# $\Lambda_b \to J/\Psi p \, K$ Production Correction Run2 {'16, '17, '18}

M. Ferrillo UZH

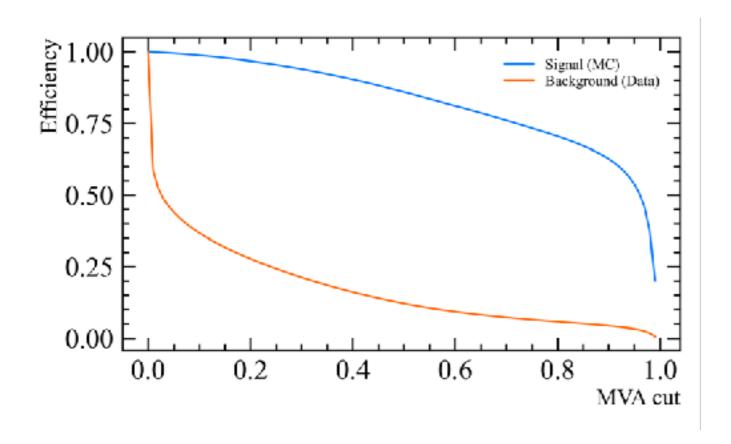
#### NOVELTY

- ♦ Enforced a cut on mu\_P, mu\_PT > 2 GeV
- ♦ Repeat the analysis chain for 2016, 2017, 2018
  - ◇ Run PIDCorr to retrieve PID vars correlations ({p, K}ProbNN, DLL)
  - ♦ Apply preselection
  - ♦ Build MC/Data samples for MVA to tackle MisID/comb bkg
  - ♦ Optimise hyperparams MVA
  - ♦ Train MVA. Post training checks on Lb\_M correlation with MVA score
  - ♦ Apply MVA cut and design a MisID selection (cut on reflection backgrounds)
  - ♦ Pre-fit MC Lb\_M to extract the shape parameters of signal (extended unbinned ML fit with Double Crystal Ball)
  - ♦ Extended unbinned ML fit data Lb M with DCB (sig) + exp (bkg)
  - Extract sWeights for signal Data
  - ♦ Calculate the Data/MC correction in bins of Lb\_P, Lb\_PT. Check pre-post correction distributions

## MVA TRAINING SET VARIABLES

# Taken from the Pentaquark analysis [PRL122(2019)222001]

- $K_{\rm PT} + p_{\rm PT}$
- $\ln p_{\rm P}$
- $\ln IPCHI2_{\Lambda_h}$
- $\Lambda_{b\,\mathrm{PT}}$
- $\ln \text{FD}_{\Lambda_b}$
- $\ln \text{CHI2OWNPV}_{\Lambda_b}$
- $ln(1 DIRA_{\Lambda_b})$
- ♦ ln MINIPCHI2<sub>h</sub>
- $\min DLL(\mu \pi)$  PIDCorr sampled
- $p_{\text{probNN}p}$  PIDCorr sampled
- $K_{\text{probNN}K}$  PIDCorr sampled

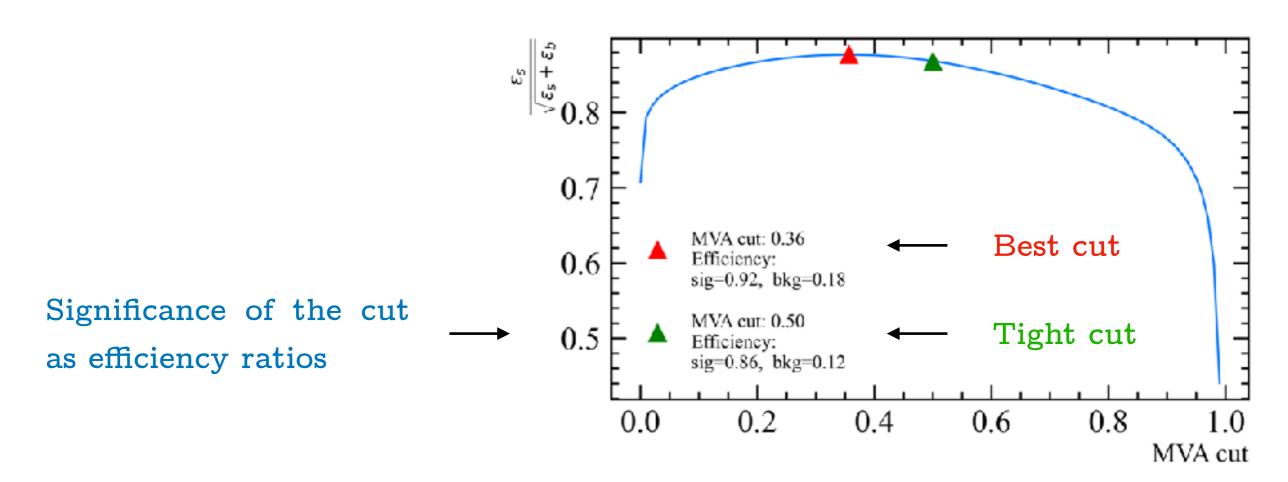


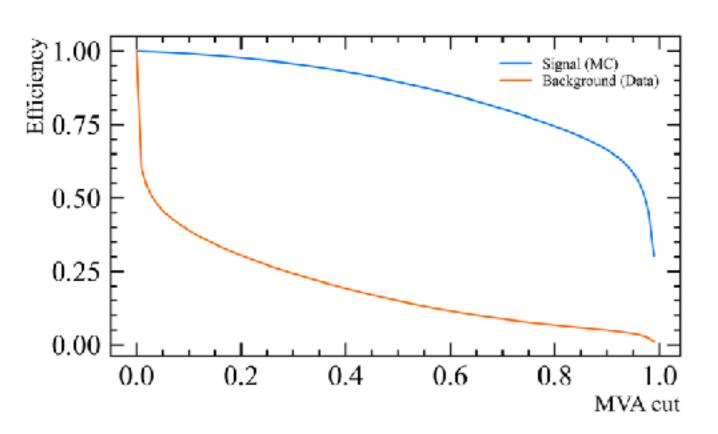
I have had a quick look at the efficiency curves. No fitting to extract the yields.

Signal: peak region

Background: lower and upper

sidebands



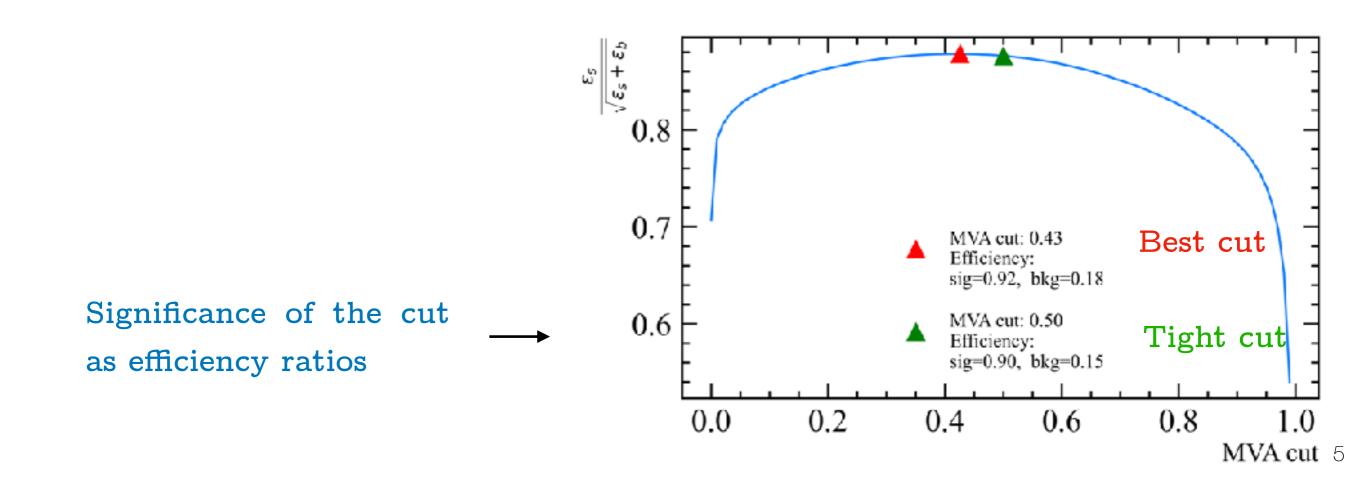


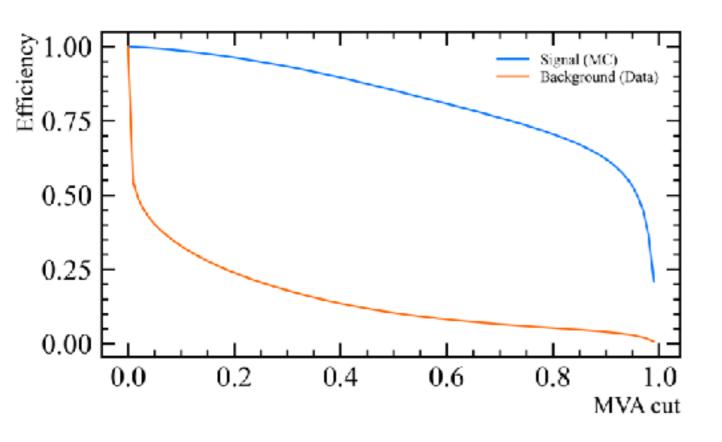
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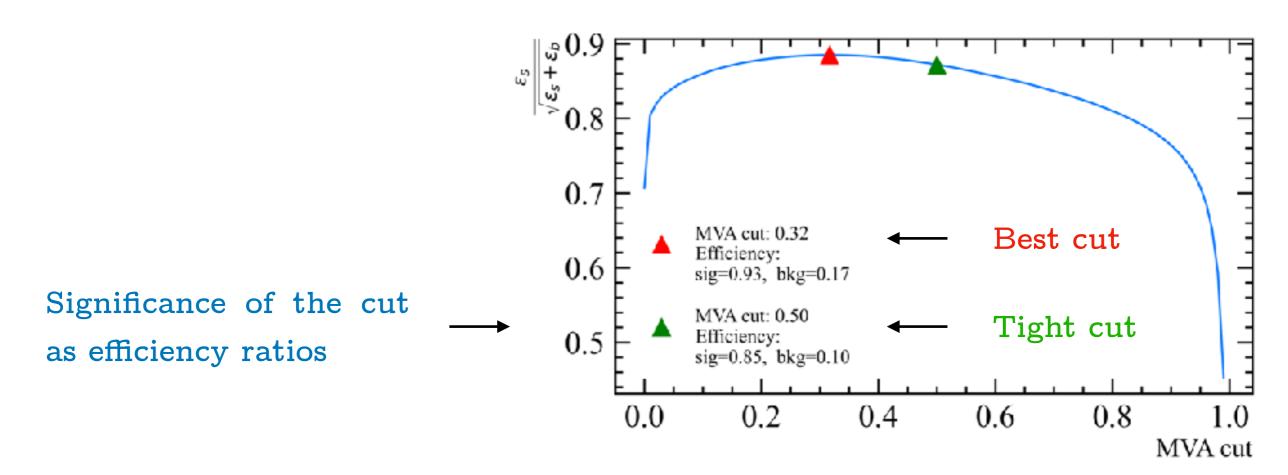


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Signal: peak region

Background: lower and upper

sidebands



## MISID COMPONENTS

- $\diamond$  The three main contributions to MisID of  $\Lambda_b^0 \to J/\Psi p K^$ 
  - a)  $\bar{B}_s^0 \to J/\Psi(\Phi \to K^+K^-)$
  - **b)**  $B^0 \to J/\Psi \pi^+ K^-$
  - c)  $\bar{\Lambda}_b^0 \to J/\Psi \bar{p} K^+$

I have checked the following:

- The reconstructed mass distribution in the replaced mass hypothesis for each MisID channel, after the MVA selection
  - $\diamond$  Case I: MVA score > best cut
  - ♦ Case II: MVA score > 0.50 (tighter cut)
- $\diamond$  The reconstructed  $\Lambda_b$  mass after a  $1\,\sigma, 2\,\sigma, 3\,\sigma$  Veto cut on the MisID reconstructed mass for each contribution ( $\sigma = 20~\text{MeV}$ ):
  - ♦ Singularly (cut on a, b or c separately)
  - ♦ Combined cut

#### MISID COMPONENTS

 $\diamond$  The three main contributions to MisID of  $\Lambda_b^0 \to J/\Psi p \, K^-$ 

a) 
$$\bar{B}_s^0 \to J/\Psi(\Phi \to K^+K^-)$$

**b)** 
$$B^0 \to J/\Psi \pi^+ K^-$$

c) 
$$\bar{\Lambda}_b^0 \to J/\Psi \bar{p} K^+$$

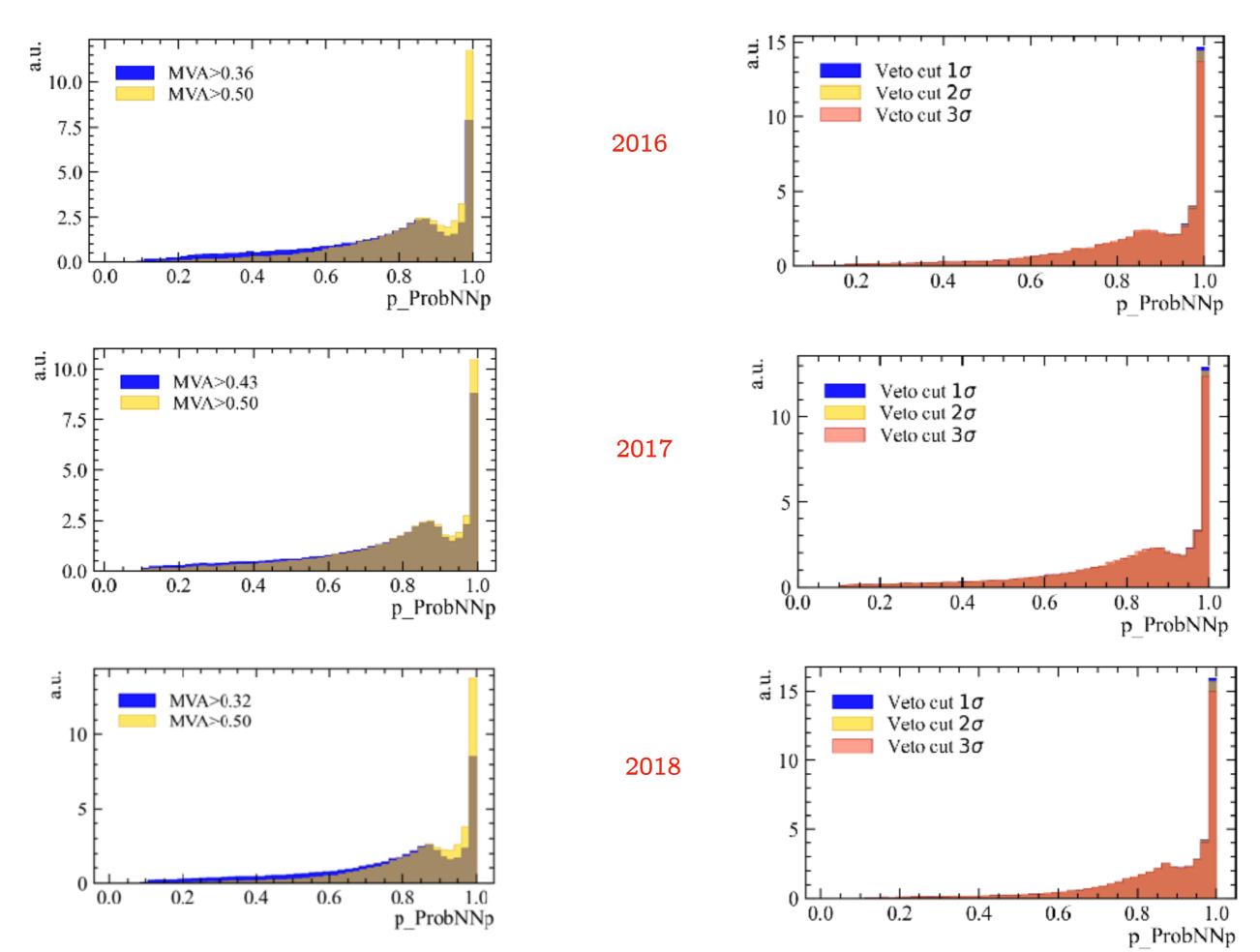
I have checked the following:

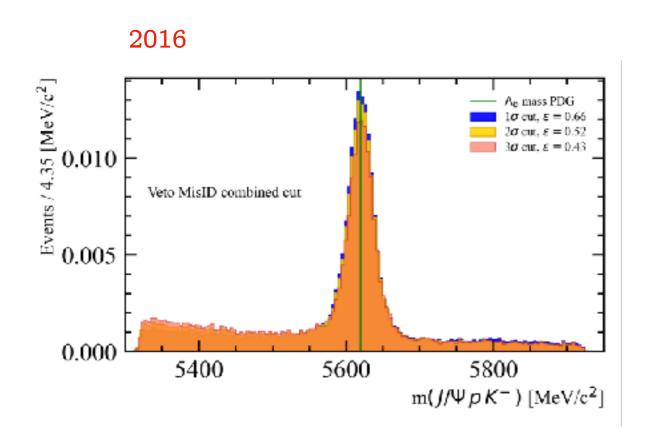
- The reconstructed mass distribution in the replaced mass hypothesis for each
  MisID channel, after the MVA selection
  - $\diamond$  Case I: MVA score > best cut
  - $\diamond$  <u>Case II</u>: MVA score > 0.50 (tighter cut)

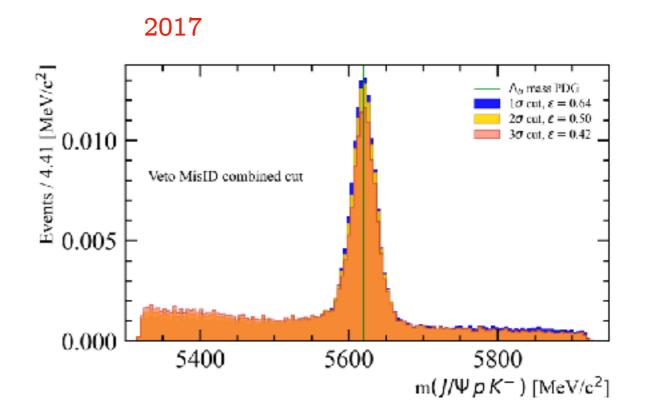
why: probNN distributions show less contamination

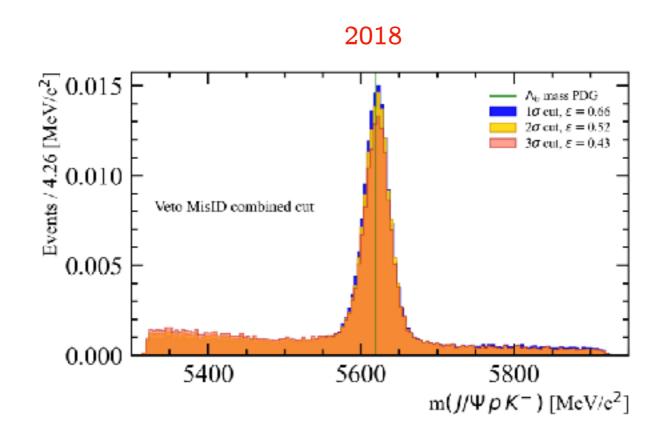
- $\diamond$  The reconstructed  $\Lambda_b$  mass after a  $1\sigma, 2\sigma, 3\sigma$  Veto cut on the MisID reconstructed mass for each contribution ( $\sigma = 20 \text{ MeV}$ ):
  - ♦ Singularly (cut on a, b or c separately)
  - ♦ Combined cut

# MISID COMPONENTS: PROBNN

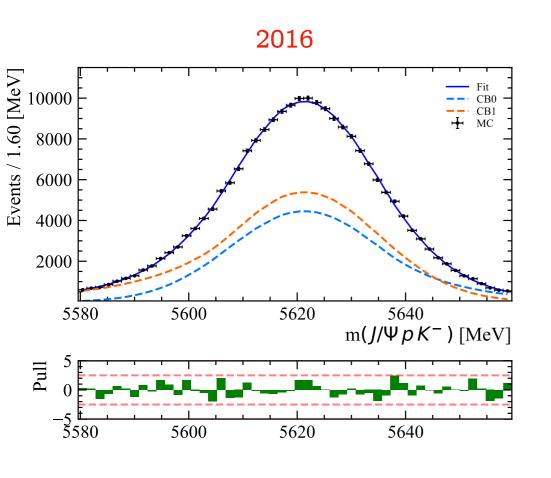


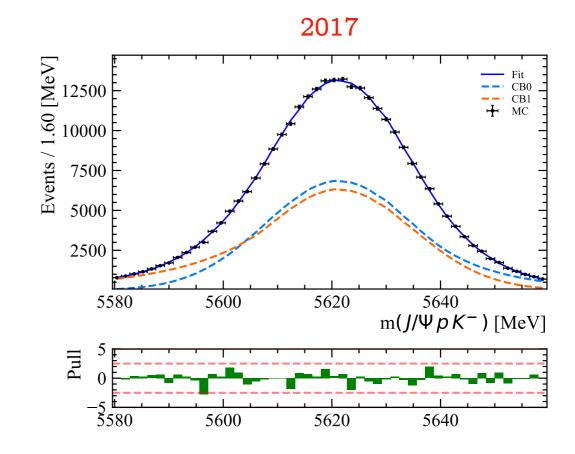


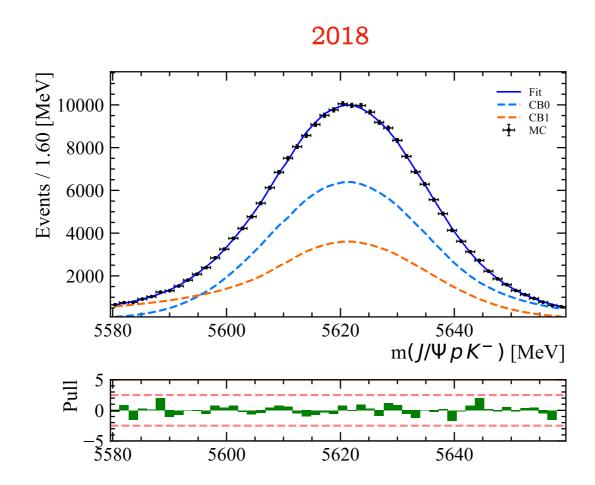




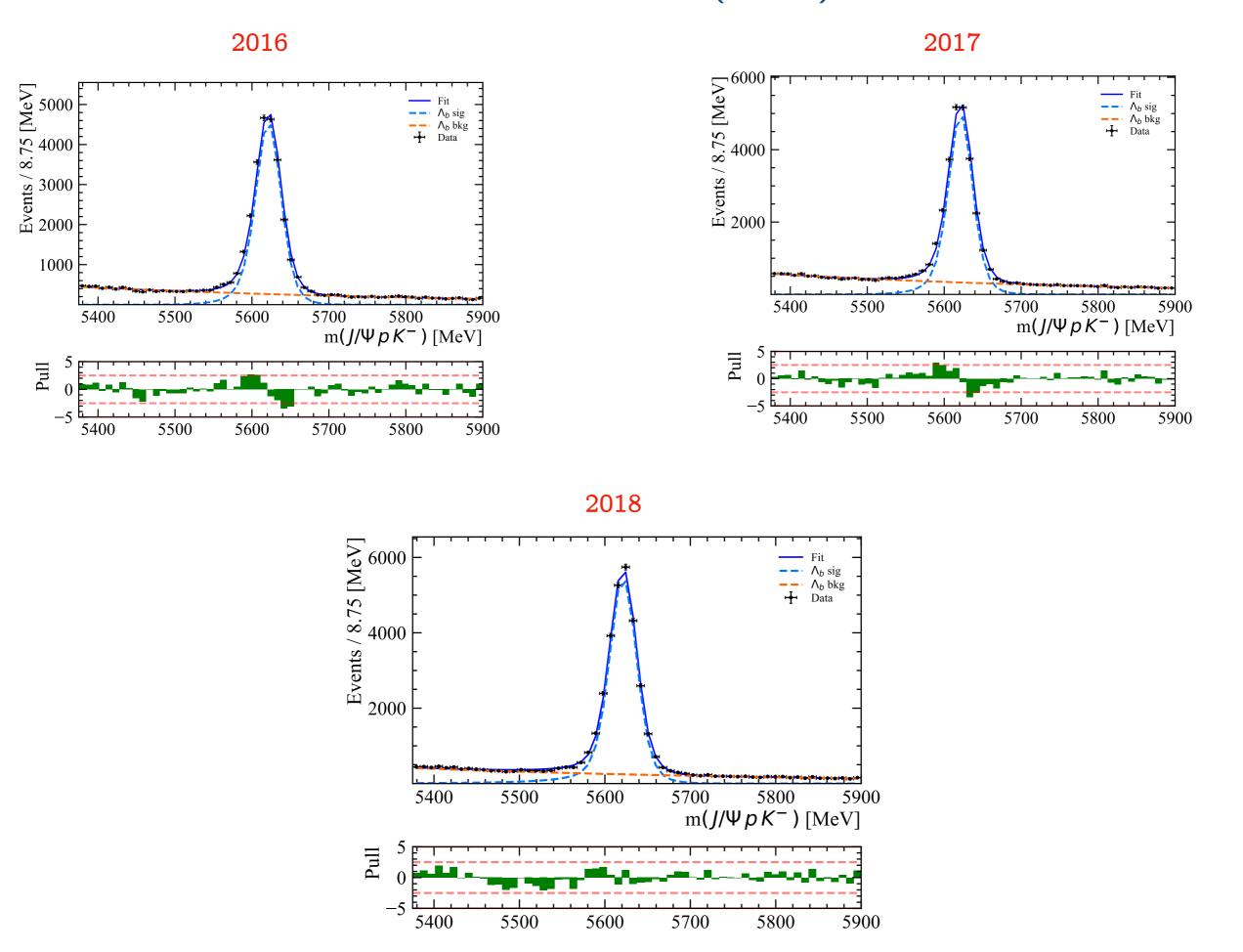
# LBM FIT (MC)







# LBM FIT (DATA)



# EVALUATE THE CORRECTION: STRATEGY

# • Option 1:

- Define a regular-sized binning scheme (Nbins = 15\*15 = 225);
- Retrieve the corrections as the ratio of the normalised Data/MC 2D (Lb P, Lb PT) distributions;
- Save to ROOT file;
- Get the MC reweighed P and PT distributions and compare them to data.

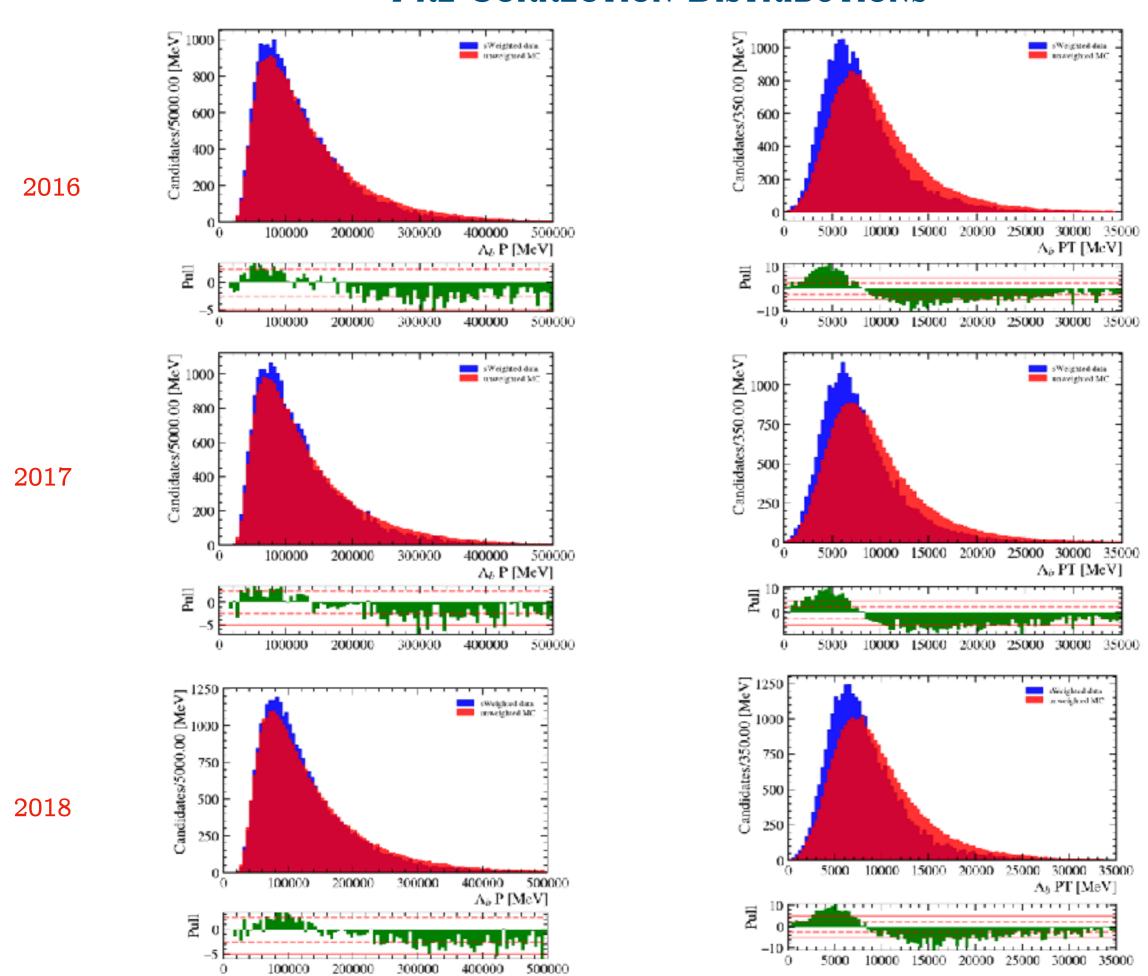
# • Option 2:

- Define a rectangular binning scheme (Nbins = 15\*30 = 450);
- Retrieve the corrections as the ratio of the normalised Data/MC 2D (Lb\_P, Lb\_PT) distributions;
- Save to ROOT file;
- Get the MC reweighed P and PT distributions and compare them to data.

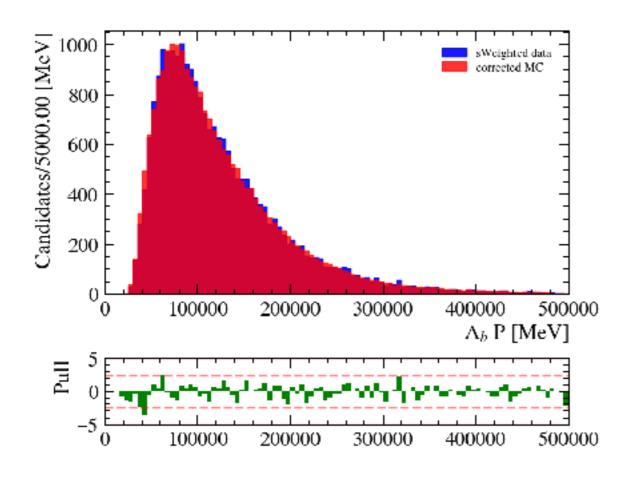
## STRATEGY

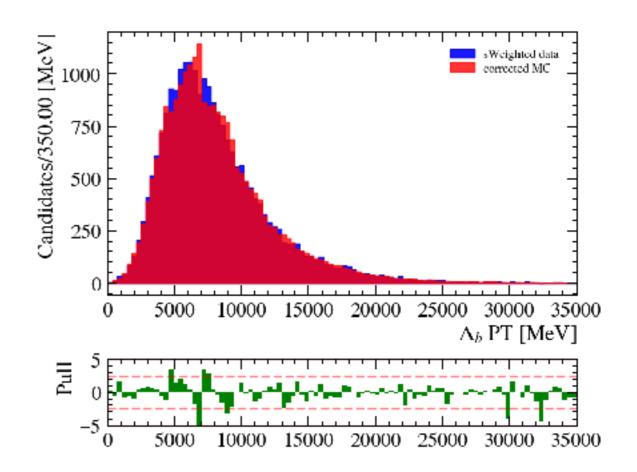
# • Option 3:

- Define a polygonal binning scheme by optimising for the number of MC entries per bin (Nentries =  $10^3 \Rightarrow \text{Nbins} \sim 250$ );
- Optimisation performed with a KDTreeBinning algorithm (ROOT);
- The binning structure of MC is adopted also for Data. TH2Poly histograms are used;
- Retrieve the corrections as the ratio of the normalised Data/MC 2D (Lb P, Lb PT) distributions;
- Save to ROOT file;
- Get the MC reweighed P and PT distributions and compare them to data.

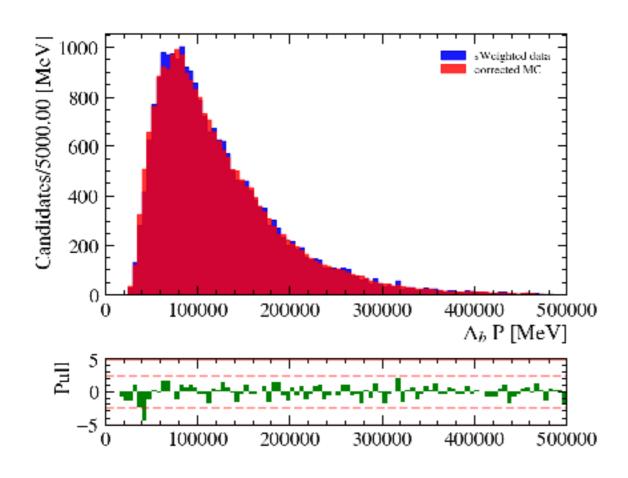


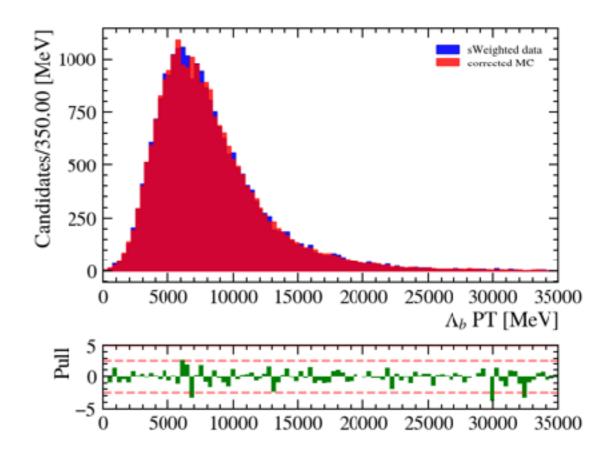
## REGULAR BINNING

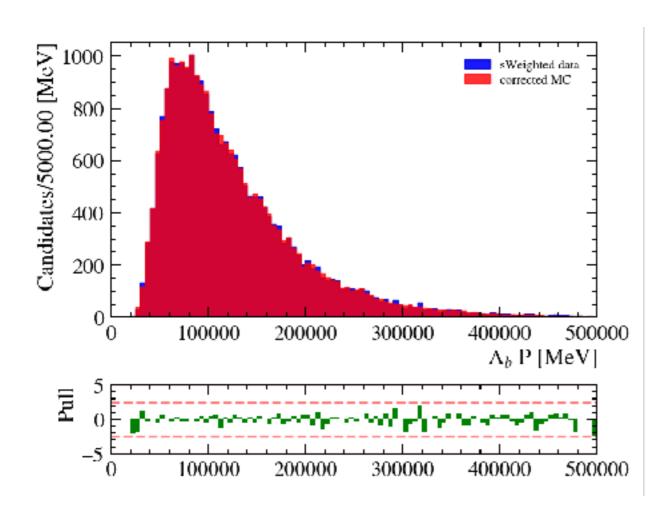


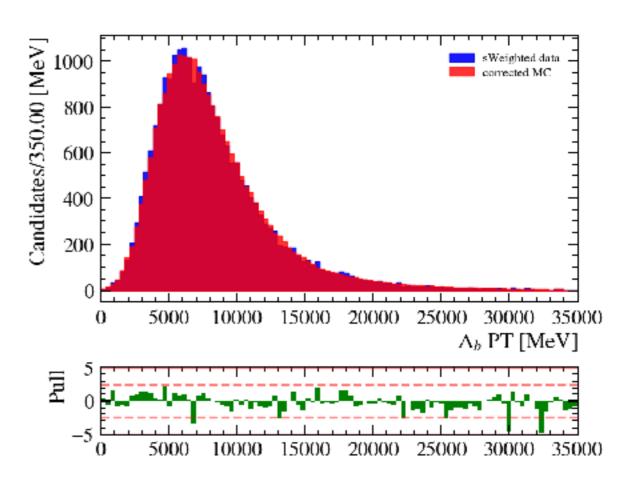


#### RECTANGULAR BINNING



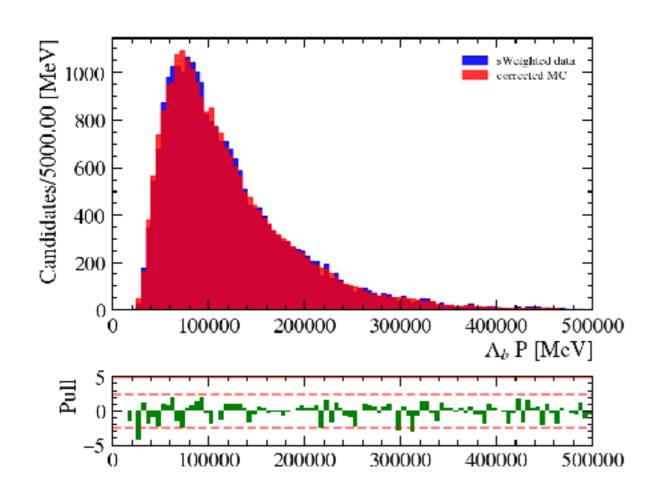


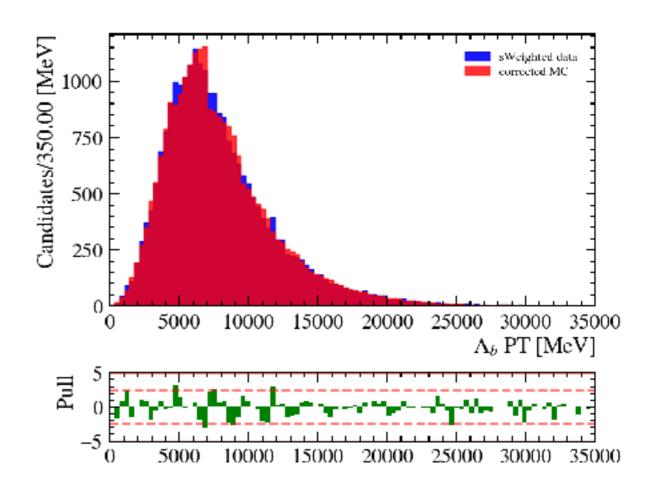




# OPTION 1: DATA/MC COMPARISON 2017

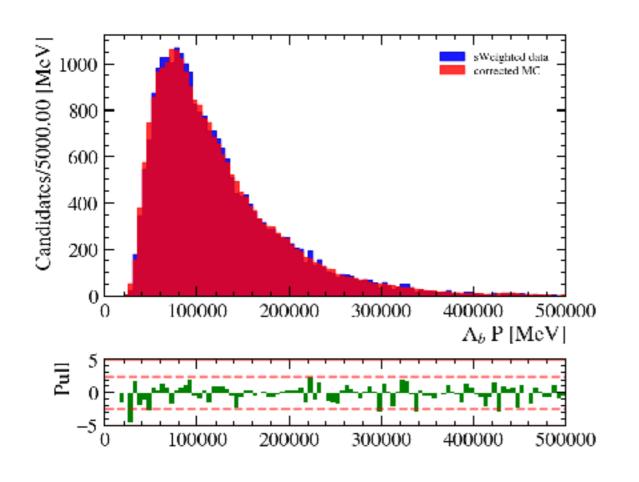
## REGULAR BINNING

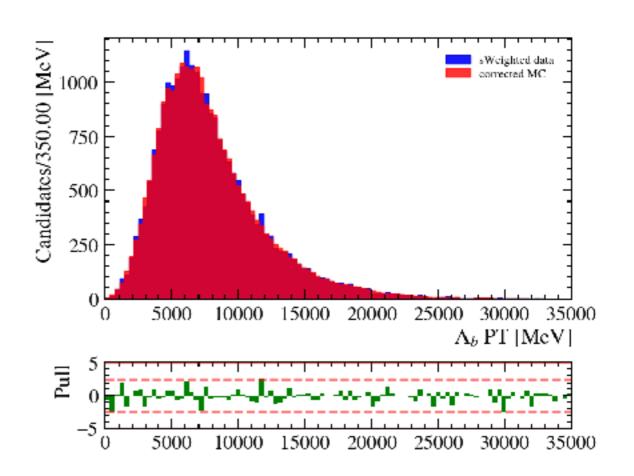




# OPTION 2: DATA/MC COMPARISON 2017

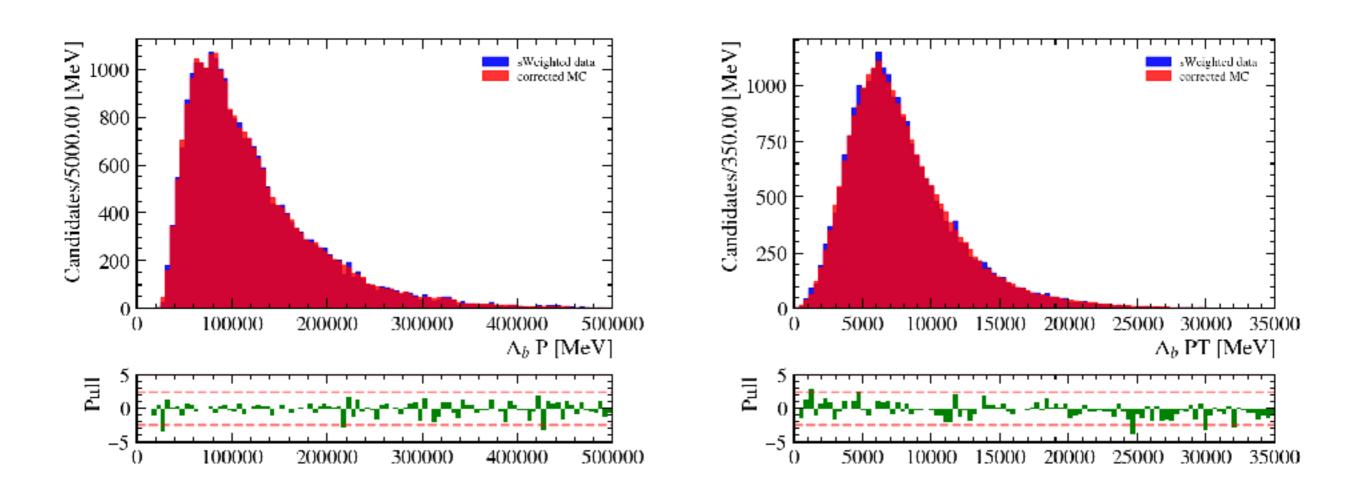
RECTANGULAR BINNING

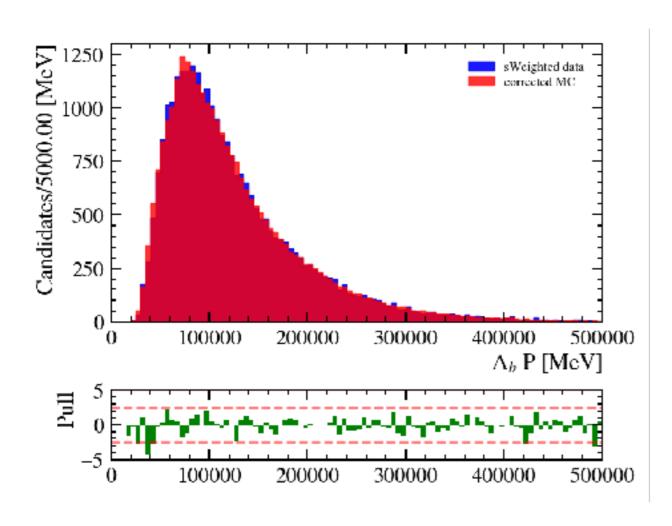


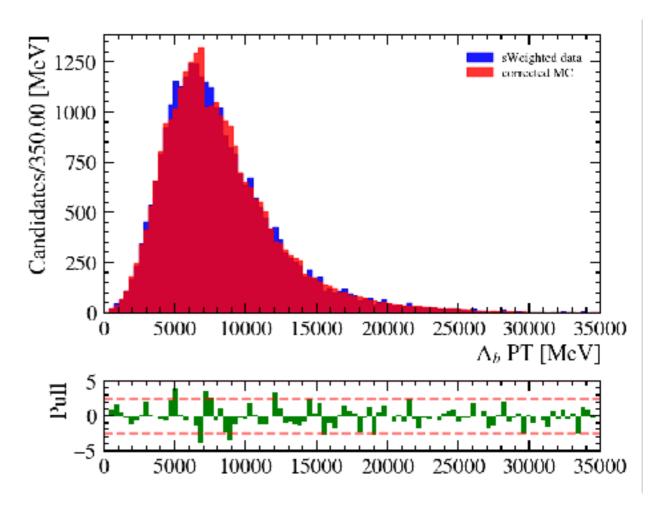


# OPTION 3: DATA/MC COMPARISON 2017

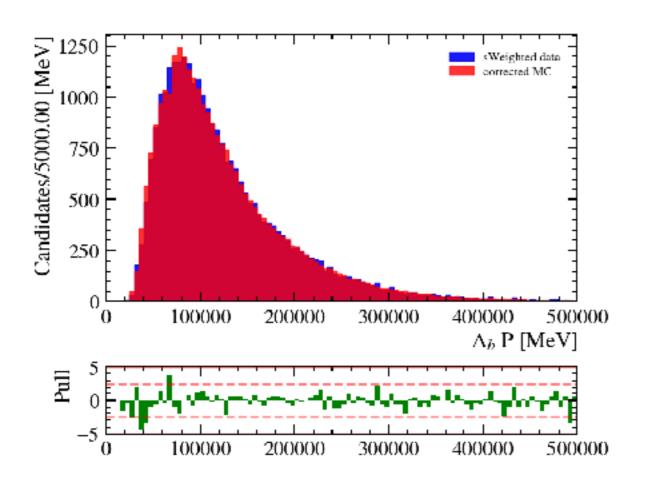
#### ADAPTIVE BINNING

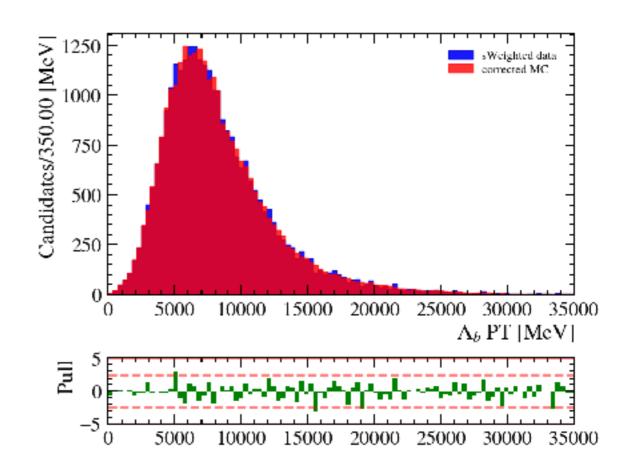






RECTANGULAR BINNING





#### ADAPTIVE BINNING

