

# Run 15 TSSA of Open Heavy Flavor Electrons at Midrapidity (5)

Dillon Fitzgerald  
Advisor: Christine Aidala  
07/01/2020



UNIVERSITY OF  
MICHIGAN



# Reminder

## Last update -- 04/29/2020

- Progress on open heavy flavor electron transverse single spin asymmetry was presented for the inclusive electron spectrum (**before background asymmetry/fraction corrections**)
  - $A_N(p_T)$  - square root formula (yellow, blue)
  - $A_N(p_T)$  - relative luminosity formula (yellow (left, right), blue(left, right))
  - Comparisons between results and extraction of systematic error from difference of sqrt and lumi formulas

## For This Talk:

- Background fraction calculation -- procedure and results
  - Closely follows procedure outlined in an1340 corresponding to PPG223



# Reminder

- Currently have  $A_N^{S+B}$  for different formulas and independent data samples
  - [ sqrt(yellow, blue), lumi(yellow left, yellow right, blue left, blue right) ]
- Need to calculate background fractions and calculate/aggregate background asymmetries such that  $A_N^S$  can be extracted as shown here
  - Here  $A_N^S = A_N^{\text{OHF} \rightarrow e}$

$$A_N^S = \frac{A_N^{S+B} - r A_N^B}{1 - r}$$

$$\sigma_{A_N^S} = \frac{\sqrt{(\sigma_{A_N^{S+B}})^2 + r^2 (\sigma_{A_N^B})^2}}{1 - r}$$

**More generally:**

$$r \rightarrow \sum_i f_i^{\text{norm}}$$

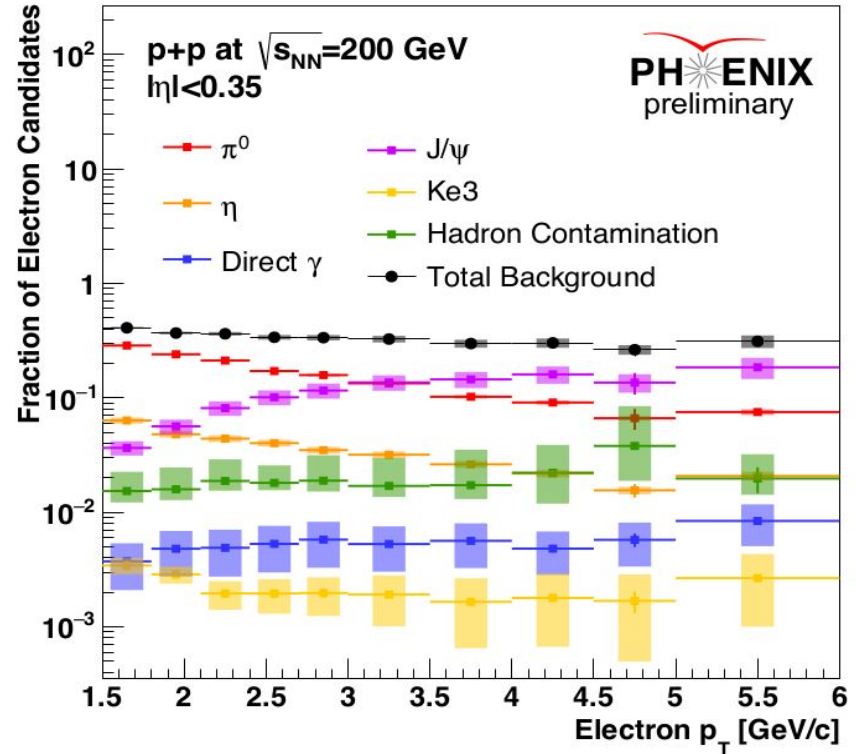
and

$$r A_N^B \rightarrow \sum_i f_i^{\text{norm}} A_N^i$$



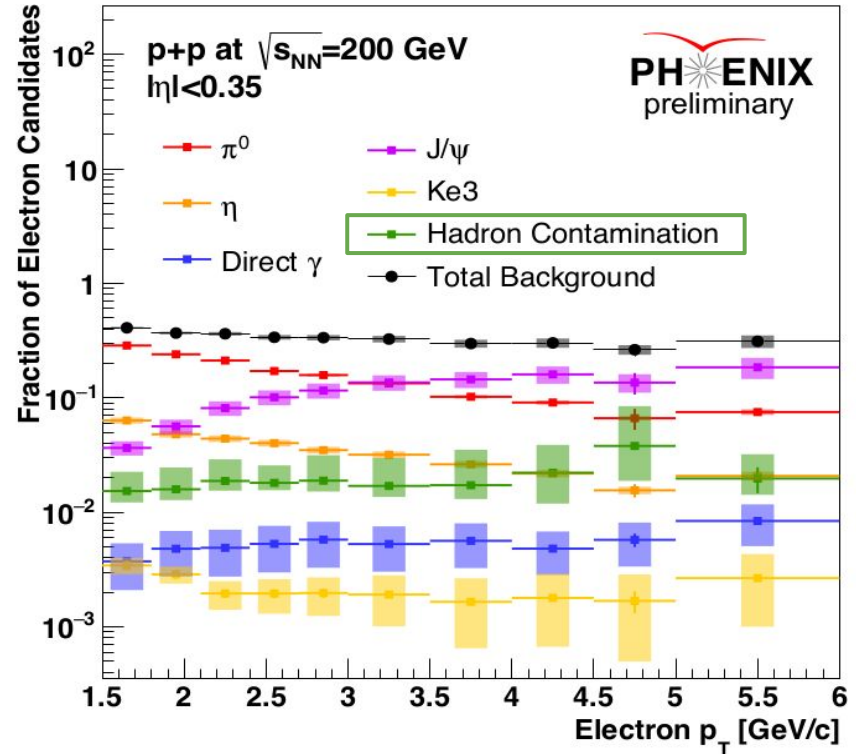
# Background Fractions from PPG223

- Background sources for this analysis and PPG223 are shown here
- Need to recalculate background fractions using my binning and curated electron candidate sample
- Consider the following backgrounds:



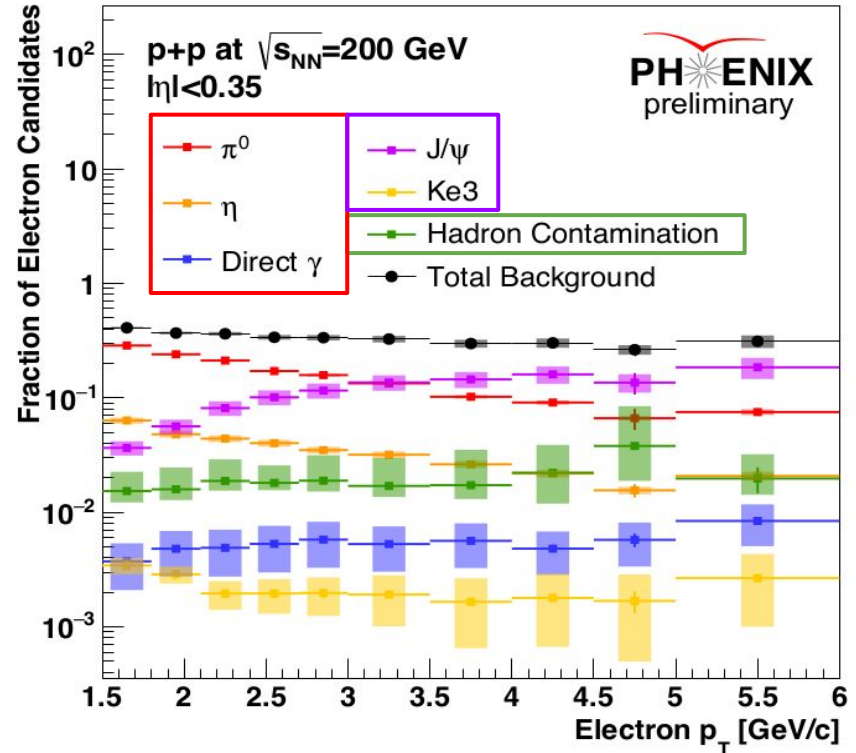
# Background Fractions from PPG223

- Background sources for this analysis and PPG223 are shown here
- Need to recalculate background fractions using my binning and curated electron candidate sample
- Consider the following backgrounds:
  - Hadron contamination (misidentified hadrons)



# Background Fractions from PPG223

- Background sources for this analysis and PPG223 are shown here
- Need to recalculate background fractions using my binning and curated electron candidate sample
- Consider the following backgrounds:
  - **Hadron contamination** (misidentified hadrons)
  - **Photonic** and **non-photonic** electrons  
-- create an electron cocktail from simulation



# Highest pT Bin -- Back to my binning...

Variable	Cut	Kinematic Range
emcdφ	< 3σ	1.5 GeV < p <sub>T</sub> < 6.0 GeV
emcdz	< 3σ	1.5 GeV < p <sub>T</sub> < 6.0 GeV
(E/p-μ) = dep * σ	< 2σ	1.5 GeV < p <sub>T</sub> < 6.0 GeV
zed	< 75	1.5 GeV < p <sub>T</sub> < 6.0 GeV
disp	< 5	1.5 GeV < p <sub>T</sub> < 6.0 GeV
χ <sup>2</sup> /ndf	< 3	1.5 GeV < p <sub>T</sub> < 6.0 GeV
nhits	> 2	1.5 GeV < p <sub>T</sub> < 6.0 GeV
quality	== (63 or 31)	1.5 GeV < p <sub>T</sub> < 6.0 GeV
conversion veto	✓	1.5 GeV < p <sub>T</sub> < 6.0 GeV
n0	> 1	1.5 GeV < p <sub>T</sub> < 5.0 GeV
prob	> 0.01	1.5 GeV < p <sub>T</sub> < 5.0 GeV
n0	> 3	5.0 GeV < p <sub>T</sub> < 6.0 GeV
prob	> 0.2	5.0 GeV < p <sub>T</sub> < 6.0 GeV

Bin #	1	2	3	4
Pt Range (GeV)	(1.5, 1.8)	(1.8, 2.1)	(2.1, 2.7)	(2.7, 6.0)

- Differing n0 and prob cuts above 5 GeV implemented to suppress hadron contamination to the few percent level in the 5-6 GeV bin of PPG223
- Spin sorting decreases statistics, 5-6 GeV bin is not viable... So currently have differing cuts within a single bin (2.7 - 6 GeV)
  - Proposal to limit range to 5 GeV in pT?  $n_{2.7-5}/n_{2.7-6} \sim 98\%$



# Pt Bins

The following Pt bins are used for this analysis -- chosen by combining bins used in PPG223:

Bin #	1	2	3	4
Pt Range (GeV)	(1.5, 1.8)	(1.8, 2.1)	(2.1, 2.7)	(2.7, 5.0)





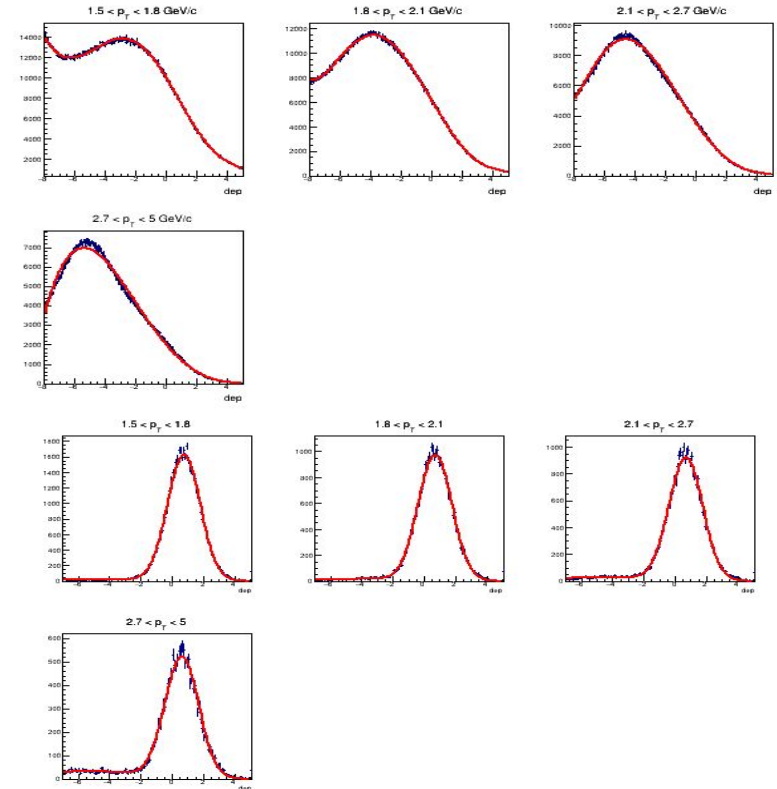
# Hadron Contamination

- Two methods used to calculate fraction of candidates attributed to charged hadrons misidentified as electrons
  - Dep fitting method
  - Algebraic calculation with  $n_0$  survival rate
- Weighted average of both methods is used to calculate hadron contamination fraction
  - Difference between average value and values from either method are considered for systematics



# Hadron Contamination -- dep Fitting

- True electrons have Gaussian dep distribution with  $\sigma \sim 1$  and  $\mu \sim 0$
- Hadron dep distribution has different characteristics
  - Accessed in data via  $n_0 < 0$  cut
  - Fit extracted for template in fitting the electron candidate distribution (normalization is the only free parameter)
- Electron candidate dep distribution fit with Gaussian + hadron dep template
  - Allows for extraction of both electron and hadron contribution
  - Hadron contamination fraction determined by ratio of the contributions in the dep cut region  $|\text{dep}| < 2$

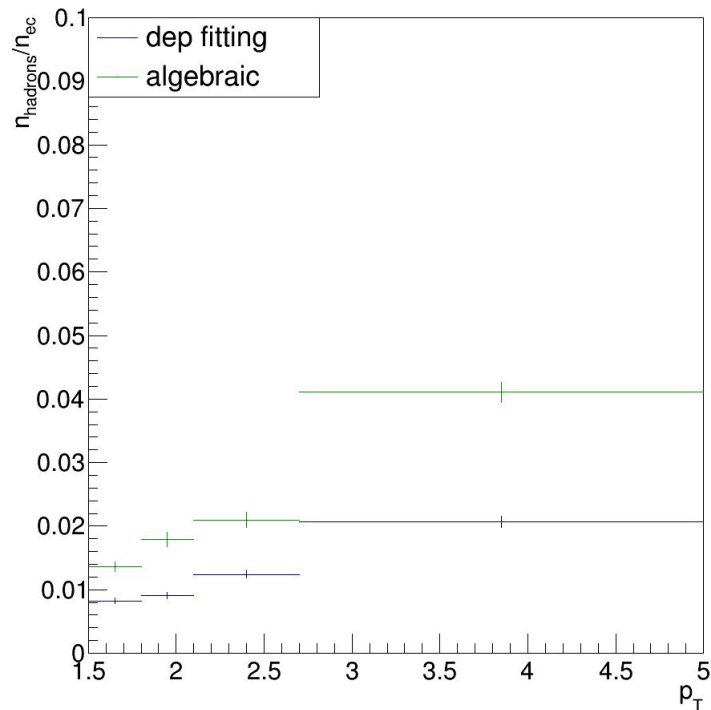


# Hadron Contamination -- Algebraic Method

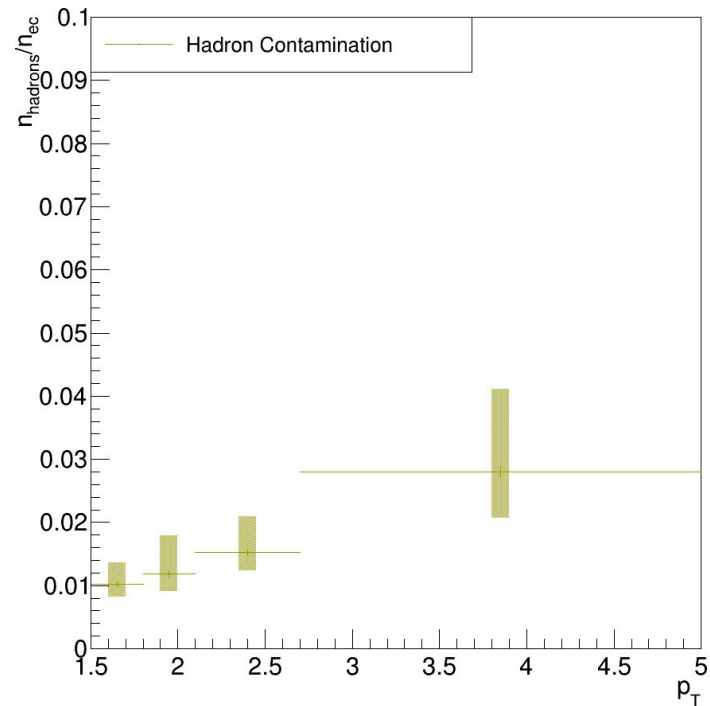
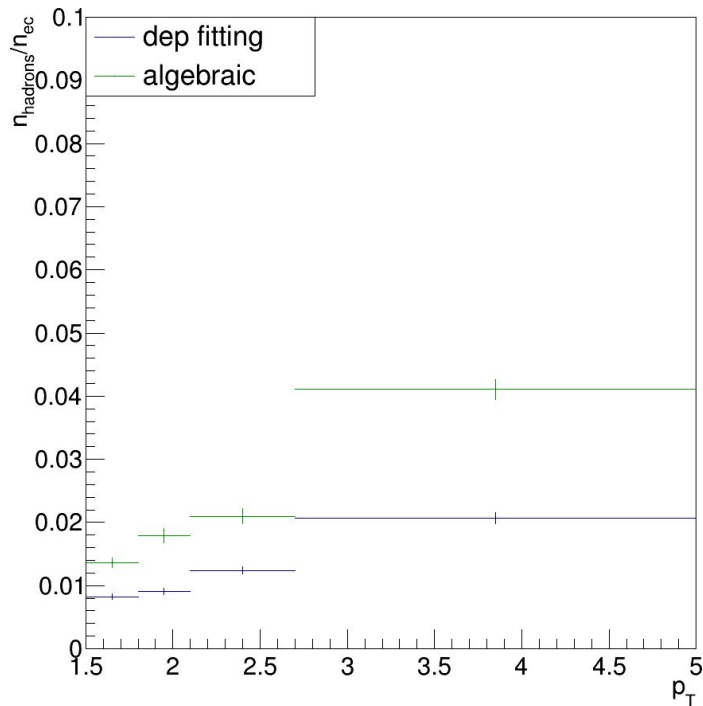
- Provides estimate that is independent of dep shape by using the survival rate of the n0 cut (accessed via electron candidate sample with dep<-6)
  - Electrons fire the RICH differently than charged hadrons, and therefore have a different survival rate
- The following system of equations can then be used to isolate the number of hadrons in our electron candidate sample:

$$n_{\text{non0}} = n_e + n_h \quad n_{h,n0} = \epsilon_h (n_{n0} - \epsilon_e n_{\text{non0}}) / (\epsilon_h - \epsilon_e)$$

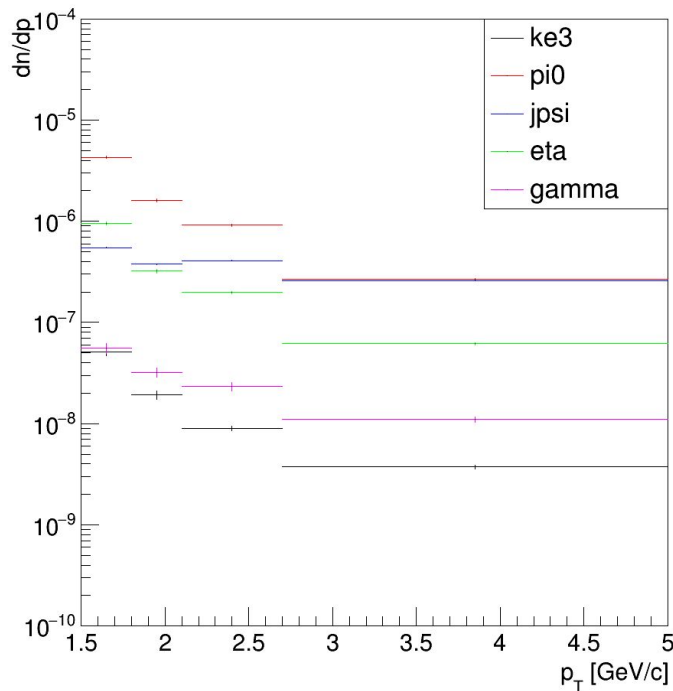
$$n_{n0} = \epsilon_e n_e + \epsilon_h n_h$$



# Hadron Contamination



# Electron Cocktail Simulation



- Electron cocktail created from single particle electron simulations
  - Simulation details outlined extensively in an1340
- $p_T$  weighting based on measured PHENIX cross sections
  - Still needs to be normalized w.r.t. Inclusive electron spectrum!
  - We have a handle on the photonic vs. non-photonic fraction of electrons in our sample via the survival rate of the conversion veto cut, and can use this fact for normalization

# Background Fraction Calculation

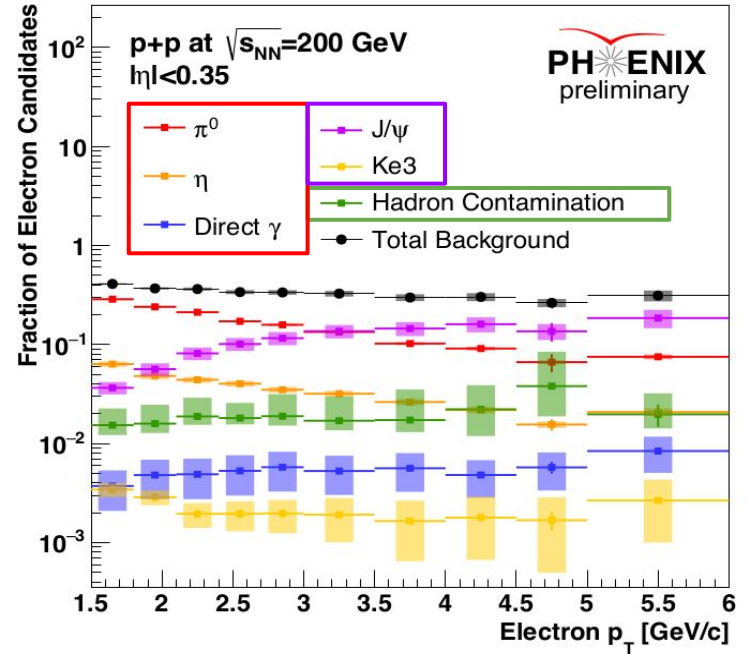
$$n_e = n_{np} + n_p + n_{hc}$$

$$\tilde{n}_e = \tilde{n}_{np} + \tilde{n}_p + \tilde{n}_{hc} = \epsilon_{uc} n_{np} + \epsilon_{uc} \epsilon_p n_p + \tilde{n}_{hc}$$

$$F_{np} = \frac{n_{np}}{n_{np} + n_p}$$

**no ~ = without conversion veto**  
**~ = with conversion veto**

$$F_{np} = \frac{\epsilon_{uc} \epsilon_p n_e - \tilde{n}_e - \epsilon_{uc} \epsilon_p n_{hc} + \tilde{n}_{hc}}{(\epsilon_p - 1)(\tilde{n}_e - \tilde{n}_{hc})}$$

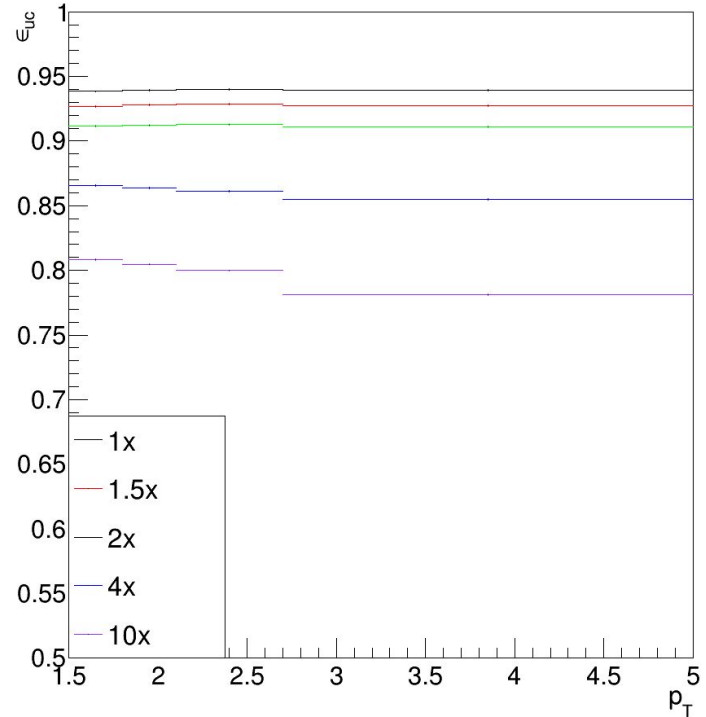
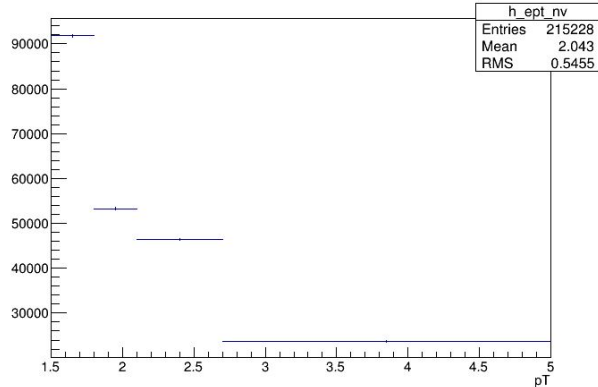


$$f_i^{\text{norm}} = (1 - \tilde{f}_{hc})(1 - F_{np}) \frac{\tilde{n}_i}{\tilde{n}_{\pi^0} + \tilde{n}_{\eta} + \tilde{n}_{\gamma}}$$

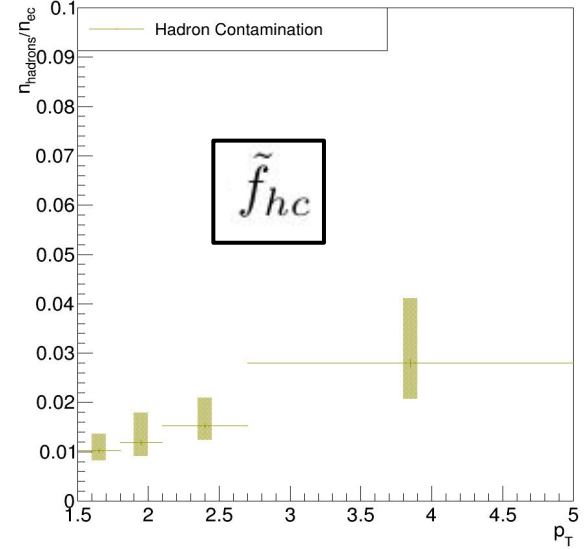
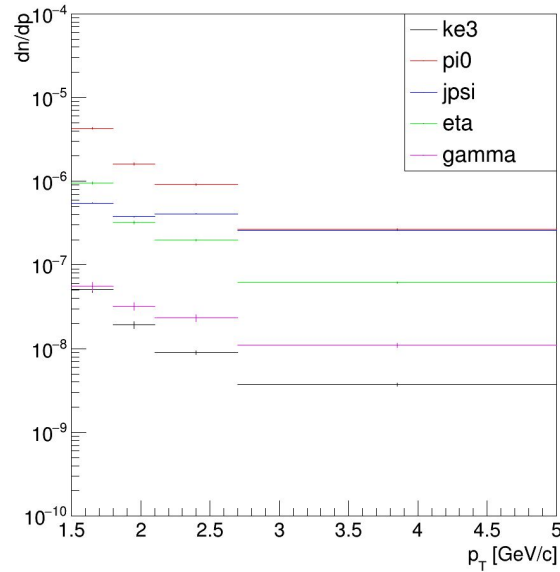
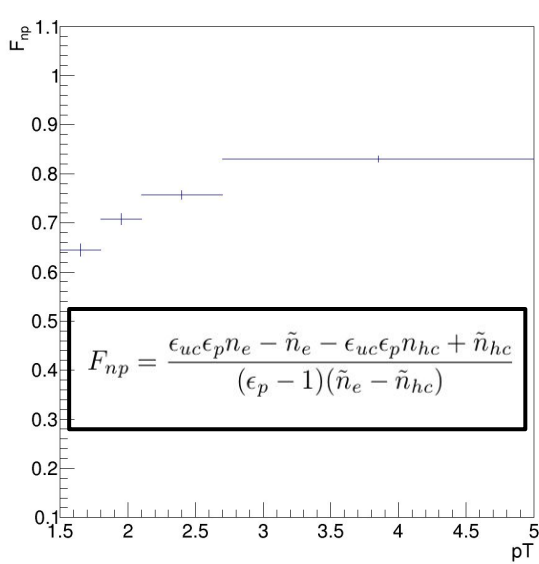


# Conversion Veto Survival Rates

- 2x conversion veto window size (green) was used in PPG223
- $\epsilon_{uc}$  extracted from survival rate histo (hadrons in data)
- $\epsilon_p$  extracted from simulation (not shown here)
- $n_e$  and  $\tilde{n}_e$  extracted from  $e p_T$  spectrum with and without conversion veto cut



# Background Fraction Calculation



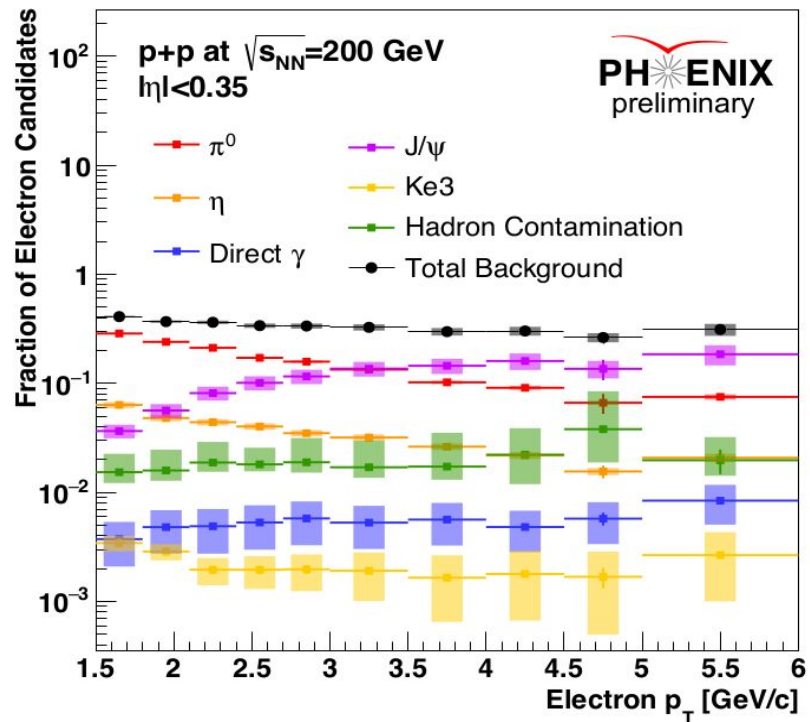
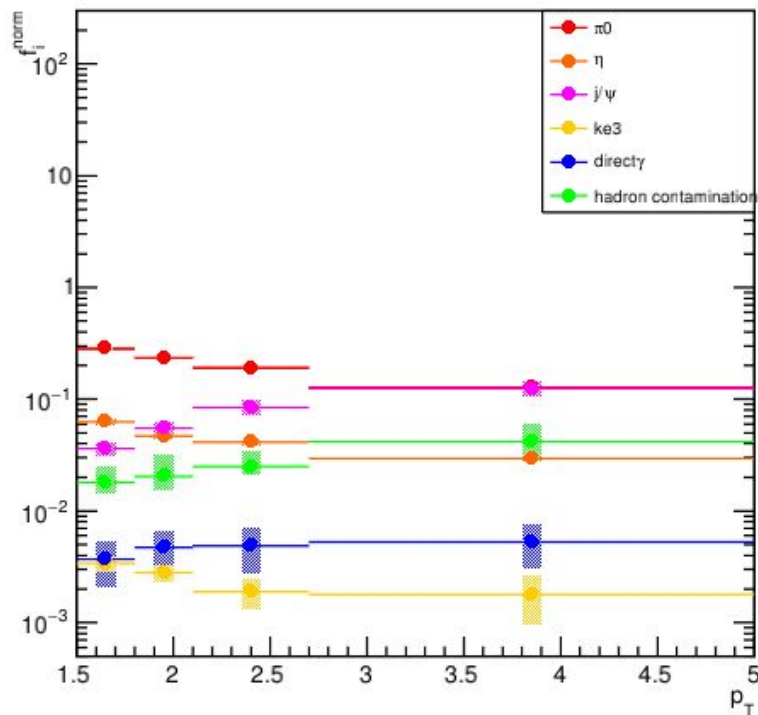
$$f_i^{\text{norm}} = (1 - \tilde{f}_{hc})(1 - F_{np}) \frac{\tilde{n}_i}{\tilde{n}_{\pi^0} + \tilde{n}_{\eta} + \tilde{n}_{\gamma}}$$

$$f_i^{\text{norm}} = f_{\pi^0}^{\text{norm}} \frac{\tilde{n}_i}{\tilde{n}_{\pi^0}}$$





# Background Fraction Calculation



# Next Steps

- Bunch shuffling cross check
- $A_N \sin\phi$  modulation cross check
- Background Correction
- Gather systematics



# Next Steps

- Bunch shuffling cross check
- $A_N \sin\phi$  modulation cross check
- Background Correction
- Gather systematics

Does the order in which these are completed matter?

# For Your Information

I have been tracking my analysis progress on the web:

[http://www-personal.umich.edu/~dillfitz/PHENIX\\_Analysis/index.html](http://www-personal.umich.edu/~dillfitz/PHENIX_Analysis/index.html)

$A_N(p_T)$  plots can be found at (see various subdirectories for formula comparisons):

[http://www-personal.umich.edu/~dillfitz/PHENIX\\_Analysis/Asymmetry\\_Ana/1.5GeV\\_6GeV/pTBins/](http://www-personal.umich.edu/~dillfitz/PHENIX_Analysis/Asymmetry_Ana/1.5GeV_6GeV/pTBins/)

For electron ID selection requirements and motivating histograms, see:

[http://www-personal.umich.edu/~dillfitz/PHENIX\\_Analysis/Open\\_Heavy\\_Flavor\\_Ana/electronID\\_and\\_spectra/lin\\_y/](http://www-personal.umich.edu/~dillfitz/PHENIX_Analysis/Open_Heavy_Flavor_Ana/electronID_and_spectra/lin_y/)

