

Update on the extraction of cos moments for kaon SIDIS

Áron Kripkó, Stefan Diehl

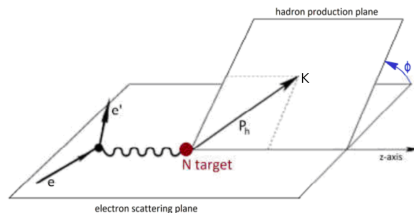
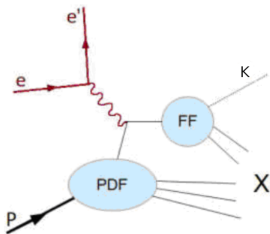
Justus-Liebig University

akripko@jlab.org

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Physics motivation

- Kaon Sidis: $e^- p^+ \rightarrow e^- K^+ X$
- Moments, averaging beam polarisation, un-polarised target:
- $d\sigma = A_0(1 + A_{UU}^{\cos\phi} \cos\phi + A_{UU}^{\cos 2\phi} \cos 2\phi)$



Process (left) and kinematics (right) of single kaon SIDIS

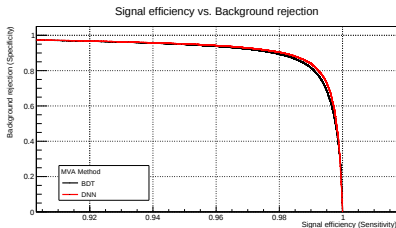
- Structure functions - moments - convolution of FFs and TMDs:
- $F_{UU} = \frac{Q^2(1-\epsilon)A_0}{\pi\alpha^2(1+\frac{\gamma^2}{2x_B})}$
- $\frac{F_{UU}^{\cos\phi}}{F_{UU}} = \frac{A_{UU}^{\cos\phi}}{\sqrt{2\epsilon(1+\epsilon)}} \quad - \quad F_{UU}^{\cos\phi} = \frac{2M}{Q}\zeta \left(-\frac{\hat{h}k_T}{M_h} x h H_1^\perp - \frac{\hat{h}p_T}{M} f_1 D_1 + \dots \right)$
- $\frac{F_{UU}^{\cos 2\phi}}{F_{UU}} = \frac{A_{UU}^{\cos 2\phi}}{\epsilon} \quad - \quad F_{UU}^{\cos 2\phi} = \zeta \left(-\frac{2(\hat{h}k_T)(\hat{h}p_T) - k_T p_T}{MM_h} h_1^\perp H_1^\perp + \dots \right)$
- Access to Boer-Mulders - $h_1^\perp$ - distribution of transversely polarized quarks in an unpolarized hadron
- Cahn effect - kinematic effect due to the intrinsic momentum of quarks in the nucleon

- Eventbuilder particle ID
- Fiducial cuts
- Electron and hadron PID refinements from the pion note:
 - PCAL minimum energy deposition
 - ECAL sampling fraction cut
 - z-vertex position cut
 - Cut on vertex difference
 - $|\chi^2_{PID}| < 3$
- Use machine learning for Kaon ID
- QA cuts
- Topology: at least one good electron and at least one good Kaon
- Use inbending 10.6 GeV (2018) RG-A dataset:
 - 5032-5419

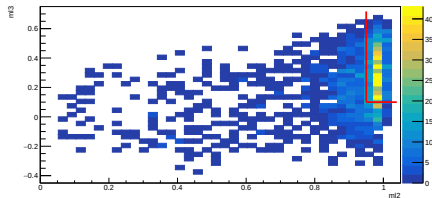
- For optimal PID:
 - $y < 0.75$
 - $1.25 \text{ GeV} < p_K < 3 \text{ GeV}$
 - Only use forward detector for Kaons:
 - $5^\circ < \theta_K < 35^\circ$
 - $5^\circ < \theta_e < 35^\circ$
- To select the deep inelastic scattering region:
 - $W > 2 \text{ GeV}$
 - $Q^2 > 2 \text{ GeV}^2$
- To reject the kaons from the fragmentation region:
 - $x_F > 0$
 - $z > 0.3$
- To reduce the contamination from exclusive processes:
 - $M_X > 1.6 \text{ GeV}$
 - $z < 0.7$

- Reduce pion contamination in the kaon sample
- Use most of the available detector information:
 - Momentum and β
 - θ and ϕ
 - ECAL energy depositions
 - ECAL path/time
 - χ^2_{PID}
 - FTOF energy
 - FTOF path/time
- These inputs are smeared and shifted before training to match data
- Significantly reduces the pion contamination in the kaon sample
- The results were cross-checked with an other MC sample
- DNN - 3 hidden layers, tanh activation function
- BDT - secondary cross check or further tightening

ROC curves and return values



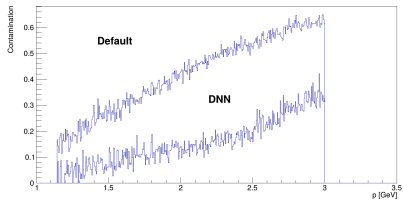
ROC Curves



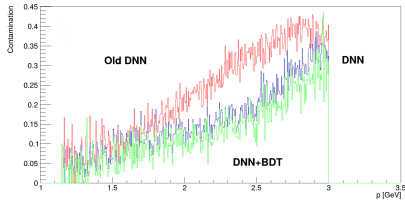
Return values

- Cuts at 0.95 and 0.1
- DNN is slightly better
- BDT can be used for ML related systematics

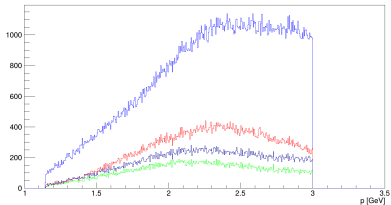
Pion contamination



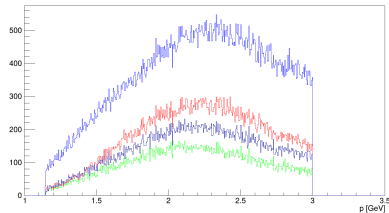
Pion contamination



Contamination with other methods



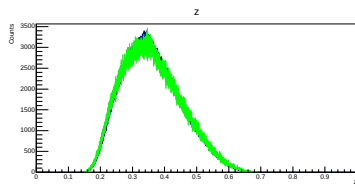
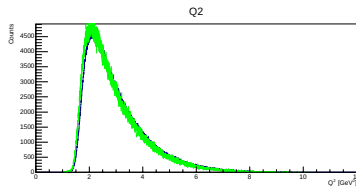
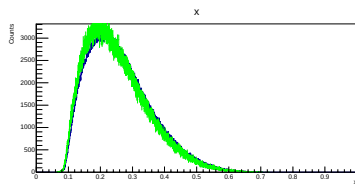
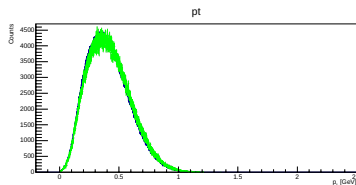
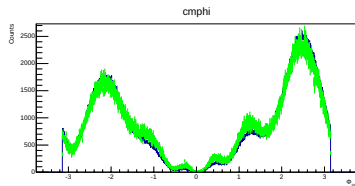
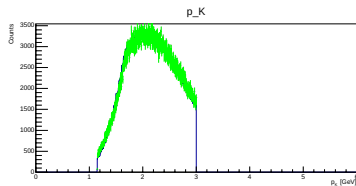
Detected kaons



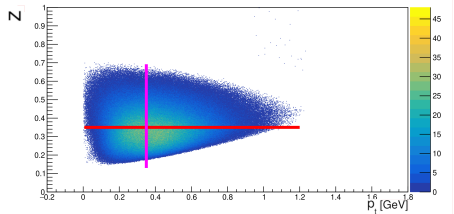
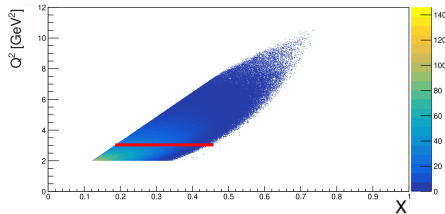
Real kaons

- Good agreement with data
- $\sim 50\%$ statistics to data - more needed
- Low statistics - 1D binning for first look
- Use smearing to get realistic resolution
- Similar acceptance - can be used for unfolding

Kinematics - MC-data comparison



Binning

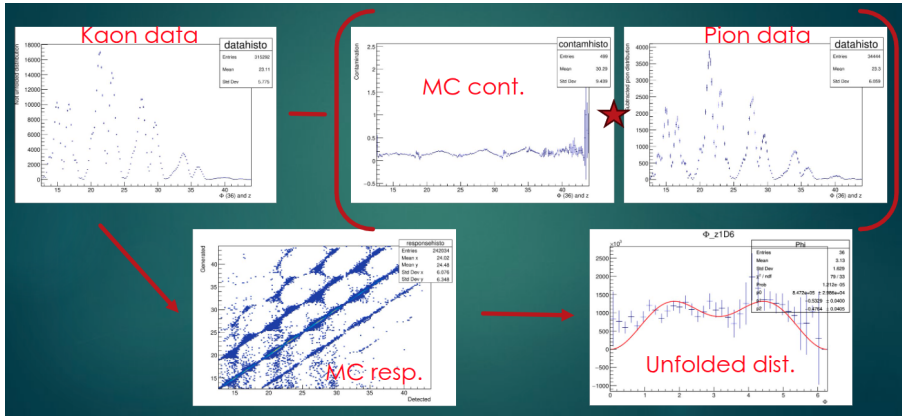


- 4 bins for multi-D
- 4-9 bins with identical widths for 1D or in the last dimension

Unfolding - acceptance corrections and bin migration

- Use the ROOTs RooUnfold package
- Acceptance correction
- Bin migration effects
- Pion contamination - handled before unfolding - MC scaled
- Φ is momentum dependent - pion data is scaled
- Response matrix - matched generated and reconstructed MC
- Generated distribution - includes missed particles
- Reconstructed distribution - does not includes fakes (pions)
- Data $\rightarrow$ Acceptance (& bin migration) corrected distribution with subtracted pions

Unfolding and analysis procedure

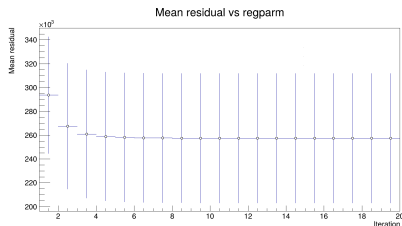
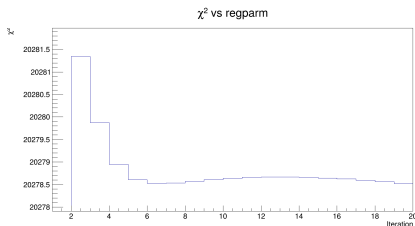


Pion contamination - subtraction

- ϕ is p -dependent
- Contamination is p -dependent
- After DNN $\sim 15\%$ of the kaons are pions
- Kaon distribution from data in flattened bin
- Pion distribution from data in flattened bin - scaled to have same p -distribution
- Misidentified pions from MC in flattened bin - scaled to have same p -distribution
- "Pure" kaon data is then unfolded with MC response only using matched kaons
- Assume that kaon contamination in pions are negligible
- Assume that MC contamination is accurate

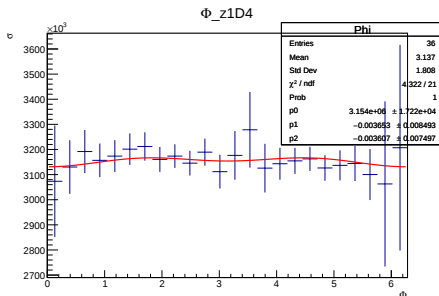
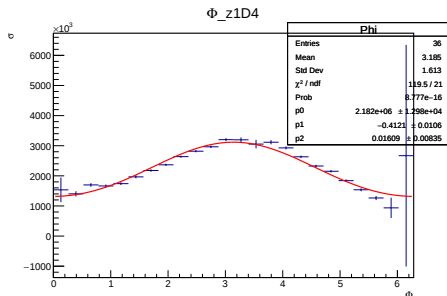
Bayesian unfolding

- Multiple unfolding methods implemented - best is Bayesian
- Flattened bins - correction factor for every kinematic and ϕ bin
- Number of iterations optimized for every binning scheme
- Example for Q^2 kaon binning scheme:

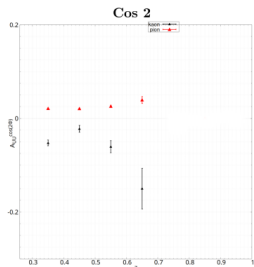
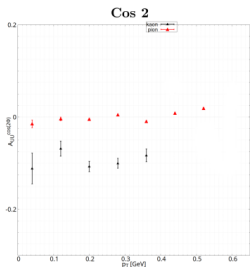
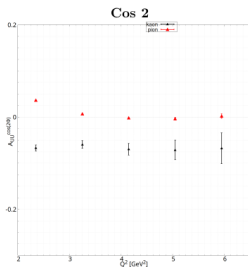
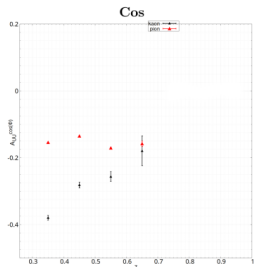
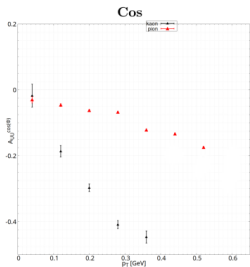
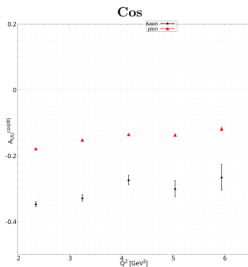


Extracting parameters

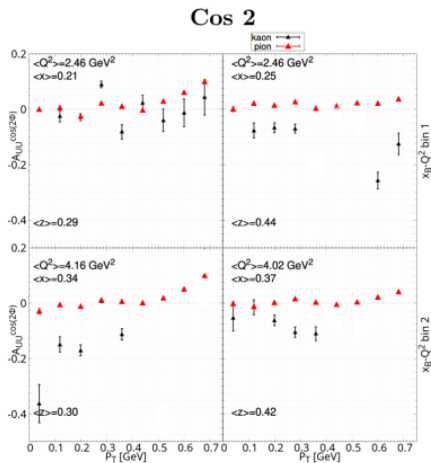
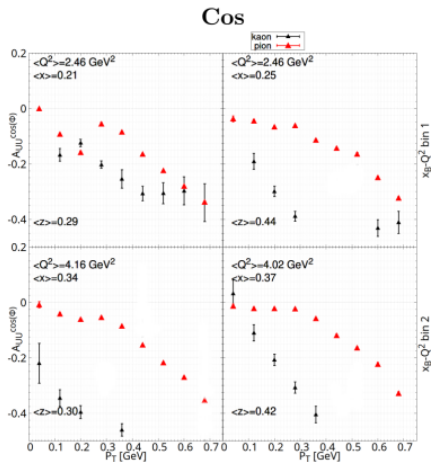
- $d\sigma = A_0(1 + A_{UU}^{\cos\phi} \cos\phi + A_{UU}^{\cos 2\phi} \cos 2\phi)$ - fit unfolded phi distribution
- Closure test - repeat whole process but on MC - should be flat
- Advanced closure test in progress - Injected moments



Results - $Q^2 - p_T - z$



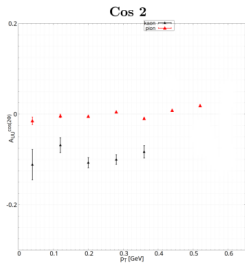
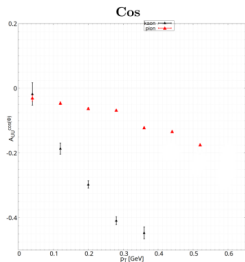
Outlook - multi-D results



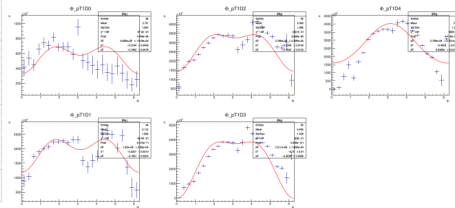
Summary and outlook

- Unfolding framework implemented and now stable
- Pion contamination is under control - new, optimized DNN
- 1D binning is possible
- More MC statistics is needed in the future - multi-D
- Closure tests are promising
- Systematic studies are in progress
- At least 10 times more MC is needed

Results - p_T



Data Fit



Closure Test

