Omega Cross-Section

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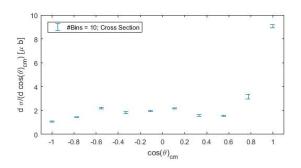


Figure 1: Olis Cross Section; Dip at about $cos(\theta) = -0.3$

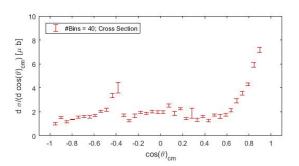


Figure 2: Increased number of bins to 40; now there is still a dip at $\cos(\theta) = -0.3$ but also a peak at $\cos(\theta) = -0.5$



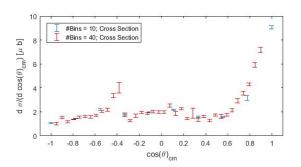
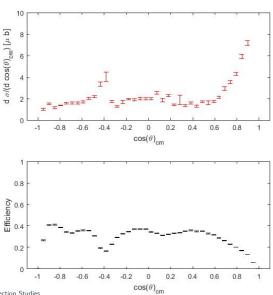


Figure 3: Both Cross Sections are shown.



Bachelor Photon

$$\omega \to \gamma \ \pi^0$$

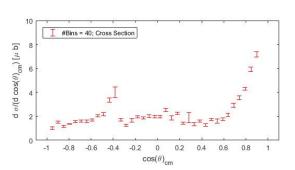
Closer look at:

• u

π⁰

γ

Proton



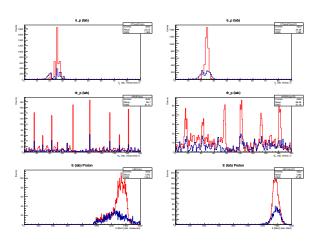


Figure 5: Red: MC; Blue Beamtime Data; Protons for $cos(\theta_{\omega}) = [-0.35, -0.25]$; Right Side are fitted data

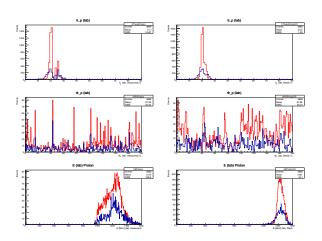


Figure 6: Red: MC; Blue Beamtime Data; Protons for $cos(\theta_{\omega}) = [-0.45, -0.35]$; Right Side are fitted data

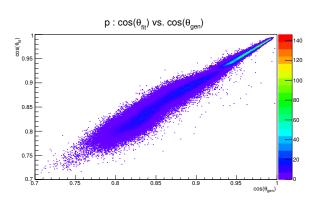


Figure 7: $cos(\theta_{fit})$ vs. $cos(\theta_{gen})$ for all protons.

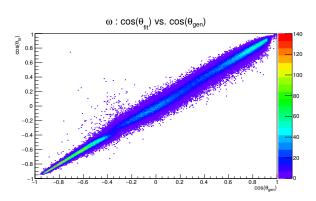


Figure 8: $\cos(\theta_{fit})$ vs. $\cos(\theta_{gen})$ for all ω .

Unfolding

Motivation

- \bullet μ is the *true* distribution given by nature
- detector effects are then described by the response function R. (inefficiencies, bias and smearing)
- This results in the distribution ν .

$$\nu_i = \sum_{j=1}^M R_{ij} \mu_j$$

 With infinite statistics, it would be possible to recover the original distribution by inverting the response matrix

$$\mu = R^{-1}\nu$$

What is Unfolding?



- Using MC we can train the unfolding algorithm
- Create a 2D-Hist with $\cos(\theta_{\omega})$ of all generated and all reconstructed ω (ω which are generated but not reconstructed are label miss)
- \bullet Then we can solve for μ iteratively

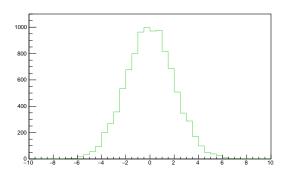


Figure 9: Example for a working Unfolding Algorithm

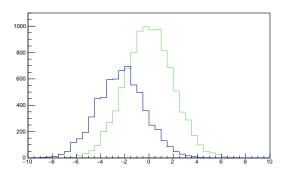


Figure 9: Example for a working Unfolding Algorithm

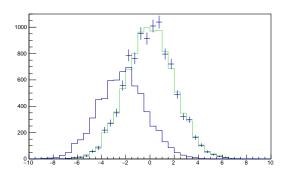


Figure 9: Example for a working Unfolding Algorithm

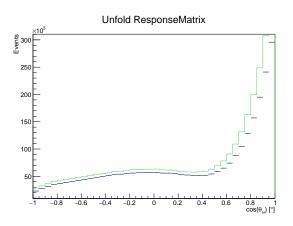


Figure 10: Folded; same cuts

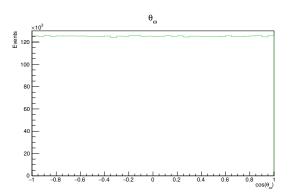


Figure 11: Distribution of the ω in center of mass frame



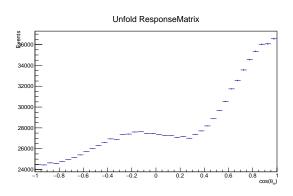


Figure 12: Flat ω was used. MC fitted data were folded.