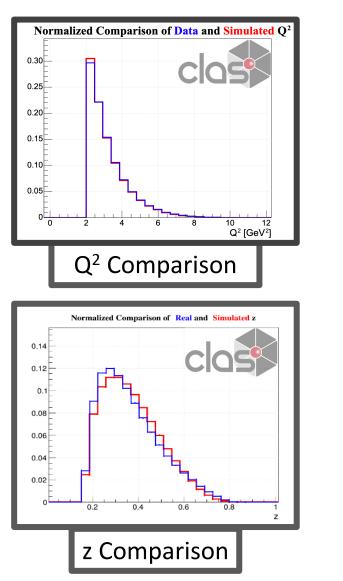
- N = Bin Yields
- N₀ = Life-time corrected incident electron flux
- BC = factor which evolves bin-averaged differential cross-section

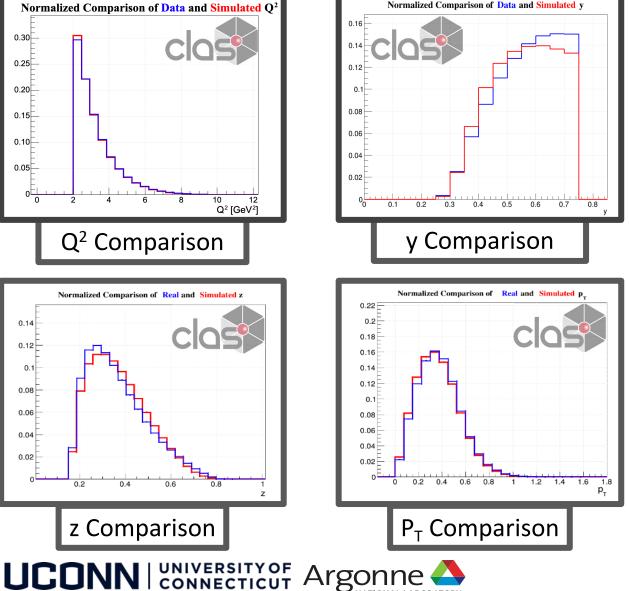
SIDIS MC are generated with LEPTO event generator

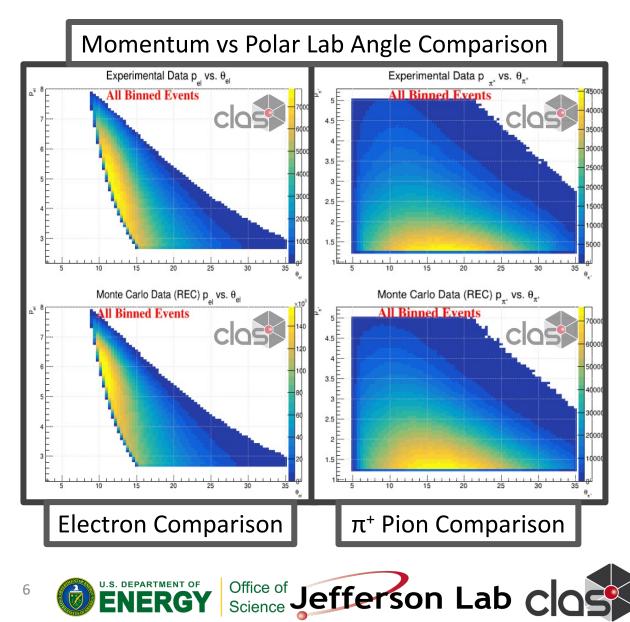




Data and Monte Carlo Comparison



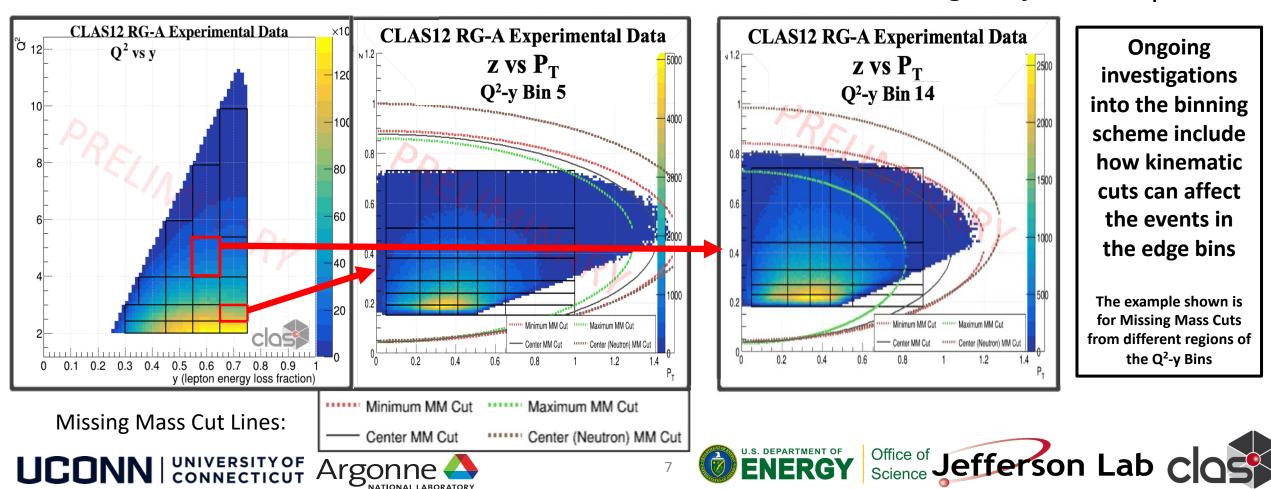




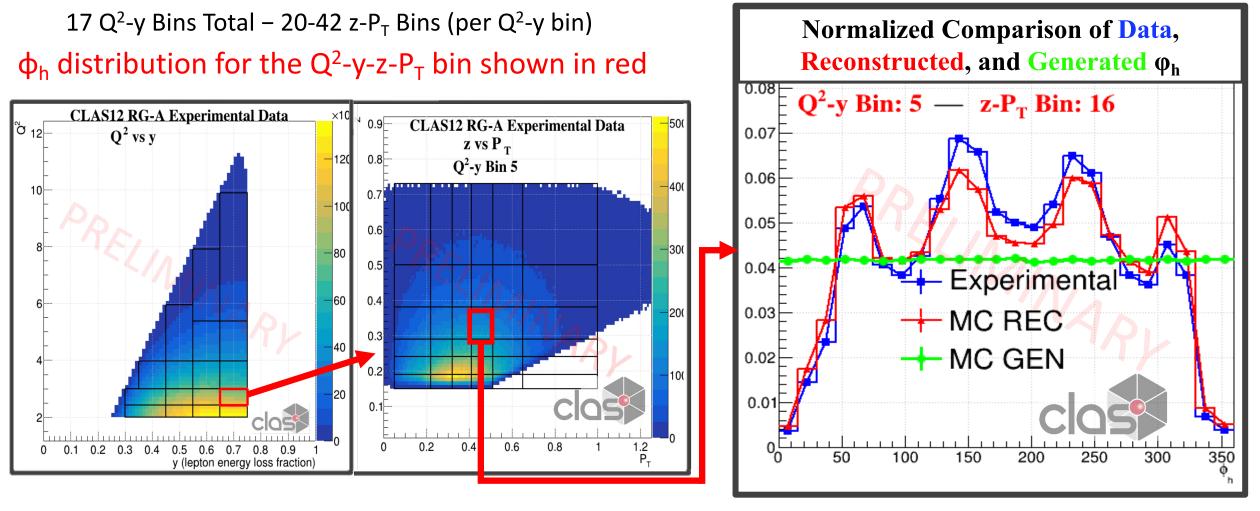
Multidimensional Kinematic Binning (4 Dimensions)

17 Q²-y Bins Total – 20-42 z-P_T Bins (per Q²-y bin)

Examples of new binning scheme using Q^2 , y, z, and P_T



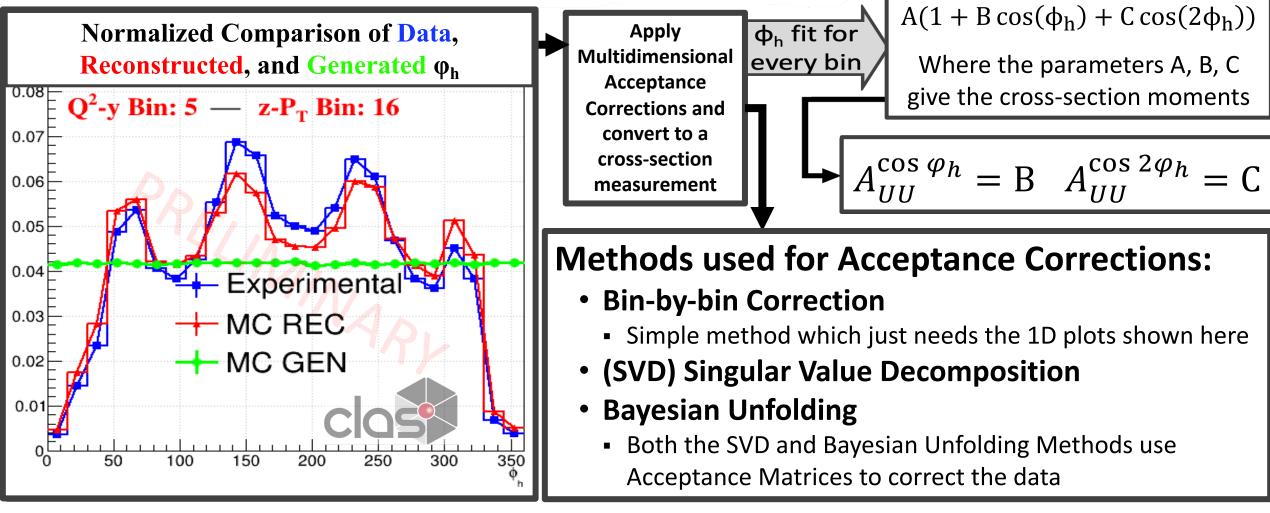
Multidimensional Kinematic Binning (5 Dimensions)





Multidimensional Kinematic Binning (5 Dimensions)

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Acceptance Corrections and Bin Migration Study

<u>Acceptance Matrix</u>: A_(i, j) describes both Acceptance (including geometric acceptance and detector efficiency) and Bin Migration

• $A_{(i, j)} = \frac{\text{Number of Events Generated in bin } j \text{ but Reconstructed in bin } i}{\text{Total Number of Events Generated in the } j \text{th bin}}$

• Acceptance Unfolding: $Y_i = A_{(i,j)}X_j \Leftrightarrow X_j = A_{(i,j)}^{-1}Y_i$

where:

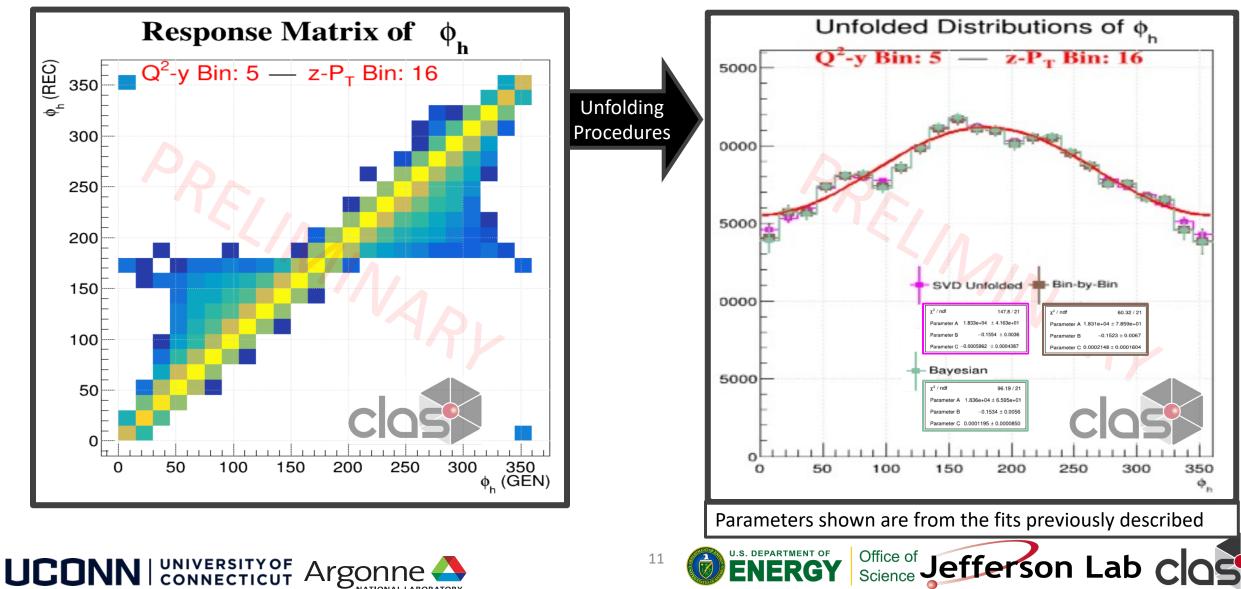
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- Y_i = Number of events experimentally measured in the *i*-th bin
- X_j = Number of acceptance-corrected events in the *j*-th bin



Example of (1D) Unfolding Procedure

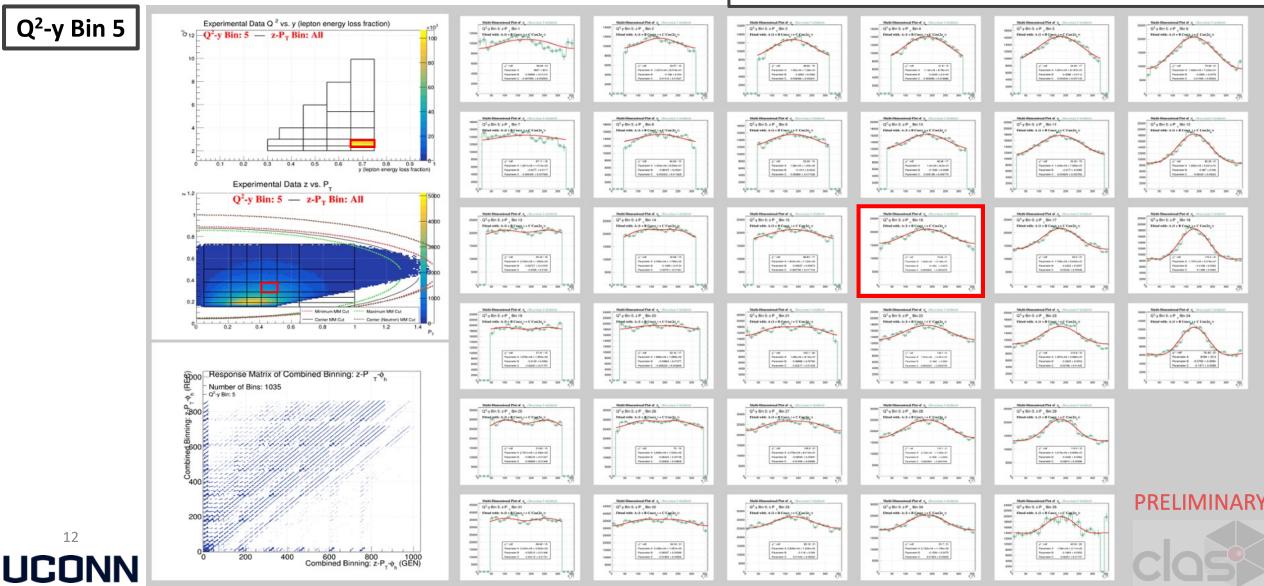
Using the Multidimensional Kinematic Bin from prior example



Example of (3D) Unfolding Procedure

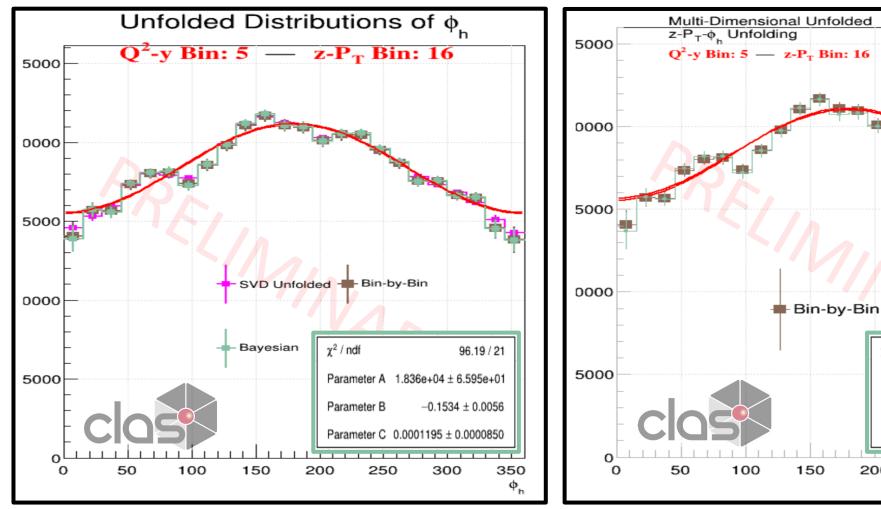
Using the Flattened z-P_T- ϕ_h Multidimensional Bins

Unfolded with Bayesian Method



Comparisons of 1D and 3D Unfolding Procedure

Using the Multidimensional Kinematic Bin from the prior example for this comparison



Bin-by-bin Acceptance Correction gives the exact same results

Bayesian

Parameter A 1.834e+04 ± 8.146e+01

Parameter C-0.0005845 ± 0.0004378

250

70.96 / 21

350 Φ.

 -0.1454 ± 0.0070

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300

 γ^2 / ndf

200

Parameter B

Bayesian Unfolding gives similar results

SVD Unfolding has not been able to work so far with the Multidimensional Unfolding procedures

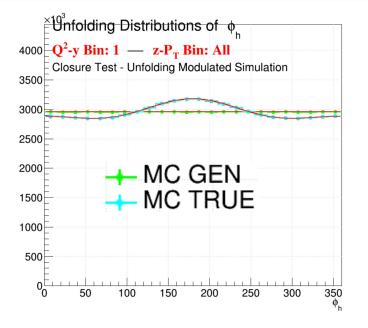


Modulated Unfolding Closure Tests

• Modulated the MC distributions using the formula:

Weight = $1 + B \cos(\phi_h) + C \cos(2\phi_h)$

- Gives the weight for each MC event based on generated $\varphi_{\rm h}$
- Parameter values currently being used in this image:
 - B = -0.05
 - C = 0.025 (Same for every $z P_T bin$)



- Modulated MC REC is then unfolded using the un-modulated response matrix (in 1D and Multi-Dim examples) and compared with 'MC TRUE'
 - MC TRUE is the modulated MC GEN distribution
 - Also performed a closure test of unfolding the un-modulated MC REC distribution with the un-modulated response matrix to ensure the method was applied properly



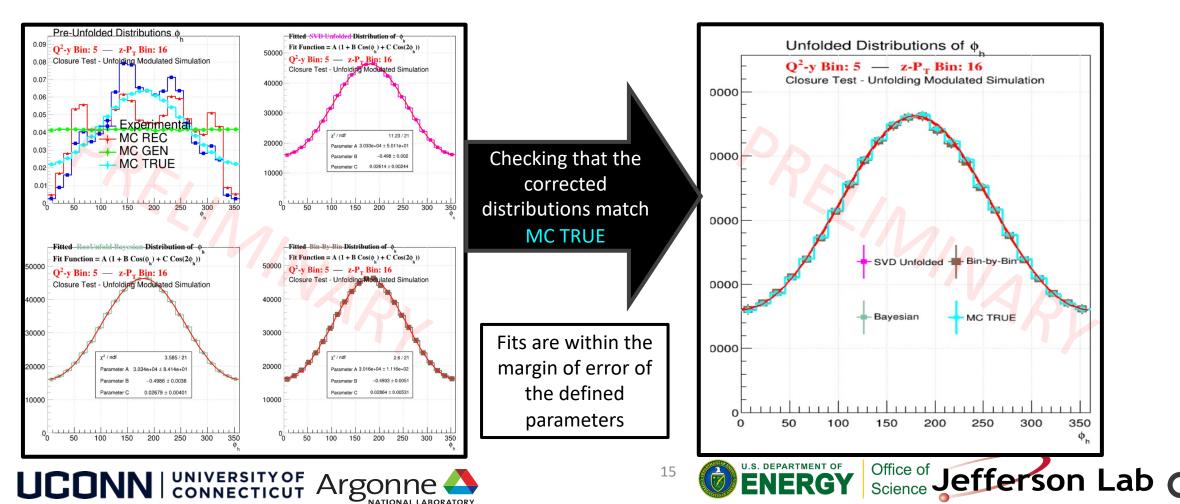


Modulated Unfolding Closure Tests

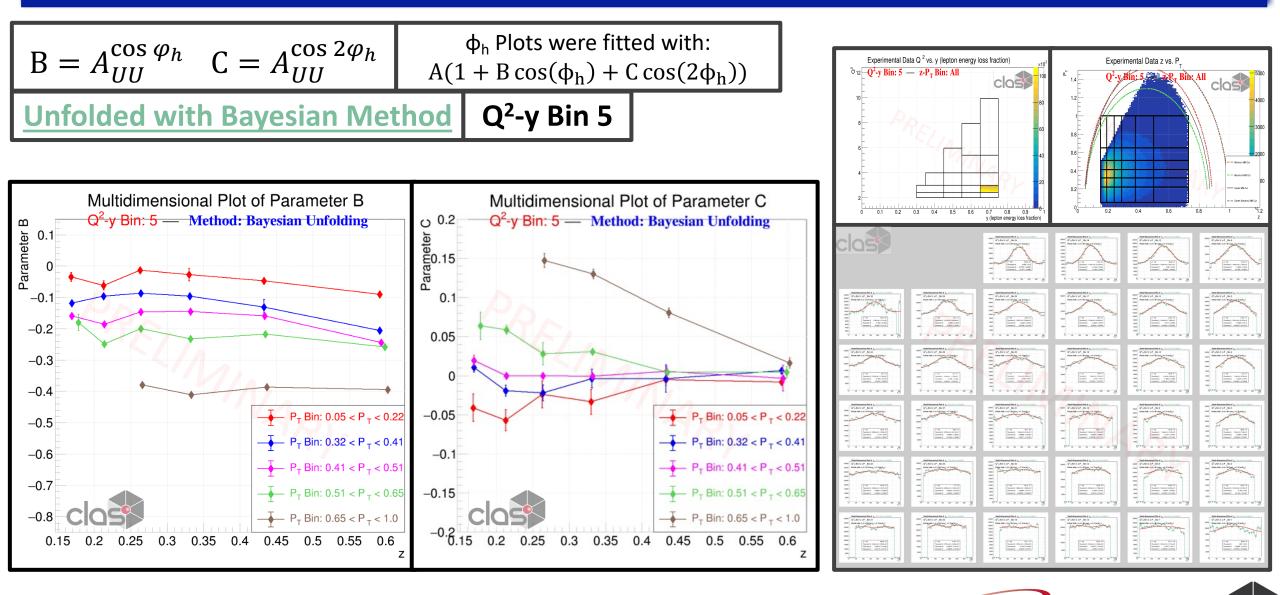
The parameters used for weighing modulations below are:

B = -0.5 and C = 0.025

Results show that an unmodulated Simulation can correct distributions with modulations



Cosine Moments as Functions of z



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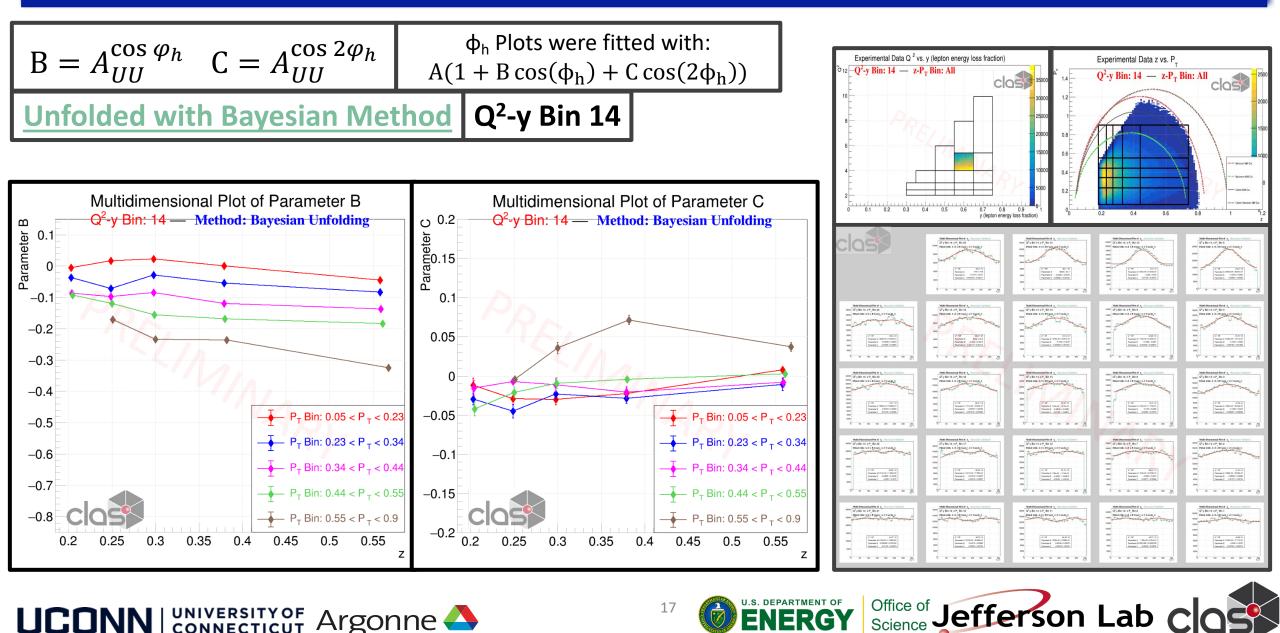
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Cosine Moments as Functions of z



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<u>Outlook</u>

- Working on Multidimensional Acceptance Corrections for the simultaneous unfolding of Q², y, z, P_T, and ϕ_h variables
- Efforts towards more realistic MC simulations, both on the detector response description and physics process
- Include Radiative and BC Corrections to this analysis
- Final goal is the extraction of multiplicity $(F_{UU,T} + \varepsilon F_{UU,L}), F_{UU}^{\cos \varphi_h}$, and

 $F_{UU}^{\cos 2\varphi_h}$ in terms of in Q², y, z, and P_T for the π^+ for all CLAS12 RG-A data







Questions?

Acknowledgments and Thanks

- Contributions made by other members of the CLAS Collaboration and researchers at Argonne National Lab
- This work is supported by the U.S. Department of Energy, Office of Science, Office of Nuclear Physics under contract number DE-AC02-06CH11357



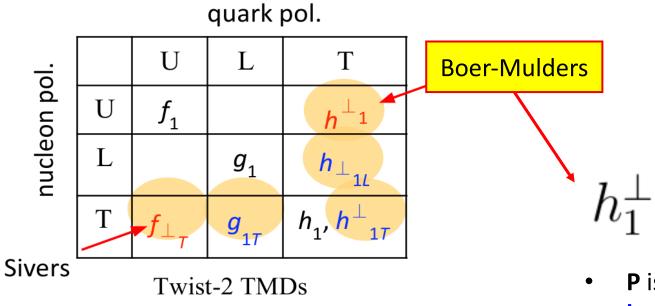


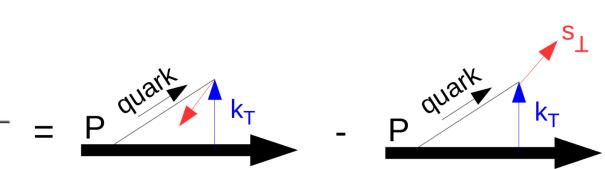
Backup Slides





More on Boer-Mulders...





- **P** is the momentum of the proton
- \mathbf{k}_{T} is the transverse momentum of the quark
- **s**₁ is the transverse spin of the quark

If the Boer-Mulders term is non-zero, then there is a net transverse quark polarization inside of unpolarized protons





Event Selection (Full PID)

The RG-A Analysis Overview and Procedures note goes into detail about the common particle identification scheme used for RG-A

(See: <u>https://clas12-docdb.jlab.org/DocDB/0009/000949/001/RGA_Analysis_Overview_and_Procedures-08172020.pdf</u>)

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Electron PID Criteria:

- Detected in Forward Detector
- > 2 photoelectrons detected in the HTCC
- > 0.07 GeV energy deposited in the PCAL
- Sector dependent sampling fraction cut
- "Diagonal cut" for electrons above 4.5 GeV (HTCC threshold)
- y < 0.75, not strictly an "electron cut", but sets the min electron energy approximately > 2.4 GeV

Pion PID Criteria:

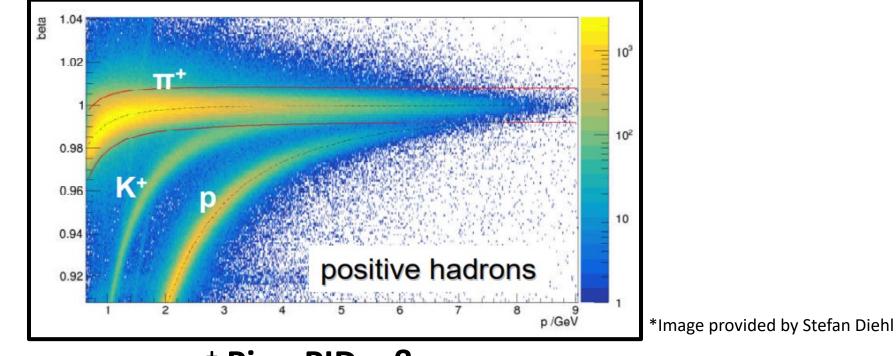
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- Detected in Forward Detector
- p > 1.25 GeV
- Refined chi2pid cuts



Particle ID (PID):

- **Electron ID:** Based on Electromagnetic Calorimeter (PCAL) and Cherenkov Counters (HTCC)
- Hadron (π^+) ID: Based on Time-Of-Flight Counters (TOF) and the correlation of velocity (β) and momentum



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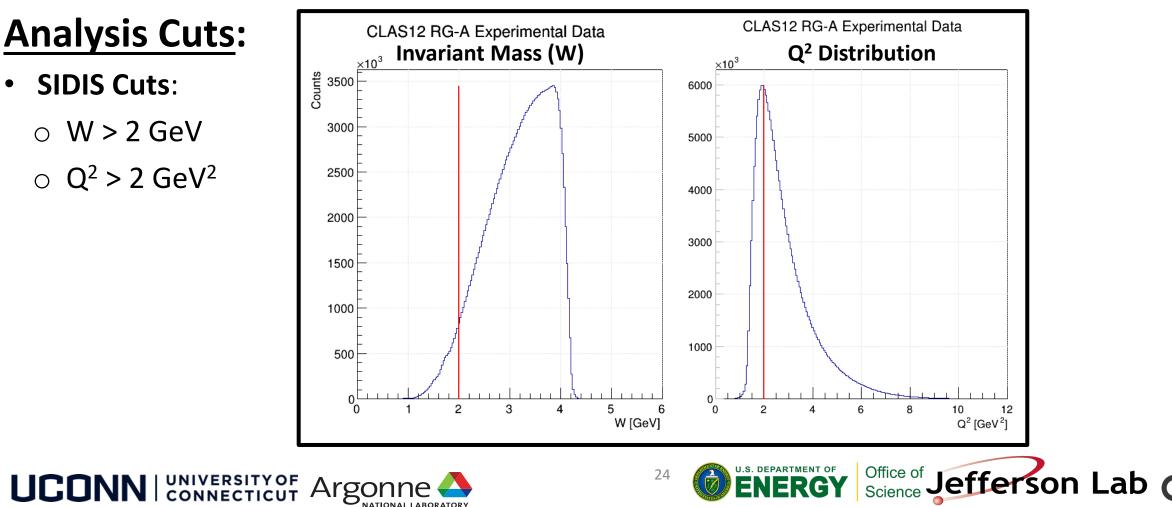
π^+ Pion PID – ß vs p





Particle ID (PID):

- Electron ID: Based on Electromagnetic Calorimeter (PCAL) and Cherenkov Counters (HTCC)
- Hadron (π^+) ID: Based on Time-Of-Flight Counters (TOF) and the correlation of velocity (ß) and momentum



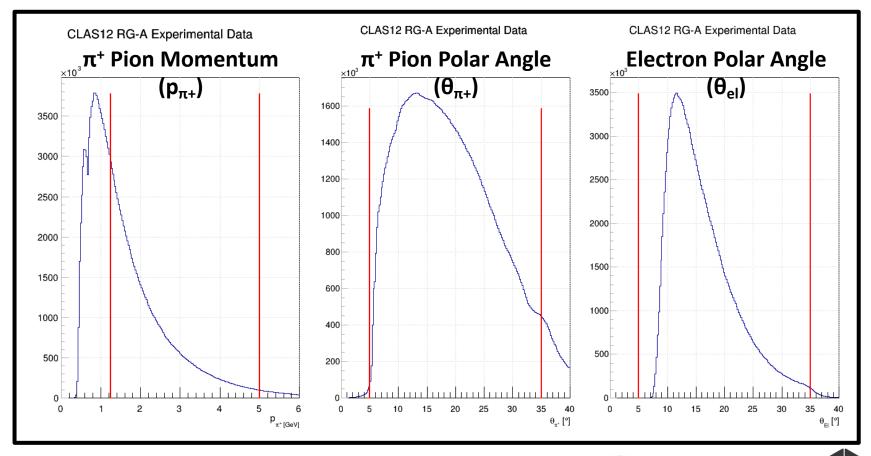
25

Particle ID (PID):

- **Electron ID:** Based on Electromagnetic Calorimeter (PCAL) and Cherenkov Counters (HTCC)
- **Hadron** (π^+) ID: Based on Time-Of-Flight Counters (TOF) and the correlation of velocity (β) and momentum ٠

Analysis Cuts:

- **SIDIS Cuts:**
 - W > 2 GeV
 - $Q^2 > 2 \text{ GeV}^2$ Ο
- **Other Analysis Cuts:**
 - $p_{\pi+}$ Cut: 1.25 GeV < $p_{\pi+}$ < 5 GeV
 - \circ θ-angle Cut: 5° < θ_{particle} < 35°



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Particle ID (PID):

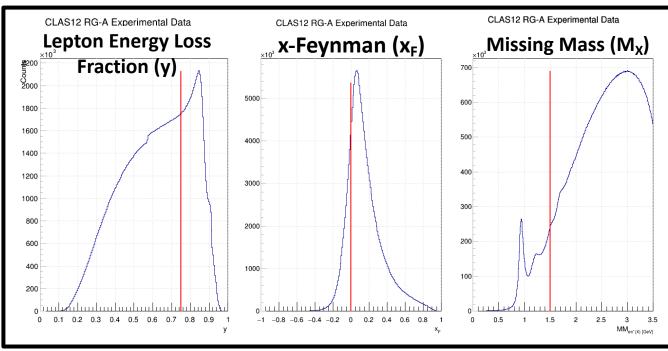
- Electron ID: Based on Electromagnetic Calorimeter (PCAL) and Cherenkov Counters (HTCC)
- Hadron (π^+) ID: Based on Time-Of-Flight Counters (TOF) and the correlation of velocity (β) and momentum

Analysis Cuts:

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 - $\circ \quad Q^2 > 2 \text{ GeV}^2$
- Other Analysis Cuts:
 - $\circ~~p_{\pi^+}$ Cut: 1.25 GeV < p_{π^+} < 5 GeV

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- \circ θ-angle Cut: 5° < θ_{particle} < 35°
- y < 0.75 (minimize other background processes)
- \circ x_F > 0 (minimize contributions from target fragmentations)
- \circ Missing Mass Cut: M_x > 1.5 GeV (limits contributions from exclusive events)



Particle ID (PID):

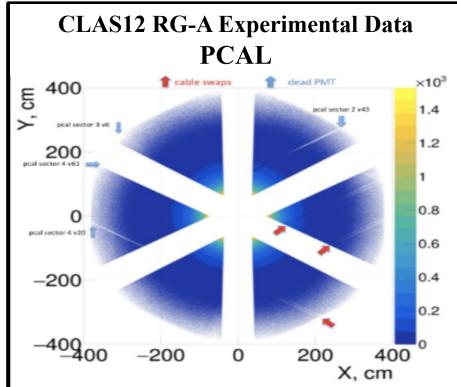
- **Electron ID:** Based on Electromagnetic Calorimeter (PCAL) and Cherenkov Counters (HTCC)
- **Hadron** (π^+) ID: Based on Time-Of-Flight Counters (TOF) and the correlation of velocity (β) and momentum

Analysis Cuts:

- **SIDIS Cuts**:
 - W > 2 GeV
 - $Q^2 > 2 GeV^2$ Ο
- **Other Analysis Cuts:**
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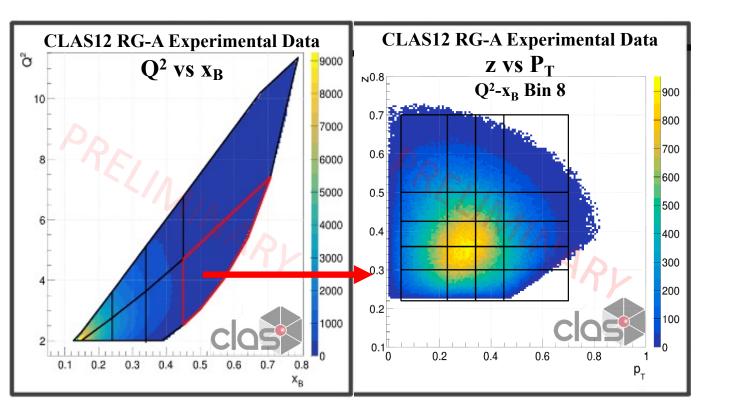
- θ -angle Cut: 5° < $\theta_{particle}$ < 35° Ο
- y < 0.75 (minimize other background processes) Ο
- $x_F > 0$ (minimize contributions from target fragmentations) Ο
- Missing Mass Cut: $M_x > 1.5$ GeV (limits contributions from exclusive events) Ο
- Fiducial Cuts (e.g., accounts for bad channels present in data) Ο



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Multidimensional Kinematic Binning (4 Dimensions)

8 Q²- x_B Bins Total – 20-49 z- P_T Bins (per Q²- x_B bin)



Example of previous binning scheme using Q^2 , x_B , z, and P_T

Main Issue was with the irregular shape of the Q²-x_B Bins

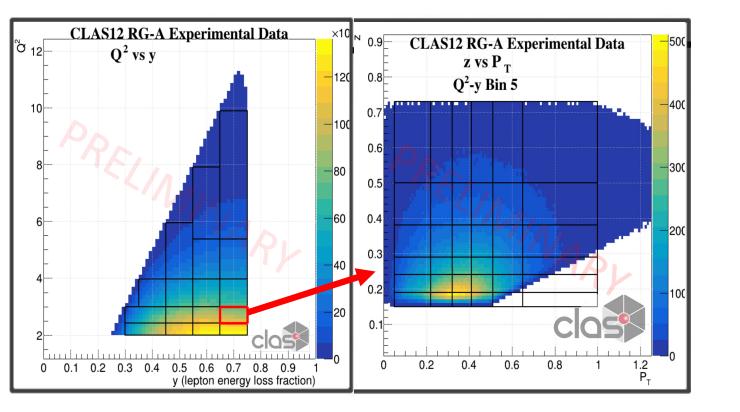






Multidimensional Kinematic Binning (4 Dimensions)

17 Q²-y Bins Total – 20-42 z-P_T Bins (per Q²-y bin)



Example of new binning scheme using Q², y, z, and P_T

Both the Q²-y and z-P_T bins are now rectangular which makes the bins easier to work with

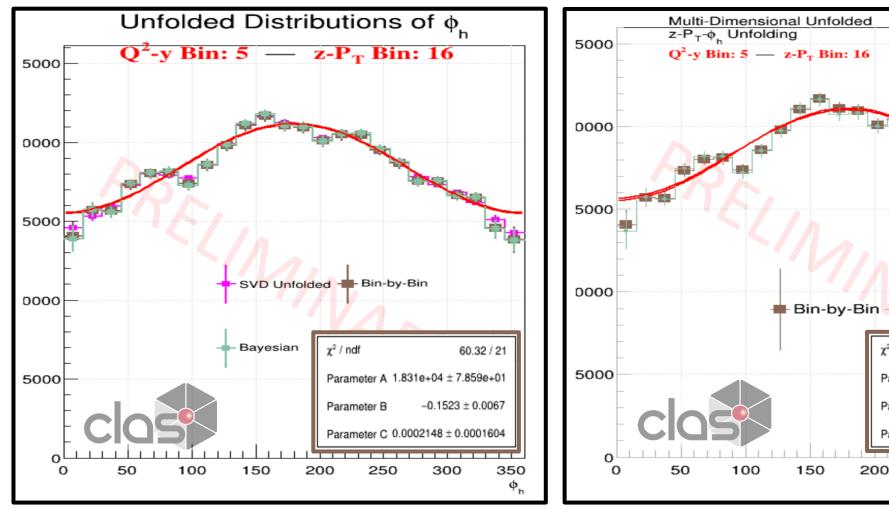






Comparisons of 1D and 3D Unfolding Procedure

Using the Multidimensional Kinematic Bin from the prior example for this comparison



Bin-by-bin Acceptance Correction gives the exact same results

Distributions of

Bayesian

250

Parameter A 1.831e+04 ± 7.859e+01

Parameter C 0.0002148 ± 0.0001604

60.32 / 21

350

Φ.

 -0.1523 ± 0.0067

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300

 χ^2 / ndf

Parameter B

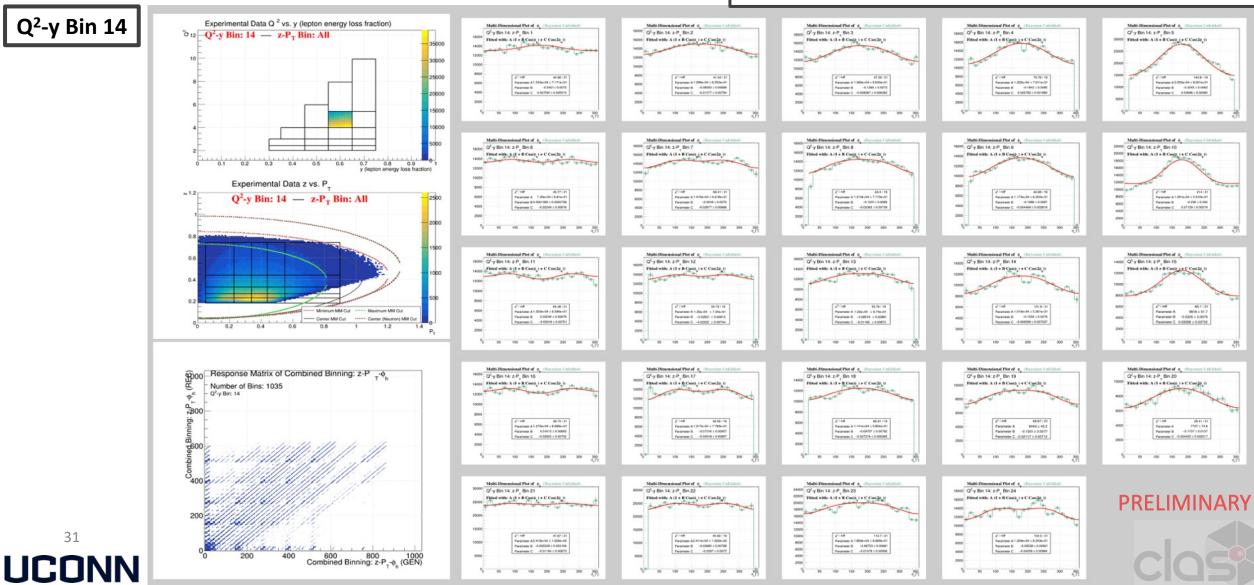
SVD Unfolding has not been able to work so far with the Multidimensional Unfolding procedures



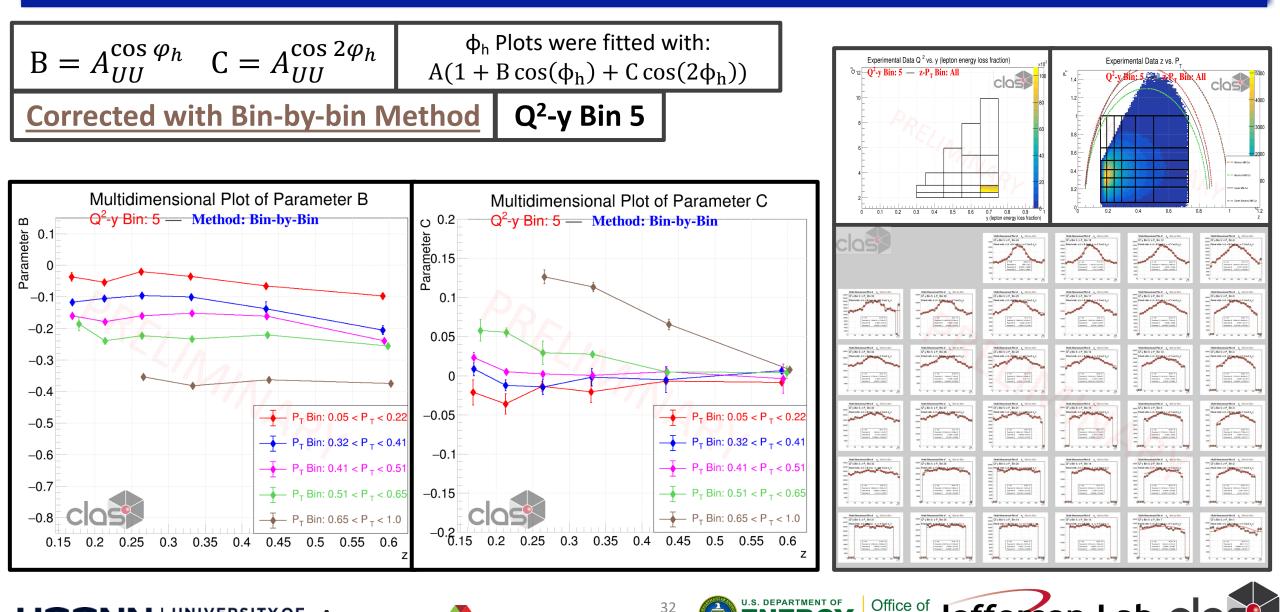
Extra Examples of (3D) Unfolding Procedure

Using the Flattened z-P_T- ϕ_h Multidimensional Bins

Unfolded with Bayesian Method



Cosine Moments as Functions of z



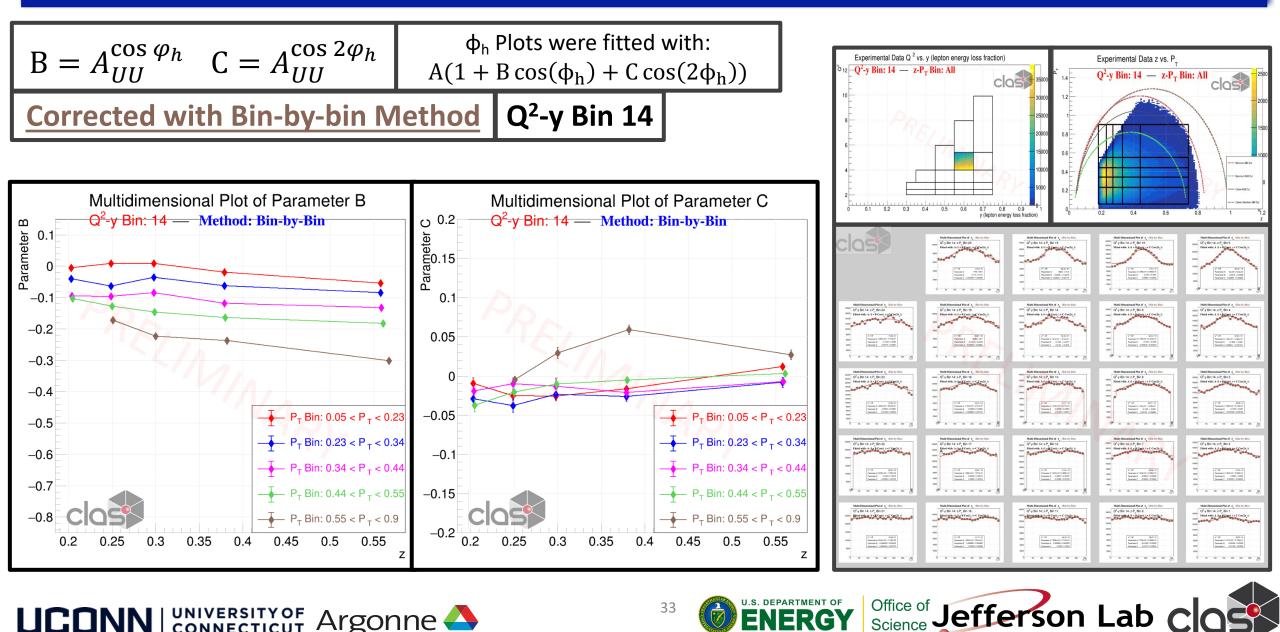
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Cosine Moments as Functions of z



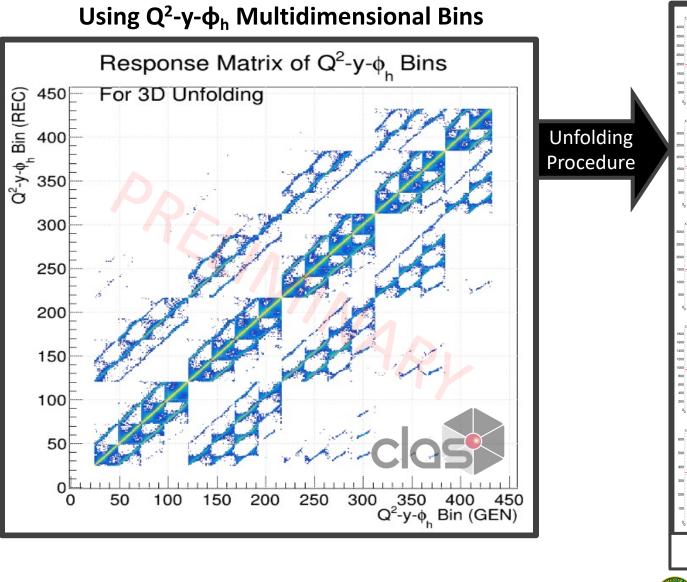
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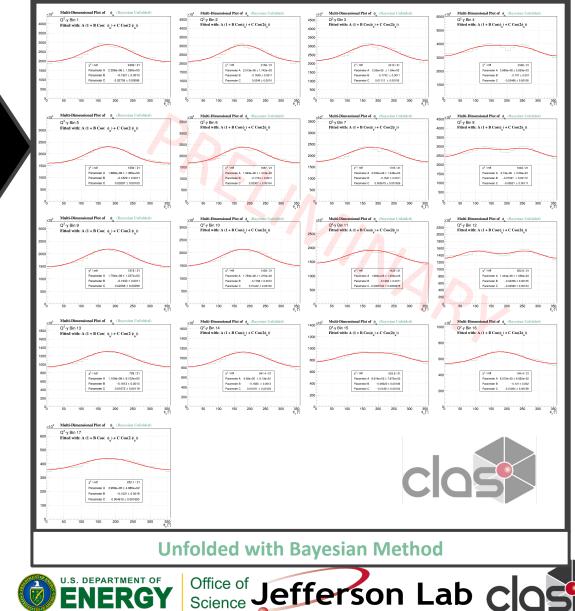
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Example of (3D) Unfolding Procedure



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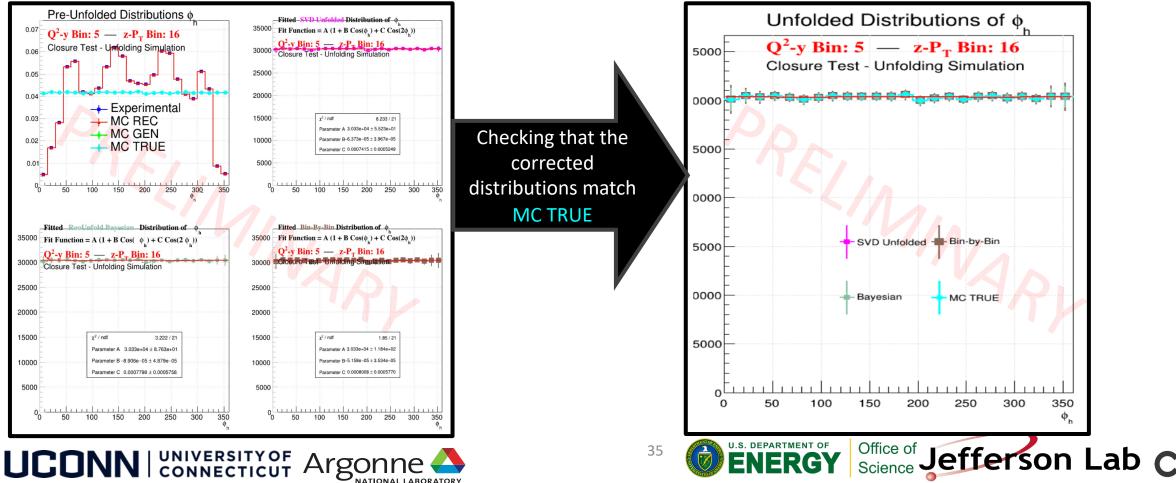


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Other Unfolding Closure Tests

Other closure tests being used to check that Unfolding is done properly:

- Replace the experimental data with the reconstructed Monte Carlo
 - Should return the generated (i.e., MC TRUE) distribution



Momentum Corrections from Exclusive Events

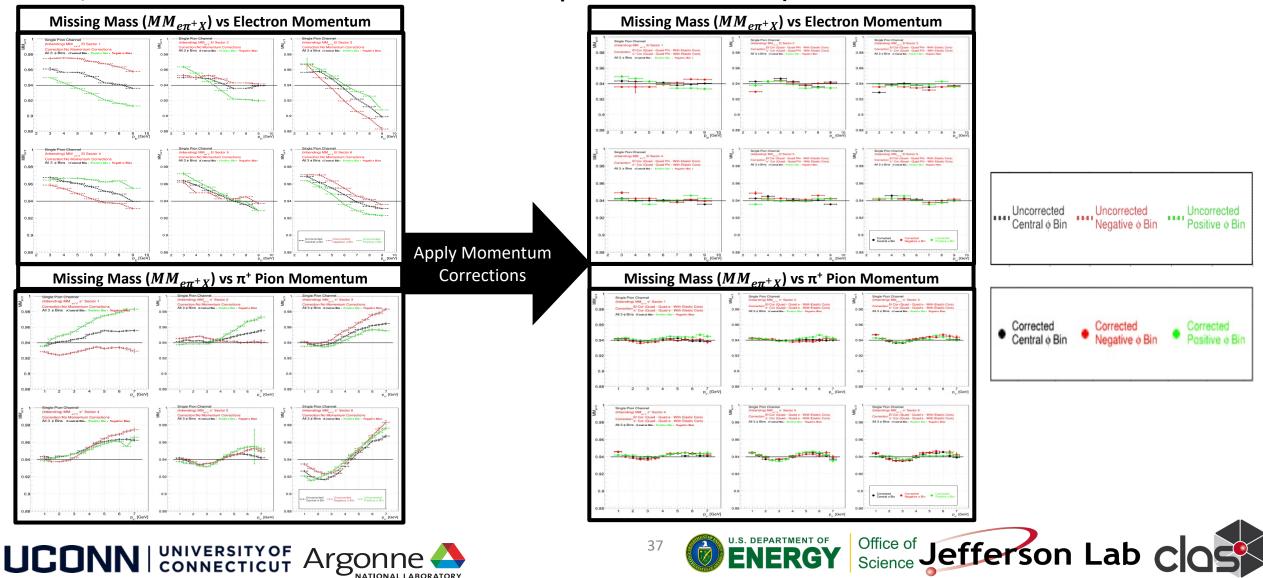
- Momentum corrections are developed for the RG-A data being used in this analysis
- Designed to correct for kinematic-dependent reconstruction issues in the experimental data using well-understood reactions
- Use exclusive reactions to correct the particles' momentum as sector-dependent functions of the particles' measured azimuthal angle (ϕ_{lab}) and momentum
 - The primary reaction used for the electron and π^+ pion is ep $\rightarrow e'\pi^+(N)$
 - Elastic scattering process also used to help correct the electron momentum
- Developed from momentum 4-vector conservation to calculate the ideal momentum of a particle from exclusive reactions based on the kinematics of the other particle(s)
 - Correction is taken by plotting the difference between this calculation and the measured momentum as functions of the measured momentum and ϕ_{lab}



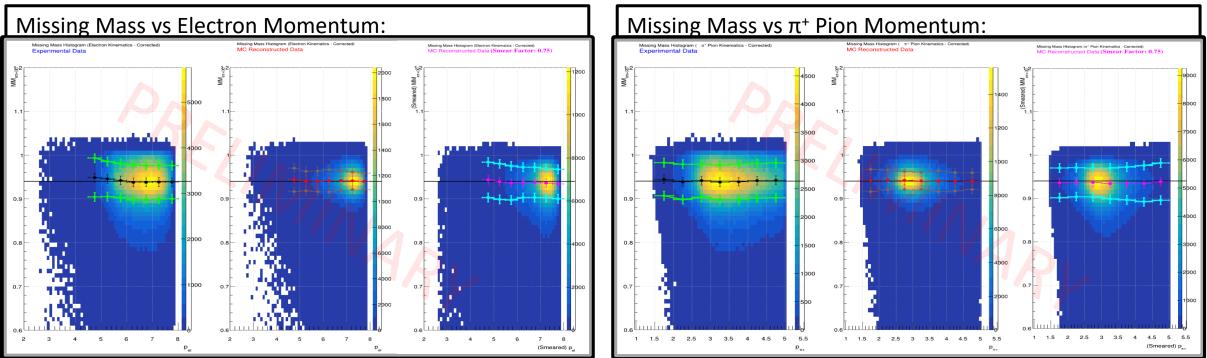


Momentum Corrections from Exclusive Events

These plots show Missing Mass vs. particle momentum in 3 ϕ bins for all 6 sectors of the detector before/after momentum corrections – **Corrections are quadratic functions of \phi and momentum**



Momentum Smearing



- The momentums of the particles in these plots are CORRECTED (see Momentum Corrections from Exclusive Events)
- Momentum Smearing is applied in addition to existing MC reconstruction processes
- The momentum smearing functions use 2D Missing Mass plots to check how it improves the MC
 - The widths of the peaks are shown in each plot above

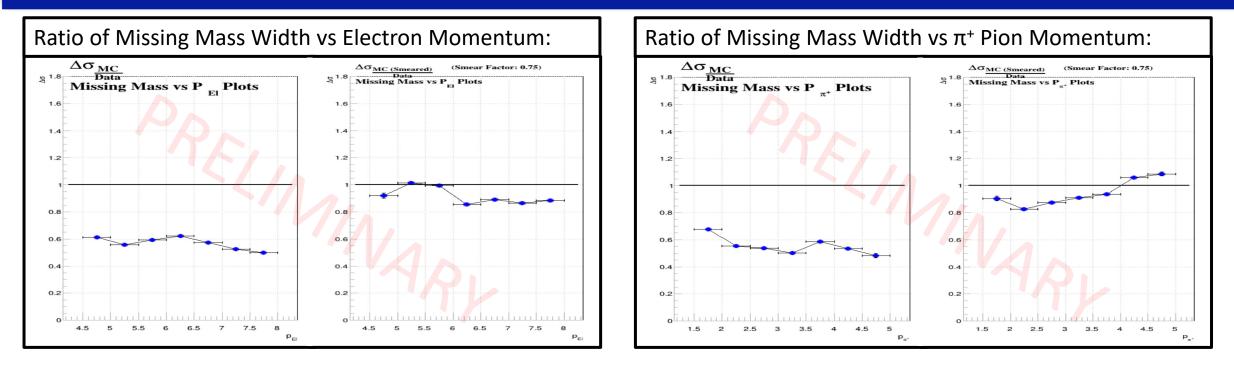
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- Momentum smearing is done with the equation: P_{Smeared} = P_{Generated} (SF + 1)*(P_{Generated} P_{Reconstructed})
 - SF is the smear factor used to modify the simulated reconstructed momentum (currently equal to 0.75)

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• A properly smeared MC distribution should have approximately the same width as the Experimental data

Momentum Corrections/Smearing



- The ratio of the Monte Carlo and Experimental data's widths should go to 1 as smearing improves
- Smearing the momentum also affects the widths of the Missing Mass vs azimuthal/polar angles of the particles
- Development of this correction calls for finding the best smearing parameter for all particle kinematics







Link to more Images:

https://userweb.jlab.org/~richcap/Interactive_Webpage_SIDIS_richcap/Interactive_Unfolding_Page_Updated.html





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