



PURDUE
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High- p_T muon resolution measurement for 2022 and 2023

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Introduction and Method

- Measure 2022 and 2023 muon p_T resolution using spark_tnp.
 - Goal is to validate resolution modeling in MC
- Can't measure the resolution for single muons in collision data
 - -> Measure the dimuon resolution at the Z peak as function of leading muon p_T
- Muon selections: $p_T > 53$ Gev, $|\eta| < 2.4$, relative momentum uncertainty < 0.3 , high-pt ID, and one of the muons need to have fired the single muon trigger ([HLT_CascadeMu100](#) or [HLT_HighPtTkMu100](#) or [HLT_Mu50_v](#))
 - Pair selection: require tag muon $p_T >$ probe muon p_T , $\text{pair_tuneP_normalchi2} < 20$ and $\cos_{\text{angle}} < \pi - 0.02$

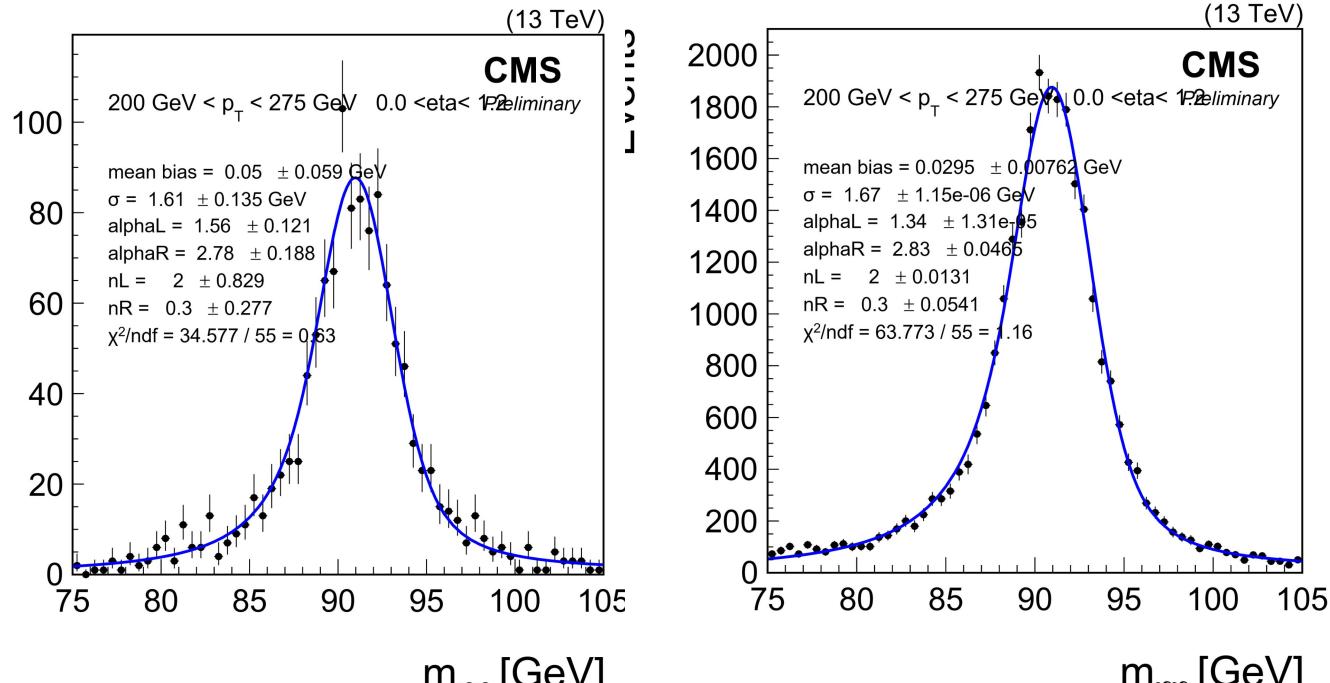
Introduction and Method

- Consider two categories: $\text{tag_abseta} < 1.2$ and $1.2 < \text{tag_abseta} < 2.4$
- Pt bins: [52, 72, 100, 152, 200, 275, 452, 800] GeV
- Data: 2022 , 2022_EE, 2023, and 2023_BPix data taking periods
- MC: Inclusive Boosted DY, binned in Z p_T

Procedures:

- 1, Create the dimuon(Z) mass histograms as a function of leading muon pT
- 2, Fit the Histograms with 3 different functions : Crystal Ball, Cruijff, and Double-sided Crystal Ball
 - We fit a convolution of the Breit-Wigner function with one of the three parametrizations of the resolution
 - Double-sided Crystal Ball is default, Cruijff and Crystal Ball used for systematic variations
- 3, Extract the resolution values from the fits and calculate the systematic uncertainties.
- 4, If necessary, **smear the MC resolution to meet the Data resolution.**
 - For simplicity, in the following plots of **comparison** between MC and Data, only the statistical error is shown.
 - The impact of systematic uncertainties on the Data/MC comparison is evaluated separately.
 - **Compared to the last update, we improved the fits and were able to improve the consistency of the results. A study of the systematic uncertainties was added**

Example fits

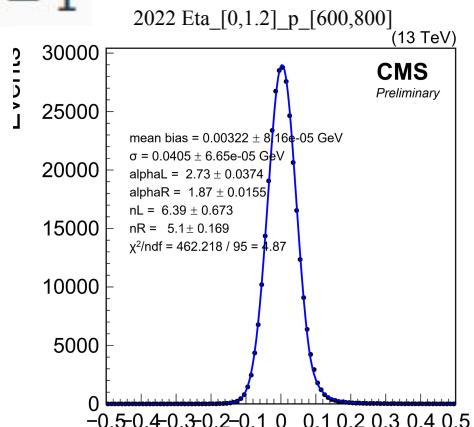
3
5
1

Data : 2022 Eta_[0,1.2] Pt_[200,275]

MC : 2022 Eta_[0,1.2] Pt_[200, 275]

Resolution Smearing

- If Data and MC disagree, we need to smear of the momenta in MC
- New resolution given by $\sigma_{\text{smeared}} = \sqrt{\sigma_{\text{default}}^2 + \sigma_{\text{extra}}^2}$,
where $\sigma_{\text{default}}(p)$ is the resolution of the residual of MC : $\frac{1/p - 1/p^{\text{gen}}}{1/p^{\text{gen}}}$
- Written in terms of desired smearing: $x = \sqrt{\text{smearfac}^2 - 1}$
 $\text{smearfac} \cdot \sigma_{\text{default}} = \sqrt{\sigma_{\text{default}}^2 + (x \cdot \sigma_{\text{default}})^2}$
- For 10% smearing : smearfac= 1.1 → x=0.46
- For 5% smearing : smearfac=1.05→ x=0.32

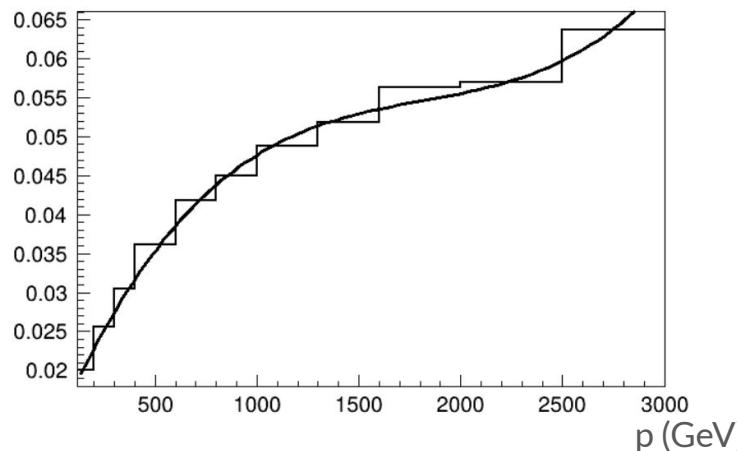


Resolution Smearing

- Need to know: $\sigma_{default}$ as a function of p .
In MC, fit the residual: $\frac{1/p - 1/p^{gen}}{1/p^{gen}}$ in bins of “ p ”:

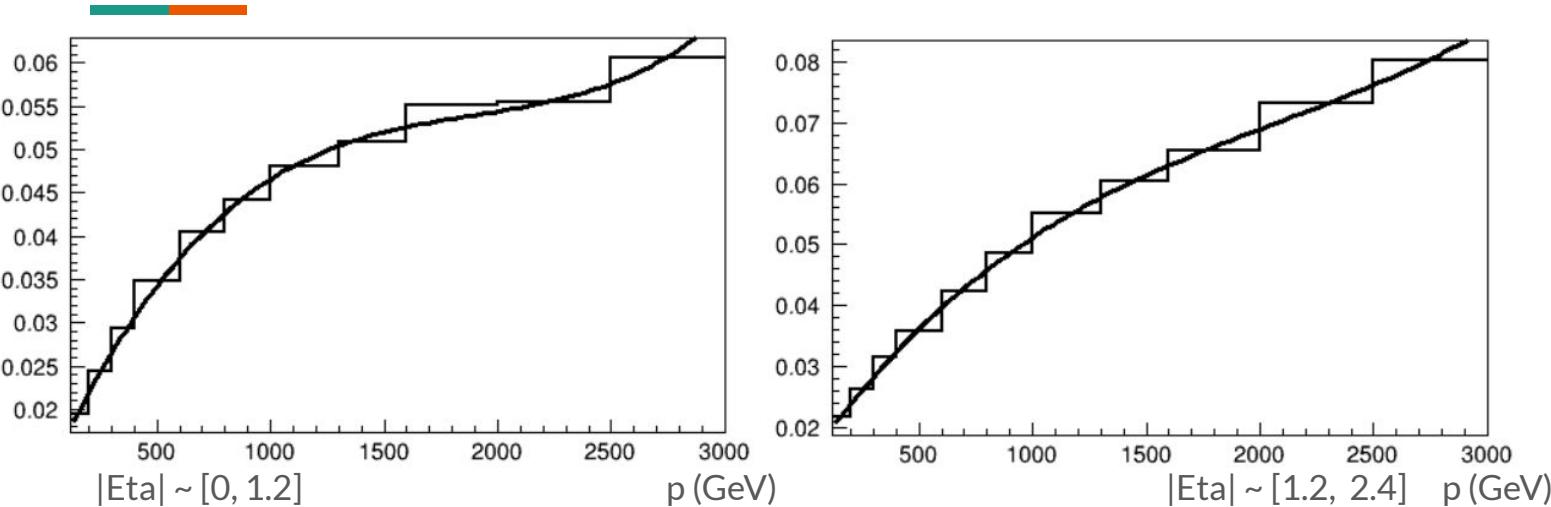
"p" :[120, 200, 300, 400, 600, 800, 1000, 1300, 1600, 2000, 2500, 3000],

Fit the resolution of residual as a function of p with polynomial



E.g. 2022 Eta_[0,1.2]

Era_2022



Paramters for Eta_[0, 1.2] :

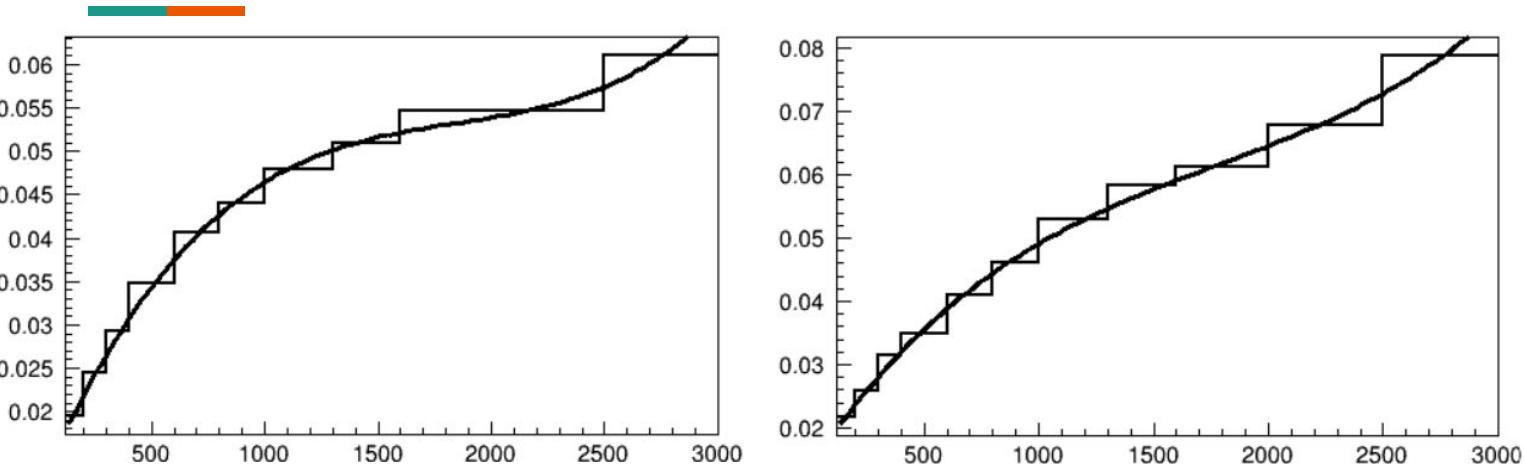
NO.	NAME	VALUE
1	A	1.11520e-02
2	B	5.94955e-05
3	C	-2.92229e-08
4	D	5.14353e-12

FCN=8.16307e-05 FROM MINOS
EDM=1.167

Eta_[1.2, 2.4] :

EXT PARAMETER NO.	NAME	VALUE
1	A	1.40545e-02
2	B	5.28169e-05
3	C	-1.87005e-08
4	D	3.00799e-12

Era 2022 , Apply q/p modification

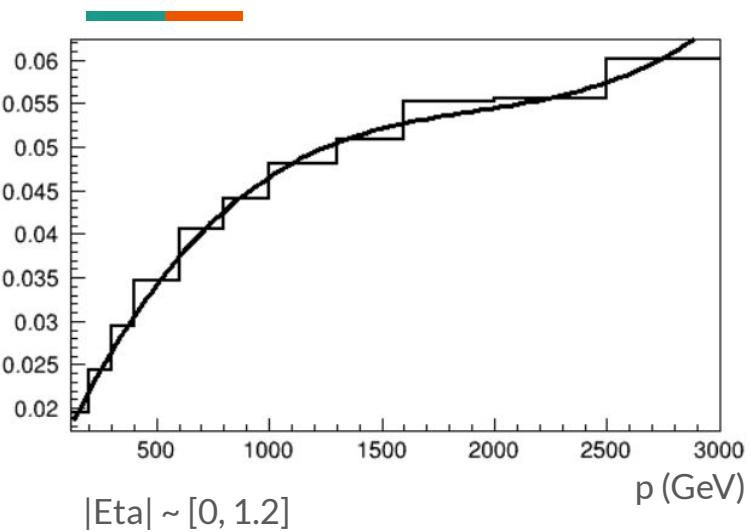


NO.	NAME	VALUE
1	A	1.09739e-02
2	B	6.06340e-05
3	C	-3.06497e-08
4	D	5.53341e-12

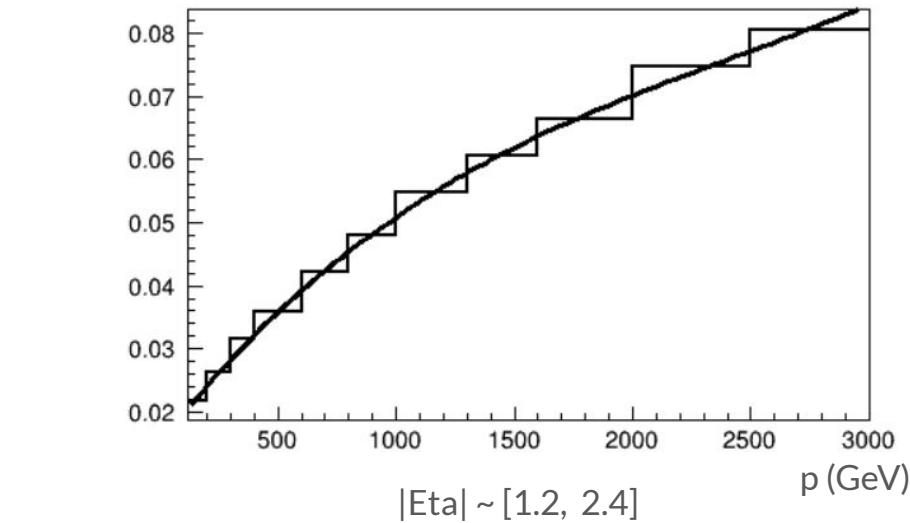
CN=0.000151367 FROM MINOS
EDM=5.470

EXT PARAMETER	NO.	NAME	VALUE
1	A	1.39045e-02	
2	B	5.35022e-05	
3	C	-2.25294e-08	
4	D	4.22415e-12	

Era 2022_EE



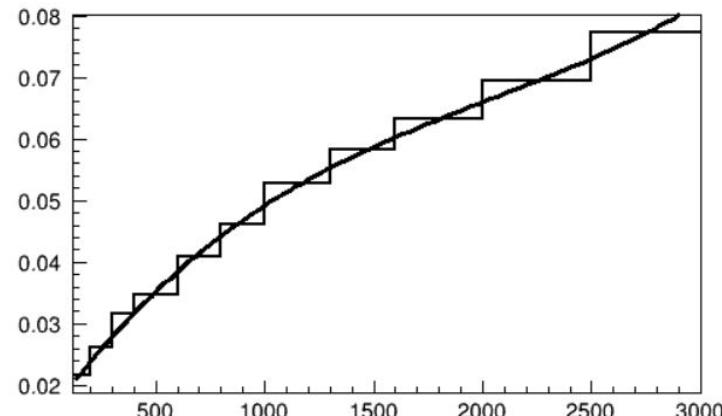
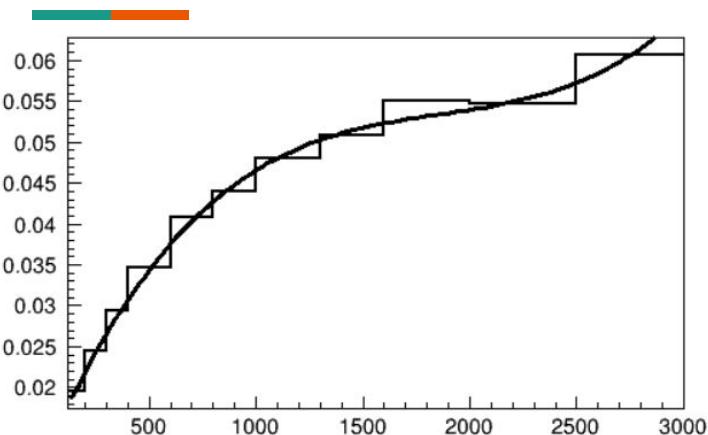
Paramters for Eta_[0, 1.2] :



Eta_[1.2, 2.4] :

NO.	NAME	VALUE
1	A	1.12593e-02
2	B	5.88829e-05
3	C	-2.84651e-08
4	D	4.92264e-12
FCN=0.00011333 FROM MINOS		
EDM=7.2284		
EXT PARAMETER	NAME	VALUE
1	A	1.50148e-02
2	B	4.81069e-05
3	C	-1.41678e-08
4	D	1.95231e-12

Era 22_EE , Apply q/p modification

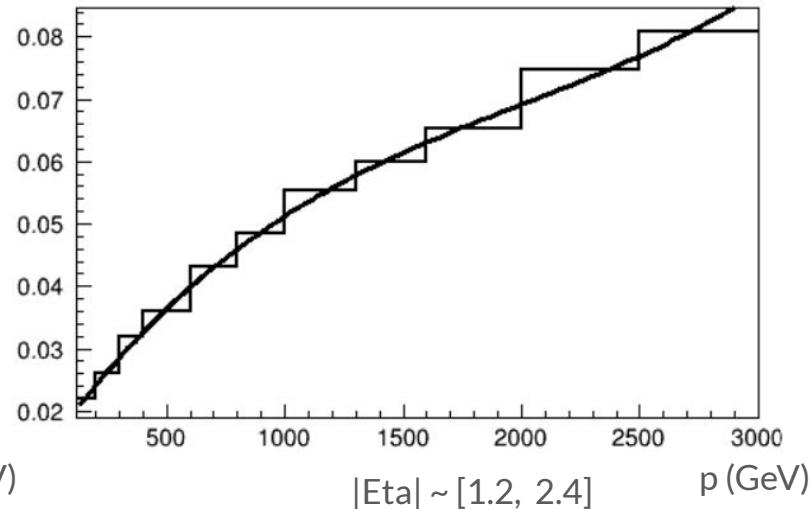
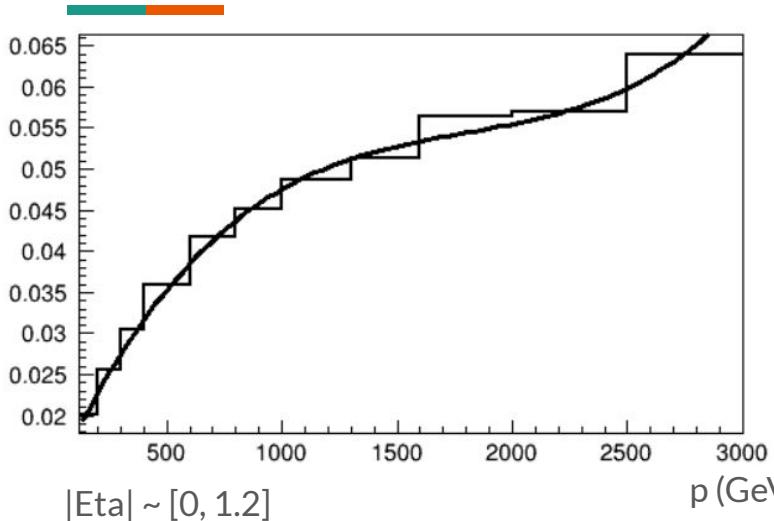


NO.	NAME	VALUE
1	A	1.09851e-02
2	B	6.05789e-05
3	C	-3.04224e-08
4	D	5.44030e-12

CN=0.000139119 FROM MINOS
EDM=6.7675

EXT	PARAMETER	NO.	NAME	VALUE
1	A	1	A	1.47765e-02
2	B	2	B	4.89326e-05
3	C	3	C	-1.73438e-08
4	D	4	D	2.84056e-12

Era_2023



Paramters for Eta_[0, 1.2] :

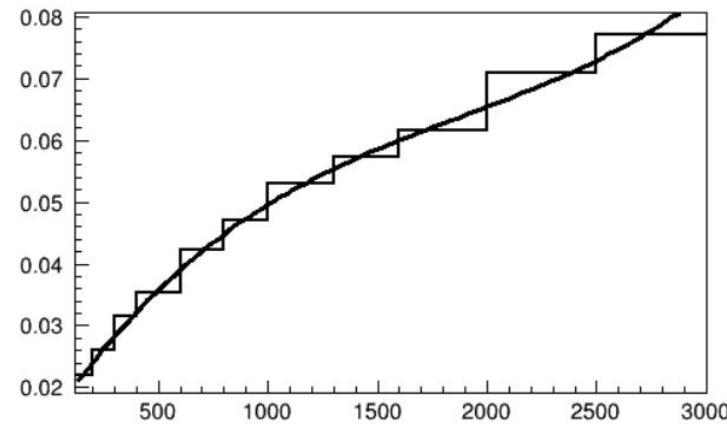
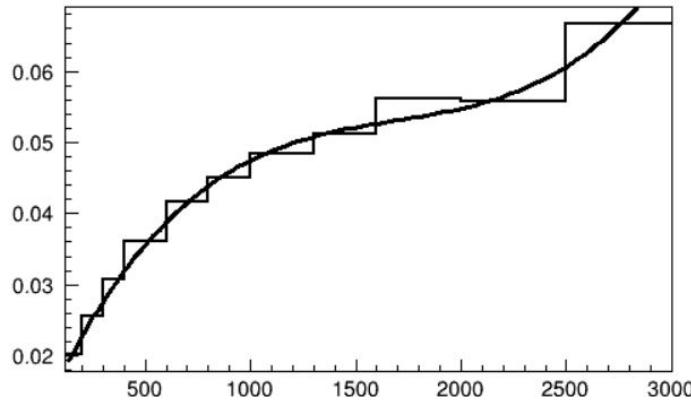
NO.	NAME	VALUE
1	A	1.17191e-02
2	B	6.14187e-05
3	C	-3.14152e-08
4	D	5.81988e-12

FCN=0.000144114 FROM MINOS
EDM=2.877

Eta_[1.2, 2.4] :

EXT	PARAMETER	NO.	NAME	VALUE
		1	A	1.42399e-02
		2	B	5.30925e-05
		3	C	-1.92247e-08
		4	D	3.20733e-12

Era_2023, Apply q/p modification

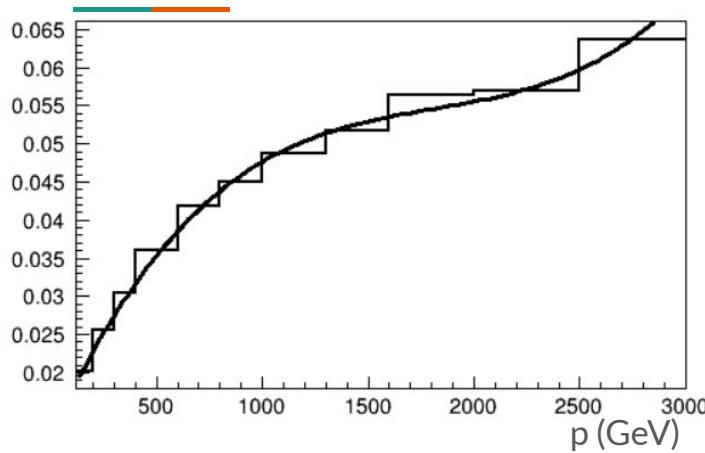


NO.	NAME	VALUE
1	A	1.12477e-02
2	B	6.46444e-05
3	C	-3.53990e-08
4	D	6.98089e-12

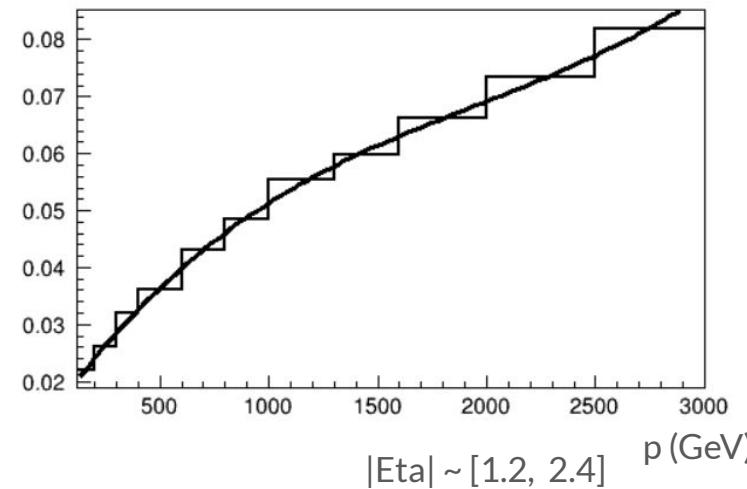
CN=0.000161953 FROM MINOS
EDM=1.173

EXT	PARAMETER	NAME	VALUE
1	A	1.43740e-02	
2	B	5.23100e-05	
3	C	-2.07144e-08	
4	D	3.66499e-12	

Era_2023BPix



$|\text{Eta}| \sim [0, 1.2]$



$|\text{Eta}| \sim [1.2, 2.4]$

Paramters for Eta_[0, 1.2] :

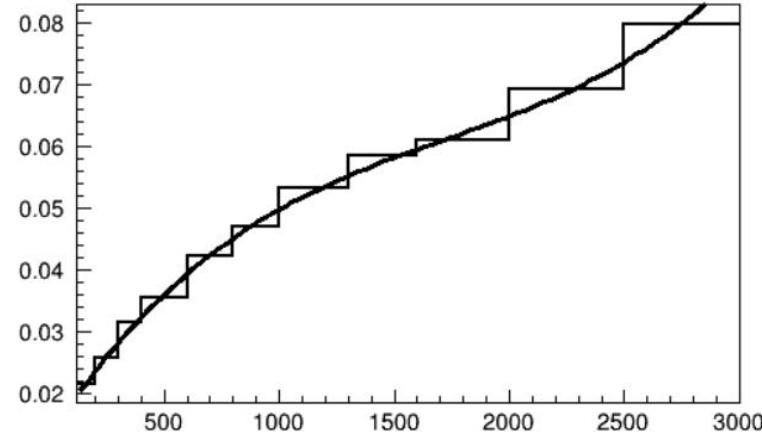
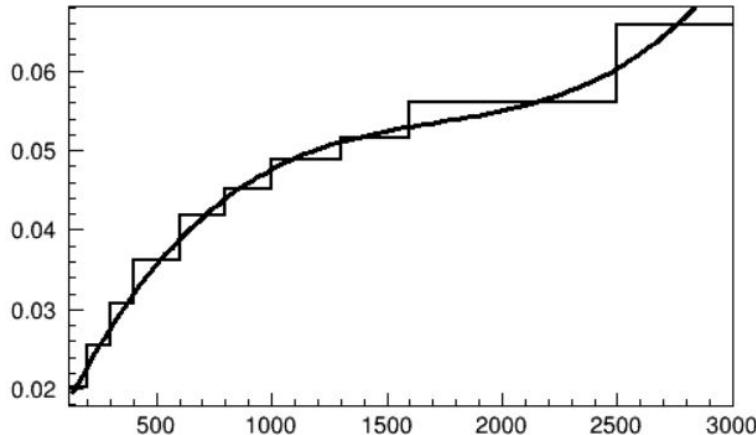
NAME	VALUE
A	1.17770e-02
B	6.13538e-05
C	-3.12168e-08
D	5.74134e-12

8.9033e-05 FROM MINOS
EDM=2.5150

Eta_[1.2, 2.4] :

PARAMETER	NAME	VALUE
A	A	1.40625e-02
B	B	5.38352e-05
C	C	-1.99658e-08
D	D	3.41569e-12

Era_2023BPix, Apply q/p modification

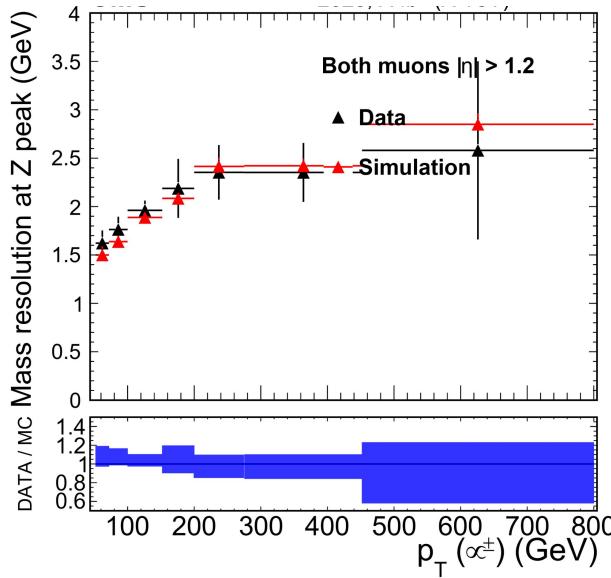
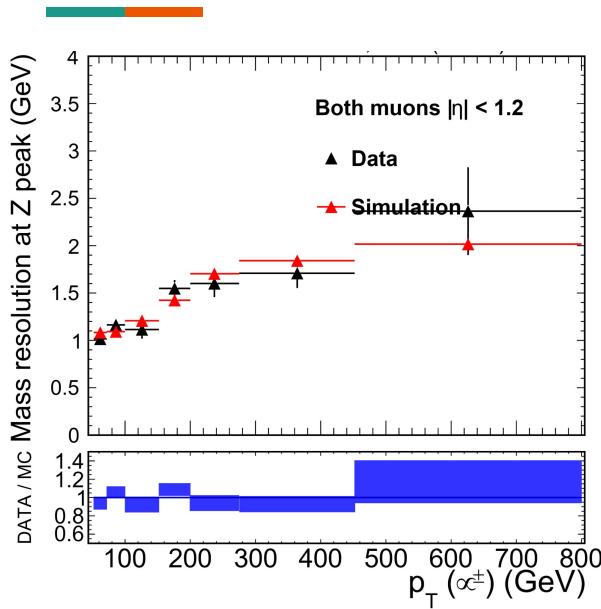


NO.	NAME	VALUE
1	A	1.13663e-02
2	B	6.41160e-05
3	C	-3.45094e-08
4	D	6.68243e-12

FCN=0.000120231 FROM MINOS
EDM=4.6188

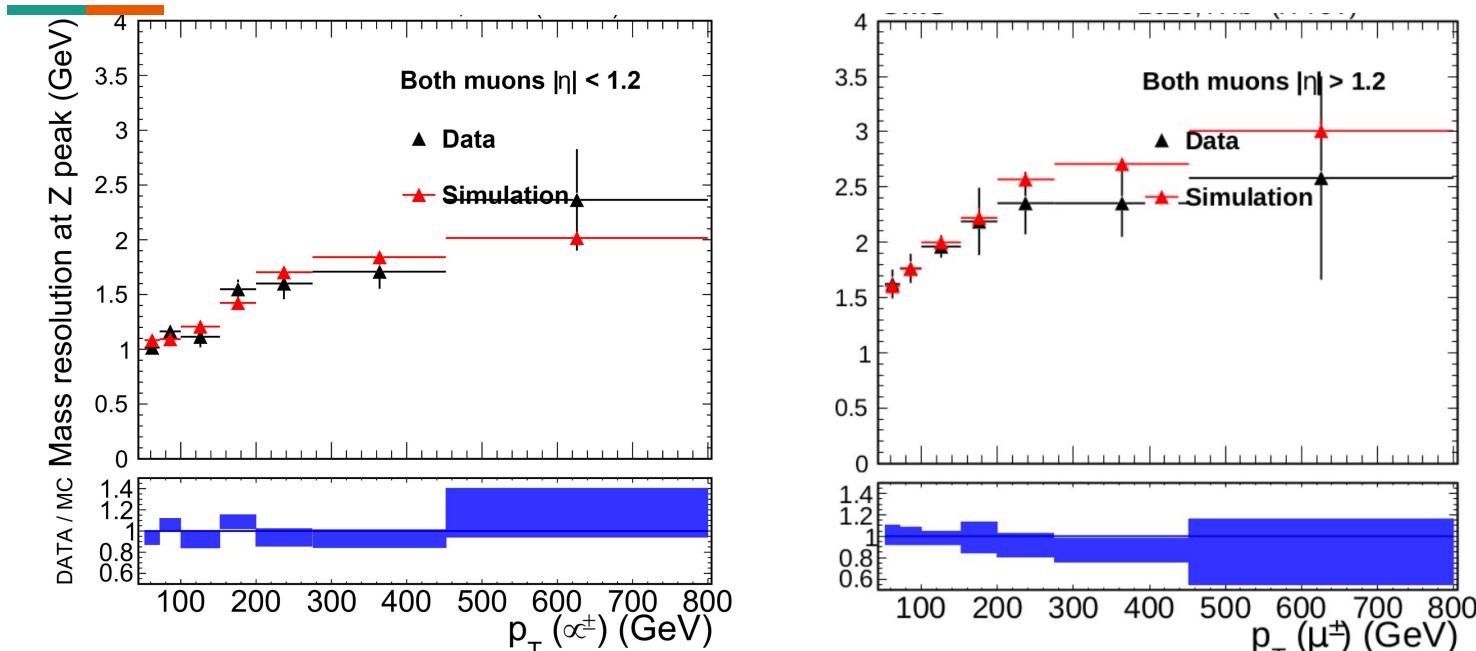
EXT	PARAMETER	NO.	NAME	VALUE
		1	A	1.32530e-02
		2	B	5.72957e-05
		3	C	-2.56409e-08
		4	D	4.95473e-12

Era 2022



- Good agreement between Data and MC for central muons ($|\eta| < 1.2$)
- Data resolution slightly worse than predicted by MC for forward muons ($|\eta| > 1.2$)
- Will recommend a smearing of forward muons in MC

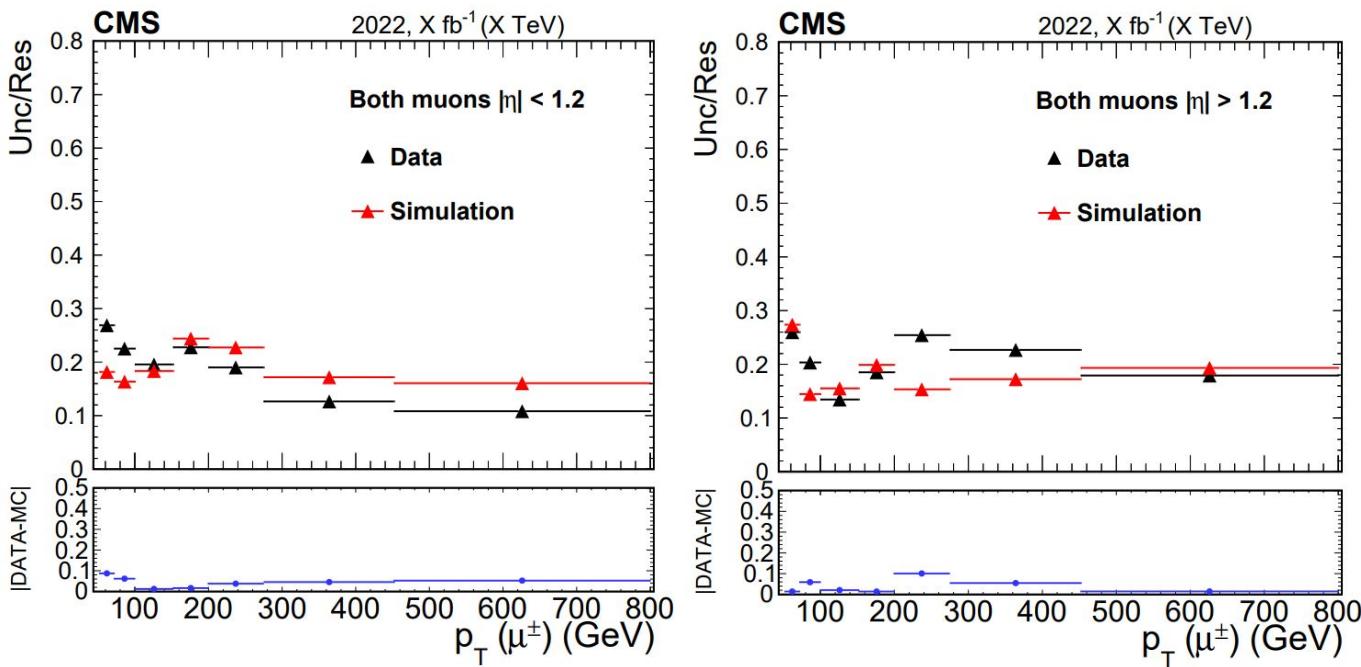
0 smearing in Barrel & 5% smear in Forward region.



- After smearing the muon in the forward region so that the resolution is worse by 5%, very good agreement is observed for forward muons in the high statistics bins
- Average remaining Data/MC difference is $\sim 10\%$
 - covered by systematic uncertainty

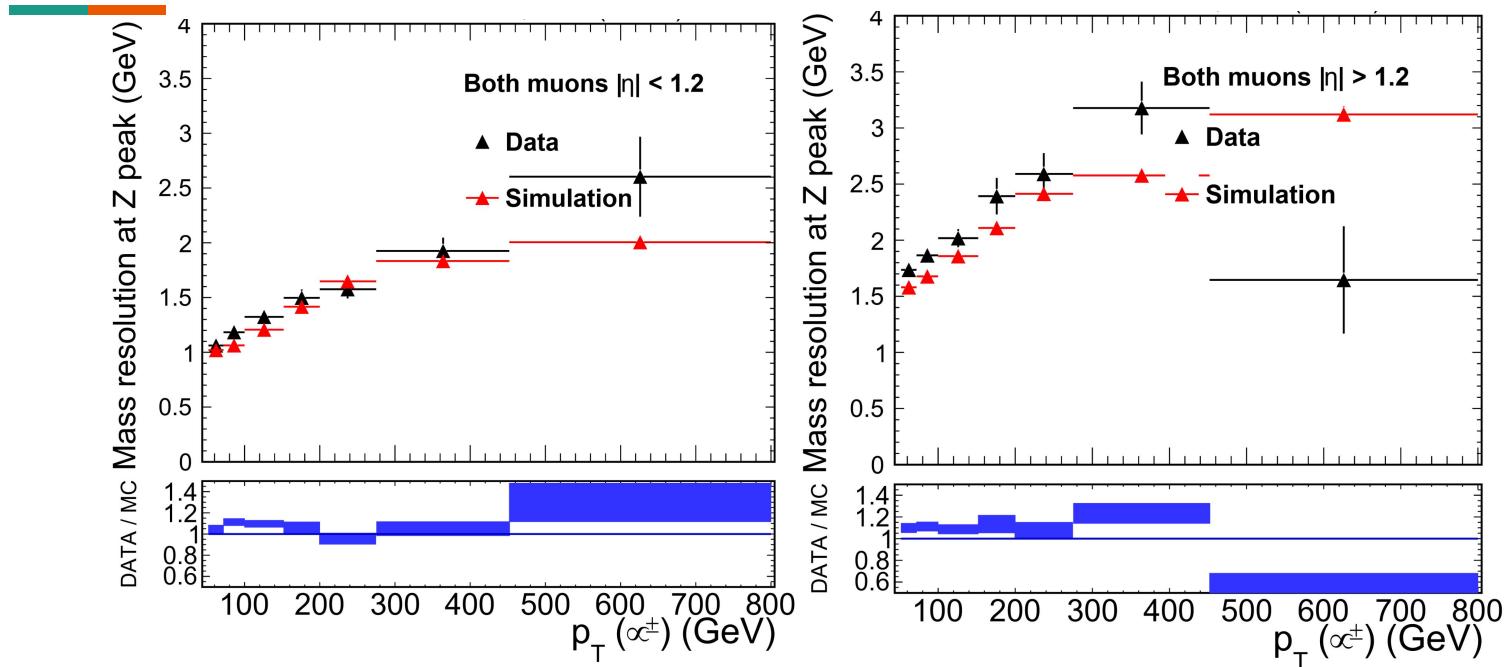
Relative Systematic Uncertainties - 2022

- Plot the total relative systematic uncertainties (quadratic sum of impact of fit functions, binning, and fit range) for both DATA and MC
- In this measurement, we care only about the agreement of DATA and MC
- We therefore look at the difference between the impact of the uncertainties in DATA and MC to assign a total systematic on the agreement



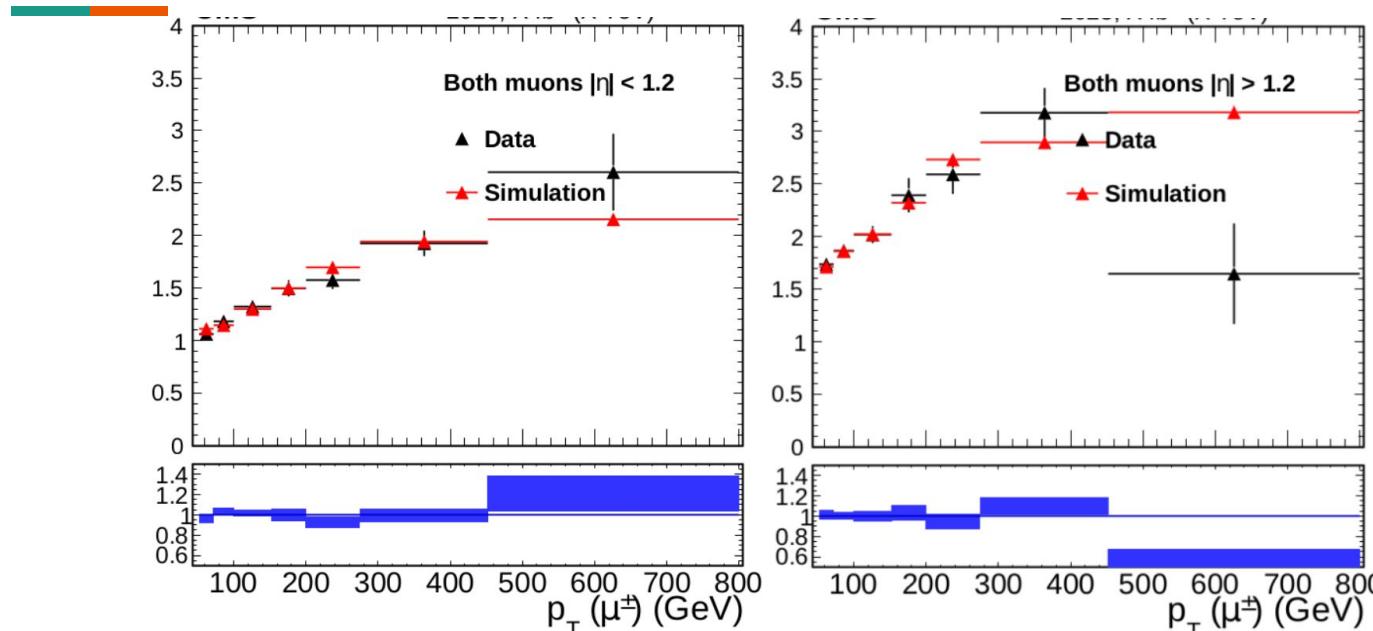
The systematic uncertainty is in general less than 10%, so we recommend a total 10% uncertainty in both Barrel and Forward region to covers the remaining differences after smearing and also covers this systematic uncertainty

Era 2022_EE comparision



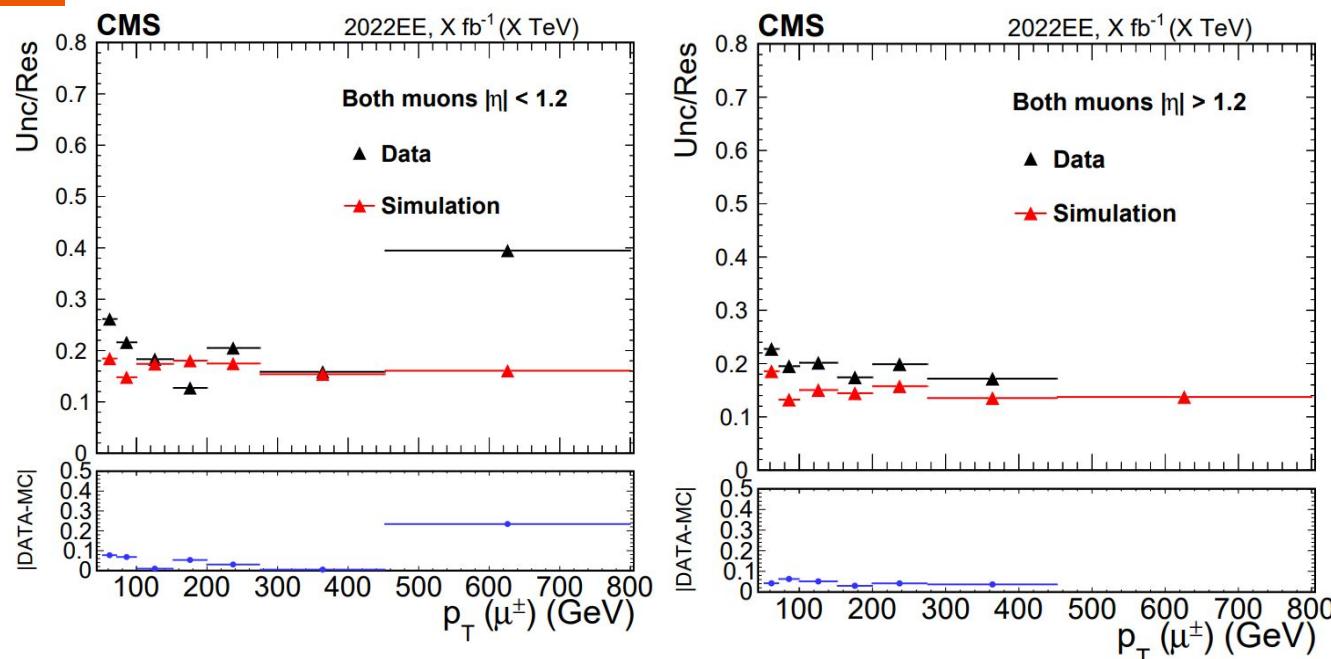
- Generally resolution in MC is better than in data -> need to recommend smearing the muon momenta in MC

5% on Barrel region & 10% on Forward region



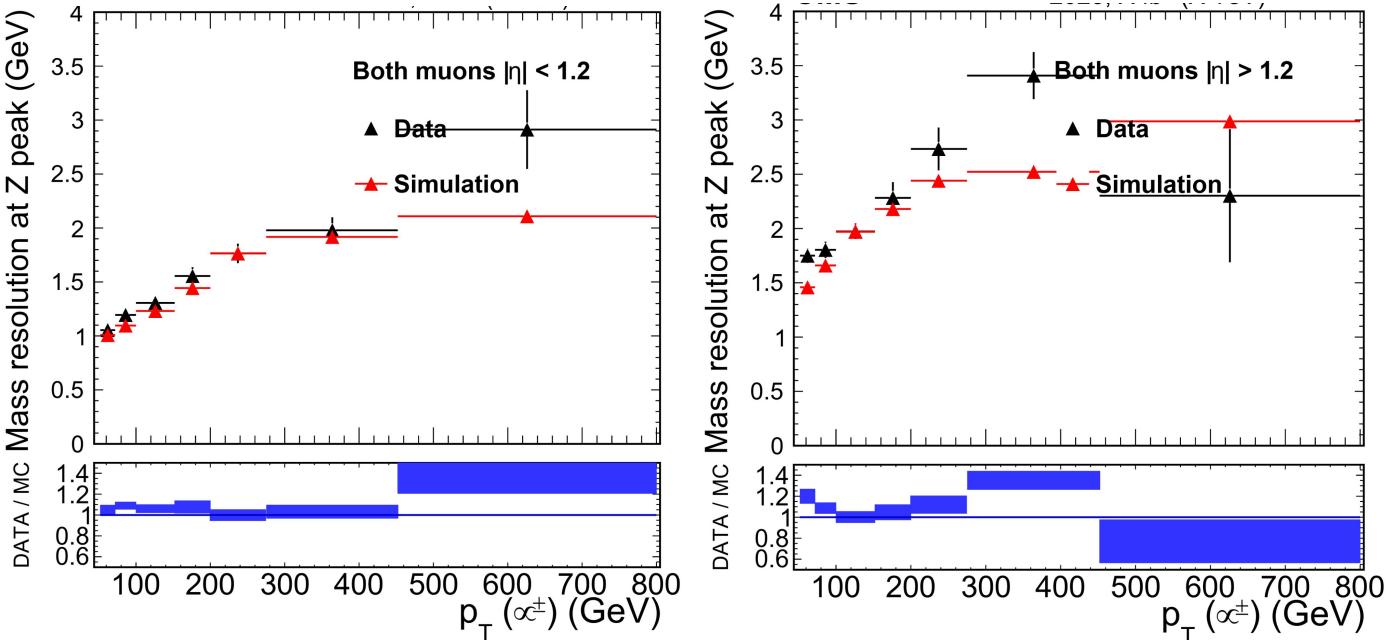
- Smearing momenta in MC by 5% in barrel and 10% in endcap results in very good Data/MC agreement subject to a limited (<5%) uncertainty.

Relative Syst Unc - 2022EE



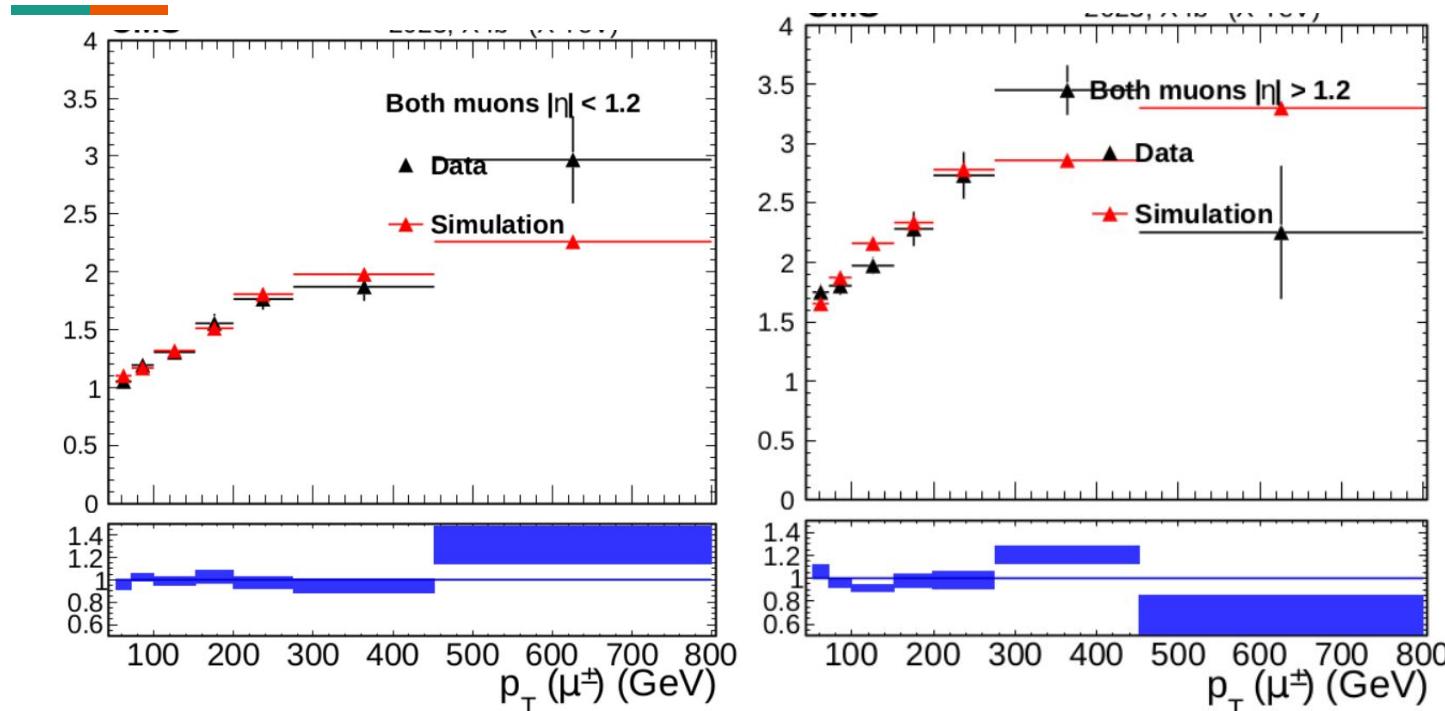
The systematic uncertainty is in general less than 10%, so we recommend a total 10% uncertainty in both Barrel and Forward region to covers the remaining differences after smearing and also covers this systematic uncertainty

Era 2023



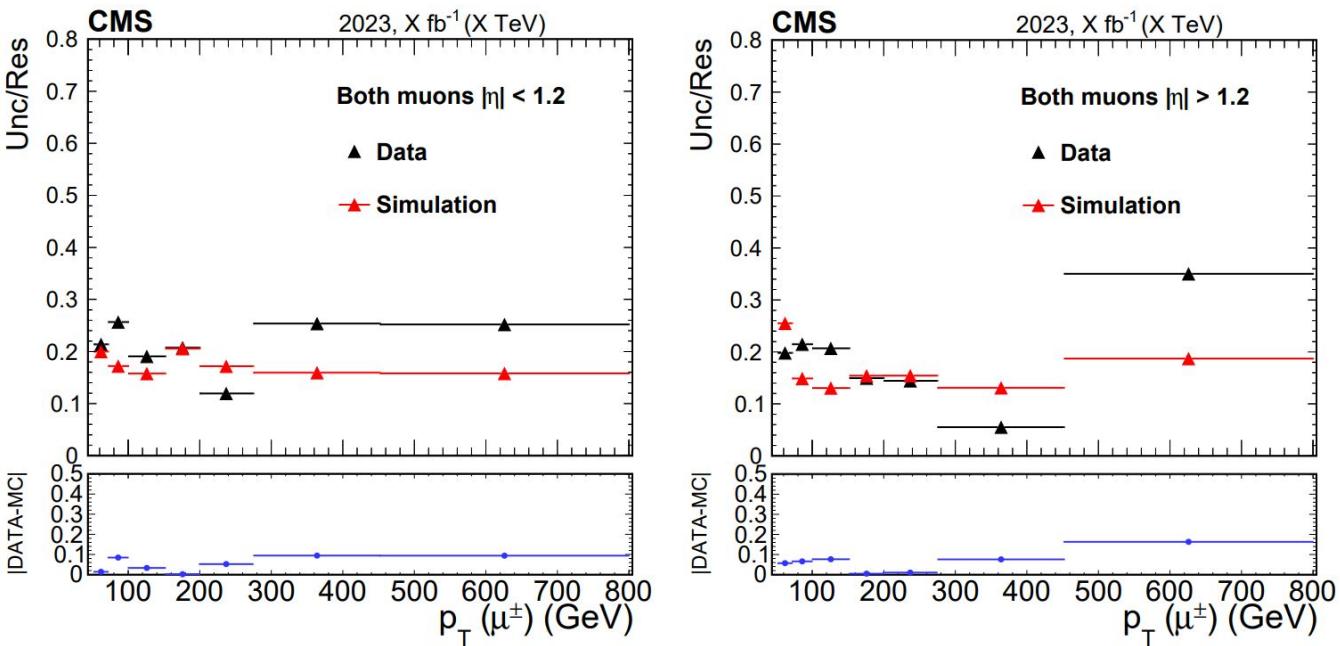
- Generally, resolution in MC is better than in data -> need to recommend smearing the muon momenta in MC

5% in Barrel region and 10% in forward region



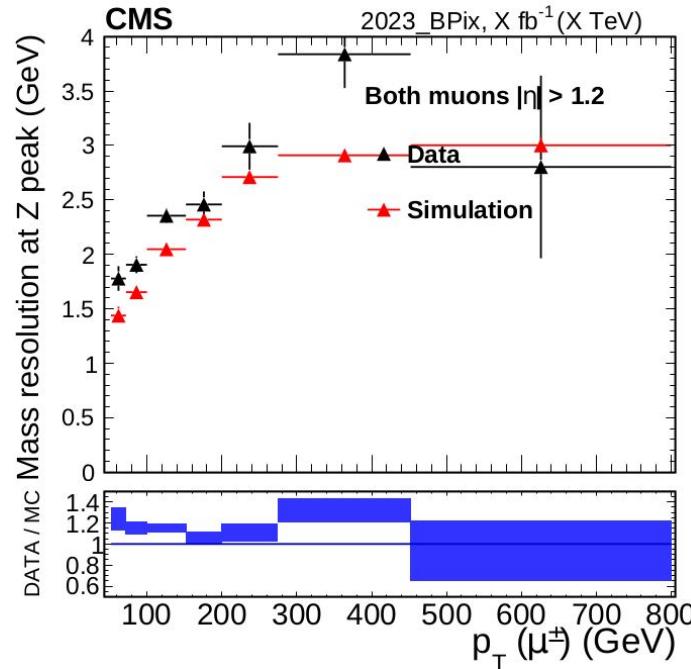
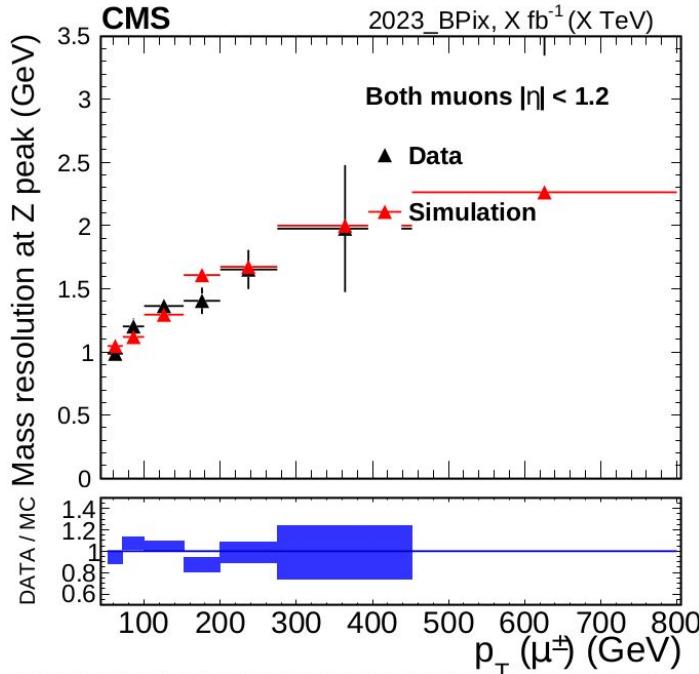
- Data/MC agree well after smearing up to a $<10\%$ uncertainty if drop the high Pt bins.

Relative Syst Unc - 2023



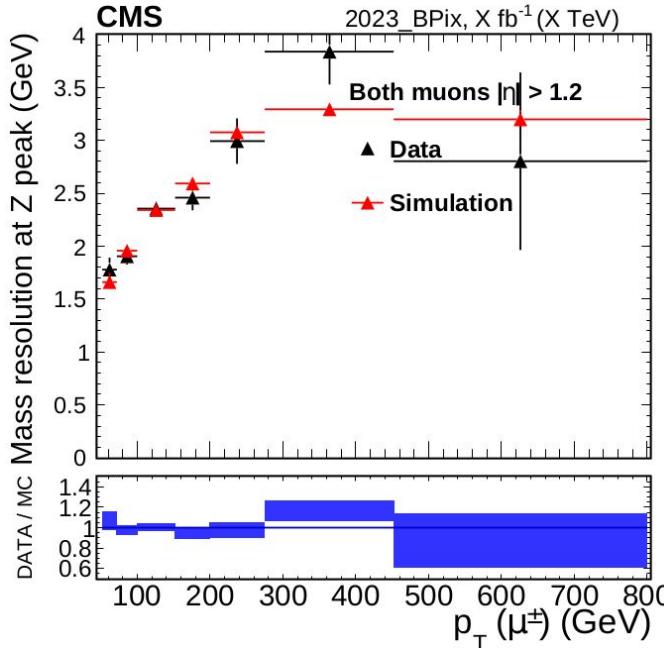
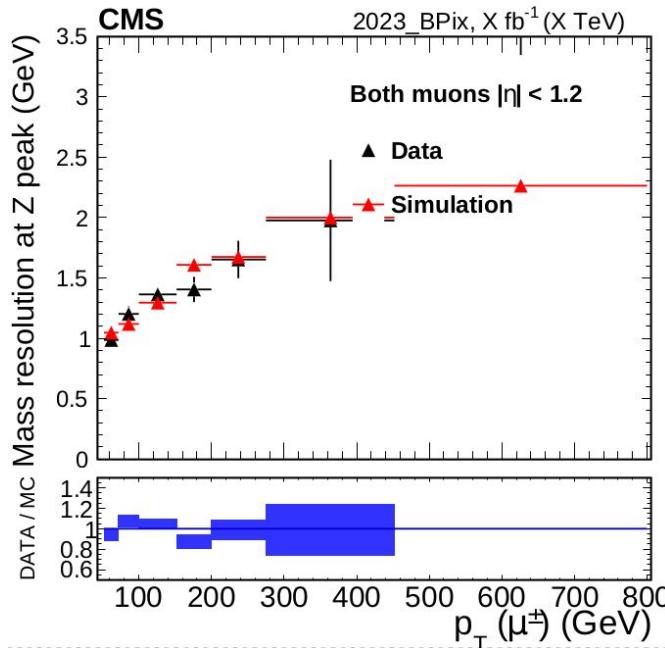
- The systematic uncertainty is in general less than 10%, so we recommend a total 10% uncertainty in both Barrel and Forward region to covers the remaining differences after smearing and also covers this systematic uncertainty

Era 2023_PBix comparision



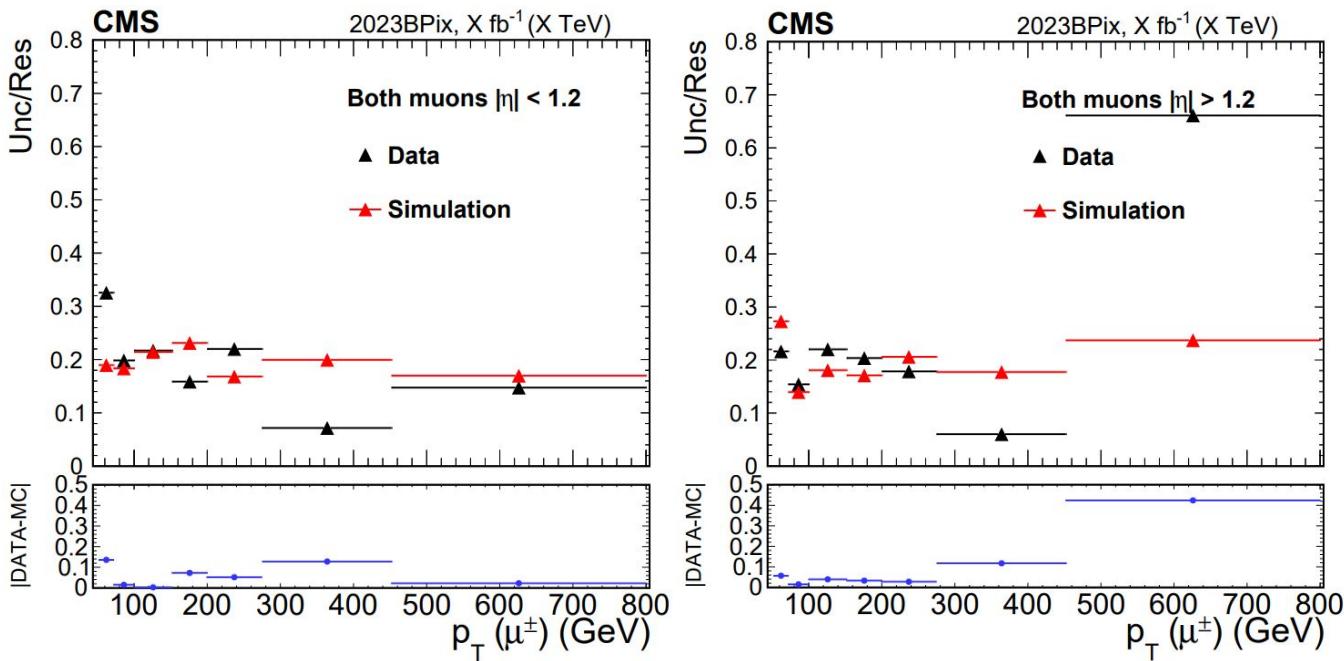
- Very good agreement in barrel region, about 15% better resolution in MC compared to data in endcap

0% on Barrel region & 15% on Forward region



- Good agreement in endcap restored after smearing up to a <10% uncertainty if drop the high Pt bins

Relative Syst Unc - 2023BPix

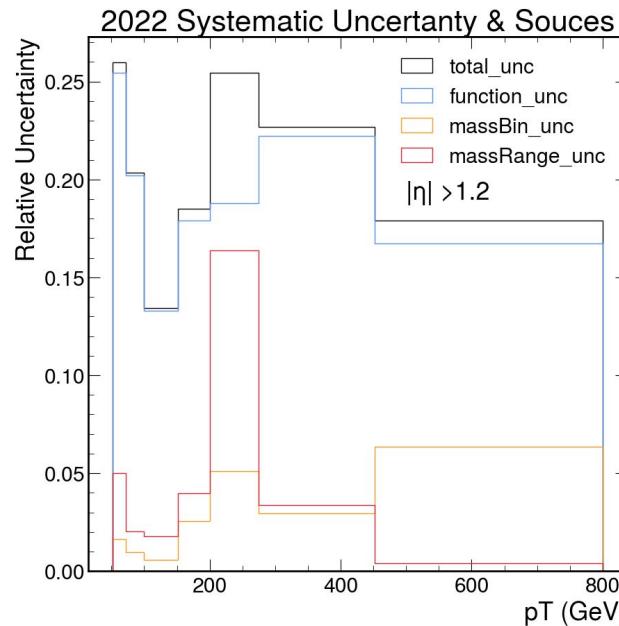
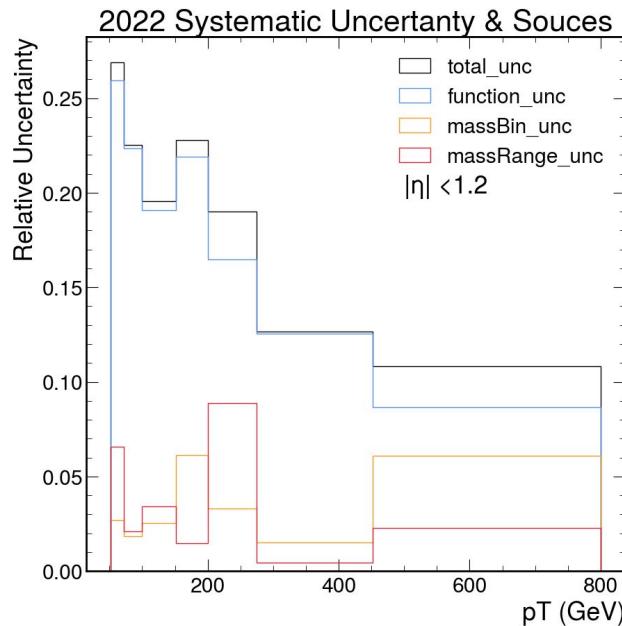


- The systematic uncertainty is in general less than 10%, so we recommend a total 10% uncertainty in both Barrel and Forward region to covers the remaining differences after smearing and also covers this systematic uncertainty
- This disregards the very first bin in the barrel region, but low p_T is less relevant for this recommendation

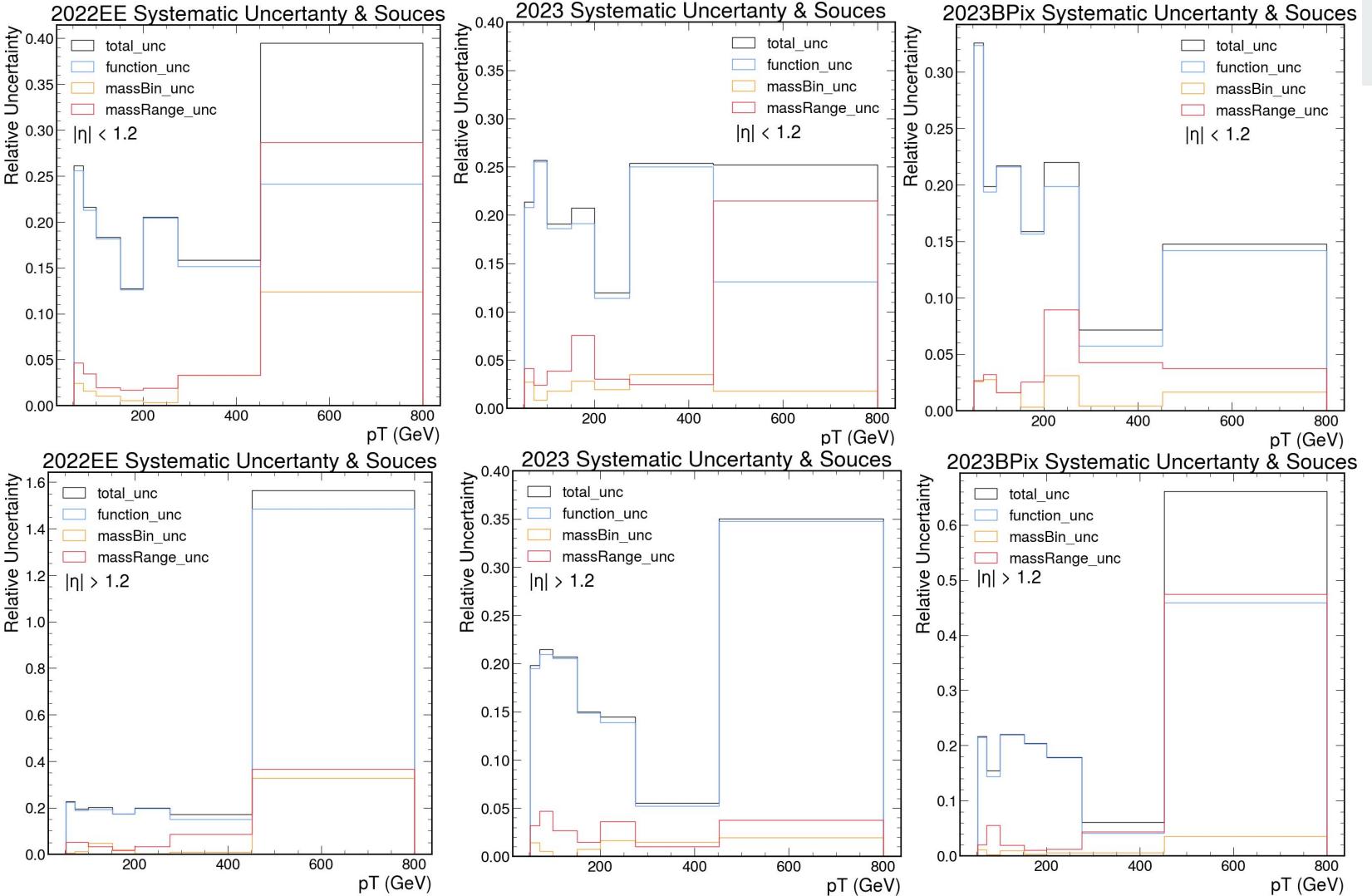
Calculation of Syst Uncertainty

- There are 3 sources :
 - S1. Fitting functions :
 $\max(\text{Crystal-Ball_Unc}, \text{Cruijff_Unc})$ (*default: Double_Crystal-Ball*)
 - S2. massRange shifts
 - S3. massBin shifts
- $\text{Total_Syst_Unc} = \sqrt{S1^2 + S2^2 + S3^2}$

Sources of Systematic Uncertainty



- Choice of fit function is by far the dominant source of uncertainty
- Binning and mass range for fits has mostly a quite small effect
- Consistent picture for all data taking periods, see next slides



Conclusion

- Study of high- p_T muon momentum resolution is complete
- In general, we found good agreement between Data and MC
 - We will recommend that analyst apply additional smearing of the muon momenta in some cases, mostly for forward muons
- An overall systematic uncertainty of 10% will be assigned, that will be applied as additional smearing of the momenta as well
- All necessary information is available to add these recommendation to the Twiki pages
- If we are satisfied with the results, we can update the code in GitHub so it can be integrated in the central POG tool

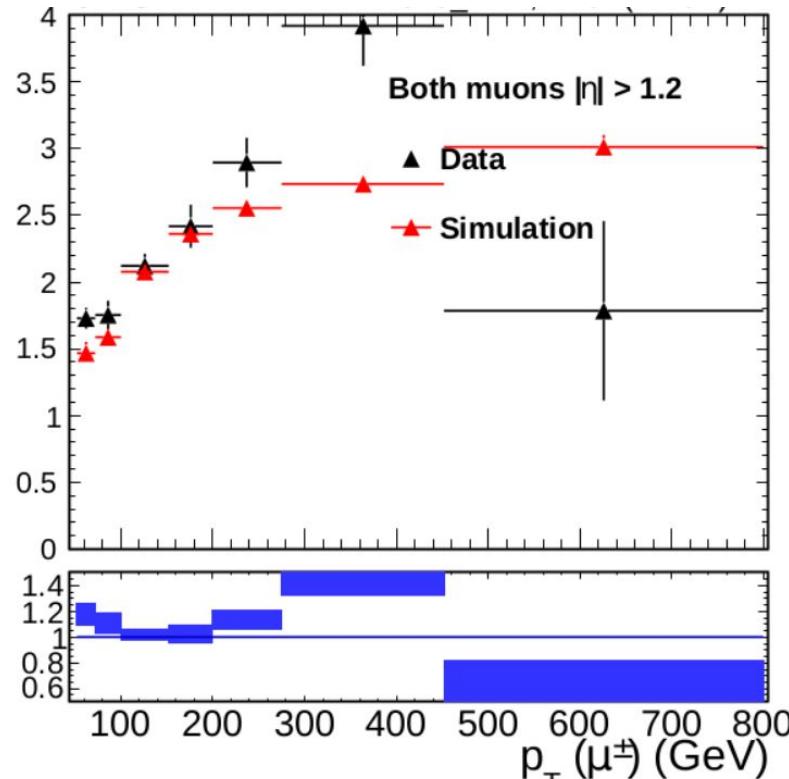
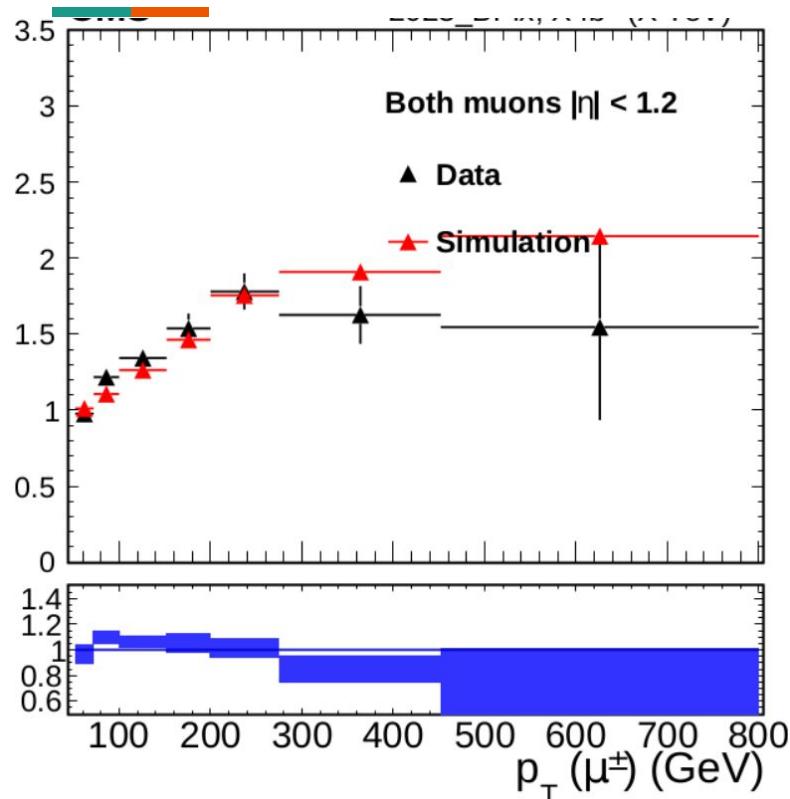


Back Up

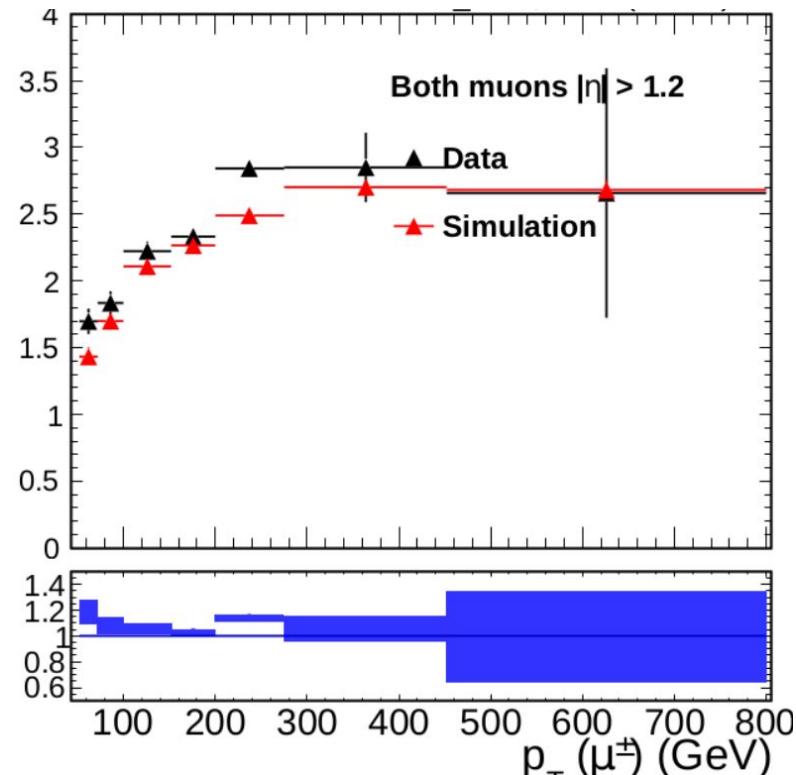
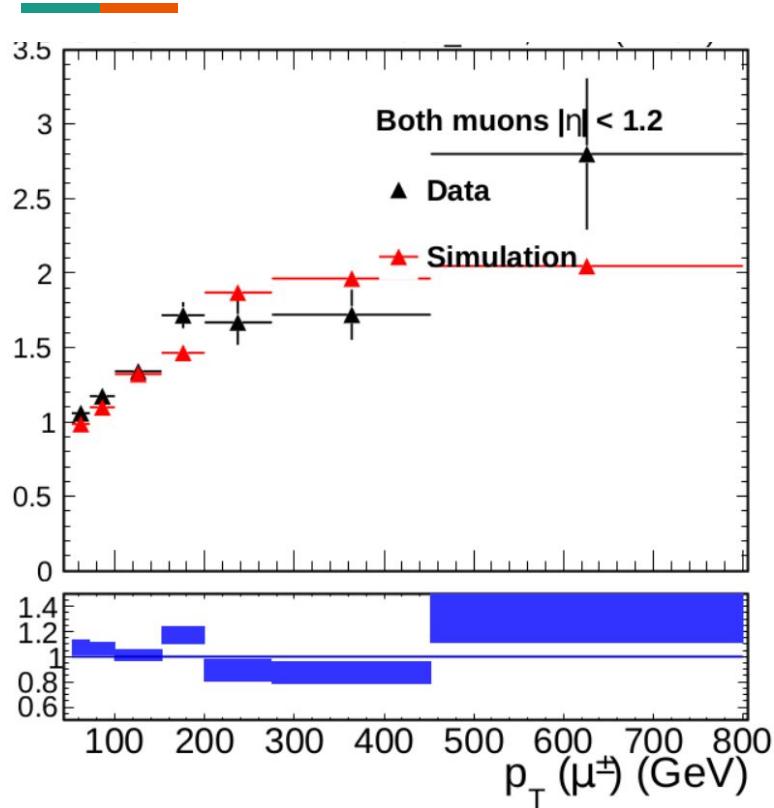
Link to the config file:

https://gitlab.cern.ch/yibo/spark_tnp/-/blob/Run2022_EE/configs/mass_resolution.json?ref_type=heads

2023 tag_charge > 0



2023 tag_charge < 0



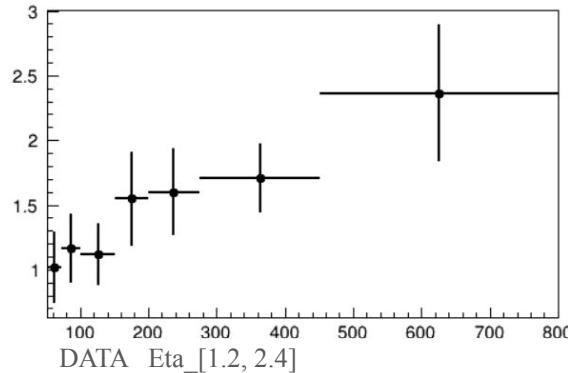
Smear the MC Resolution

- Smear momentum: $p_{\text{smeared}} = p_{\text{default}} \cdot \text{Gaus}(1, \sigma_{\text{default}} \cdot x)$
- Need to know: $\sigma_{\text{default}}(p)$
- In MC, fit the residual: $\frac{1/p - 1/p^{\text{gen}}}{1/p^{\text{gen}}}$
in bins of "p": [120, 200, 300, 400, 600, 800, 1000, 1300, 1600, 2000, 2500, 3000],
- Due to low stat of DY_Boost sample , we use DY_MassBinned for this.
- Fit the resolution of residual as a function of p with polynomial to calculate $\sigma_{\text{default}}(p)$

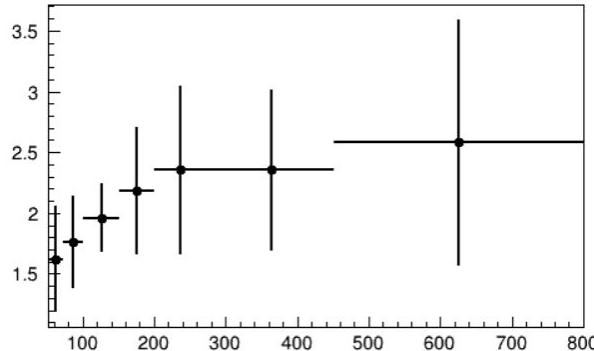
Full Errors : stat + systematic errors, Era 2022



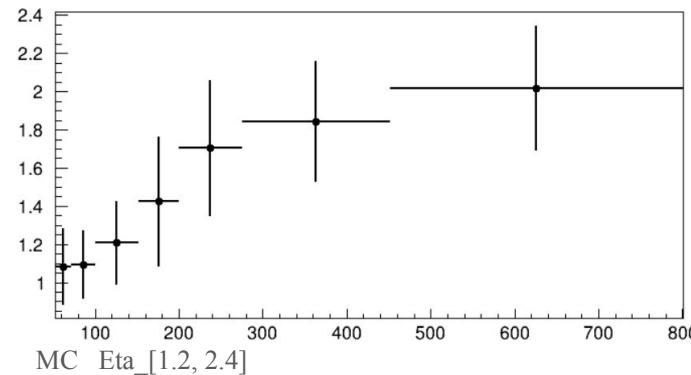
DATA Eta_[0, 1.2]



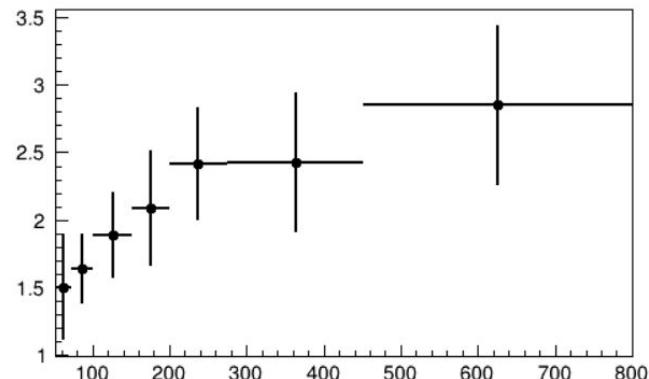
DATA Eta_[1.2, 2.4]



MC Eta_[0, 1.2]



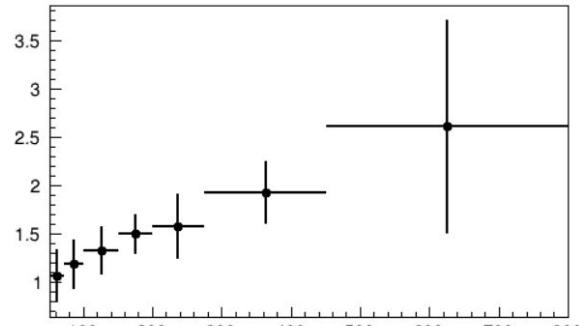
MC Eta_[1.2, 2.4]



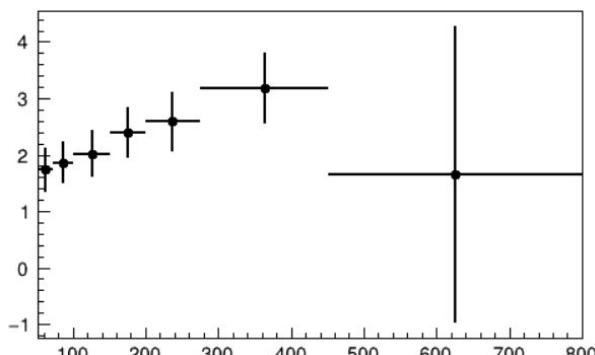
Full Errors : stat + systematic errors , Era 2022_EE



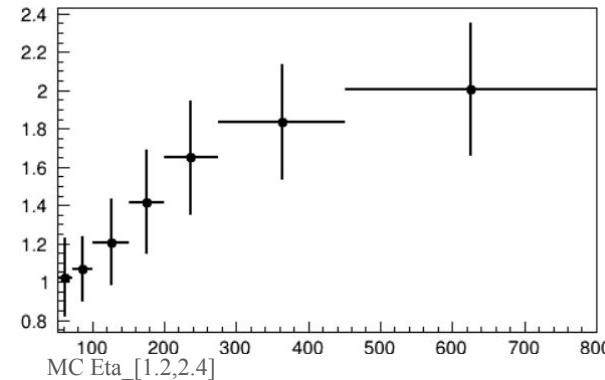
DATA Eta_[0,1.2]



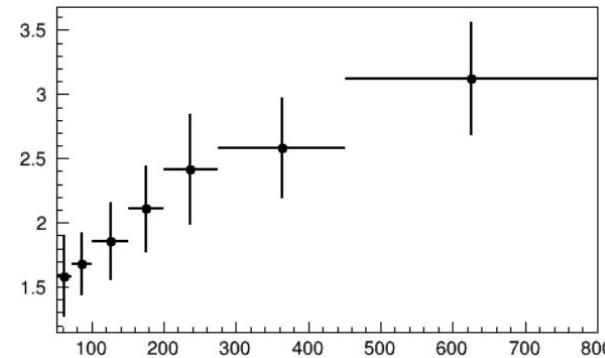
DATA Eta_[1.2,2.4]



MC Eta_[0,1.2]



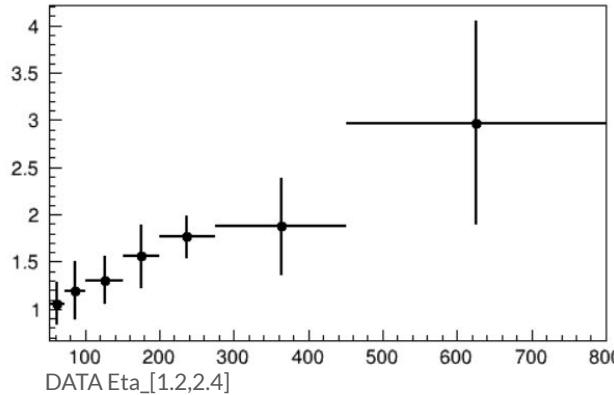
MC Eta_[1.2,2.4]



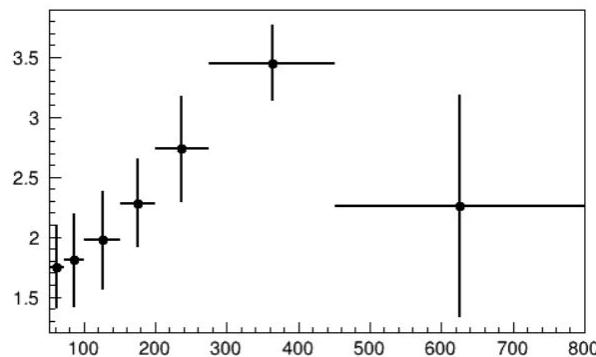
Full Errors : stat + systematic errors , Era 2023



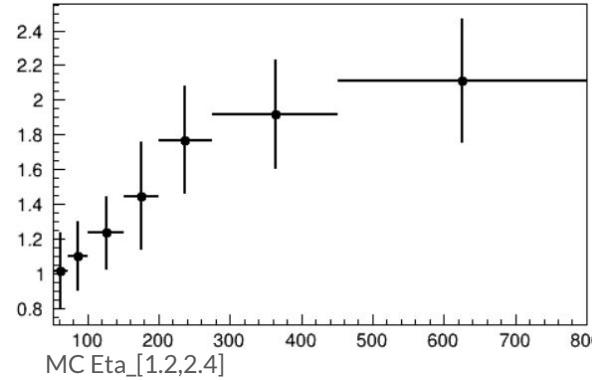
DATA Eta_[0,1.2]



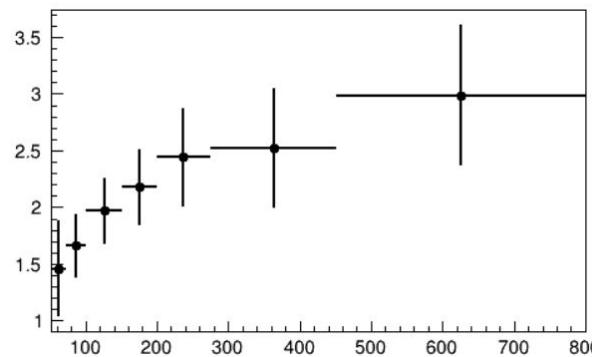
DATA Eta_[1.2,2.4]



MC Eta_[0,1.2]



MC Eta_[1.2,2.4]



Full Errors : stat + systematic errors , Era 2023_BPix

