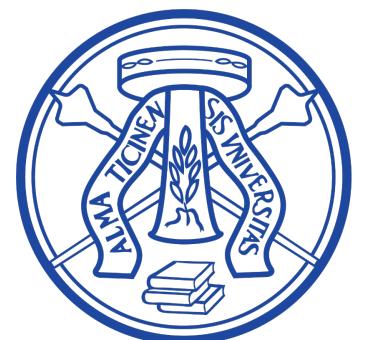


EXTRACTION OF FLAVOR DEPENDENCE IN UNPOLARIZED TMDs

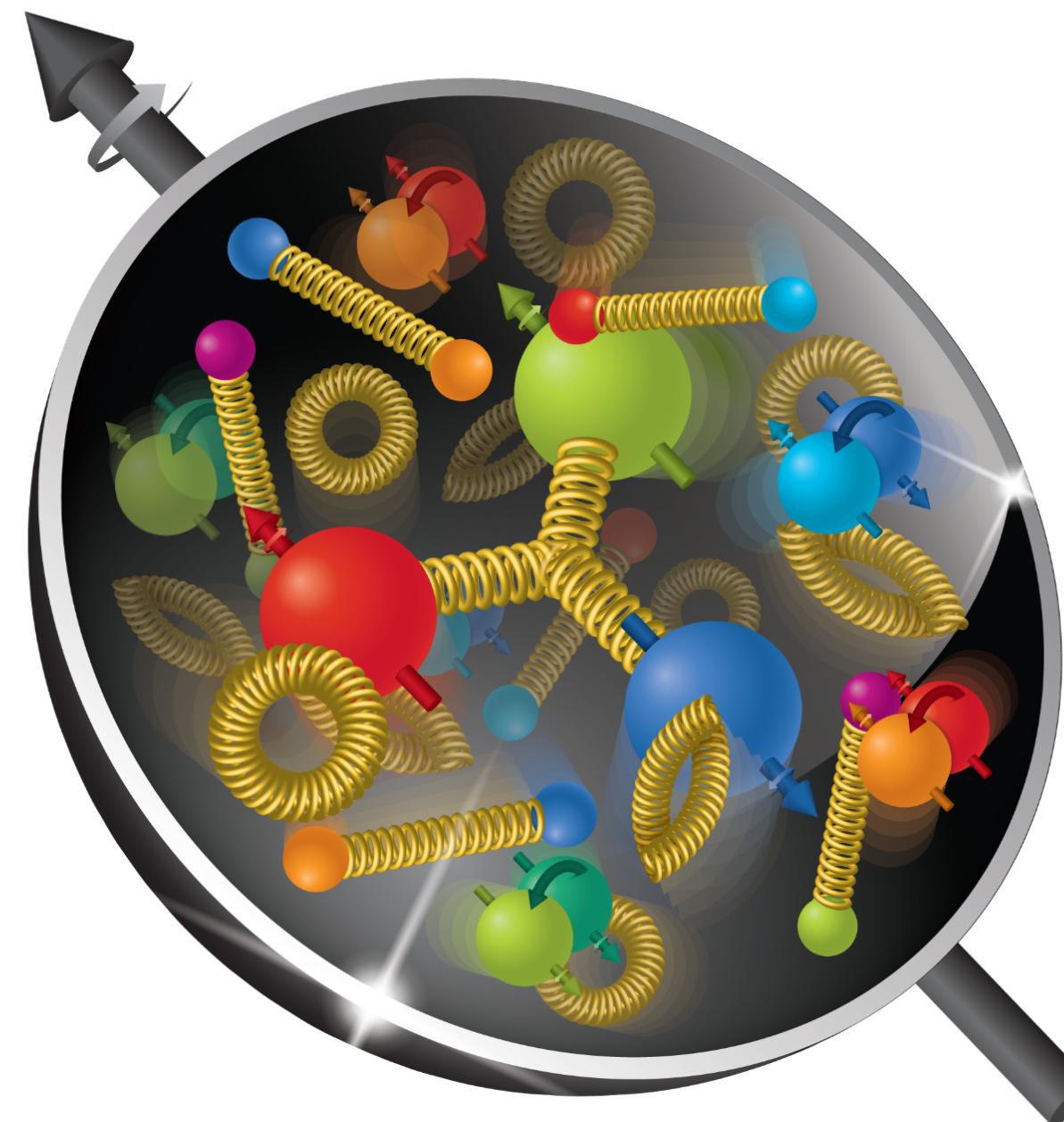
Filippo Delcarro

[JHEP08\(2024\)232](#)
(arXiv: 2405.138833)

[MAP Collaboration](#)



UNIVERSITÀ
DI PAVIA



**Flavor separation is a
fundamental step to fully explore
nucleon structure**

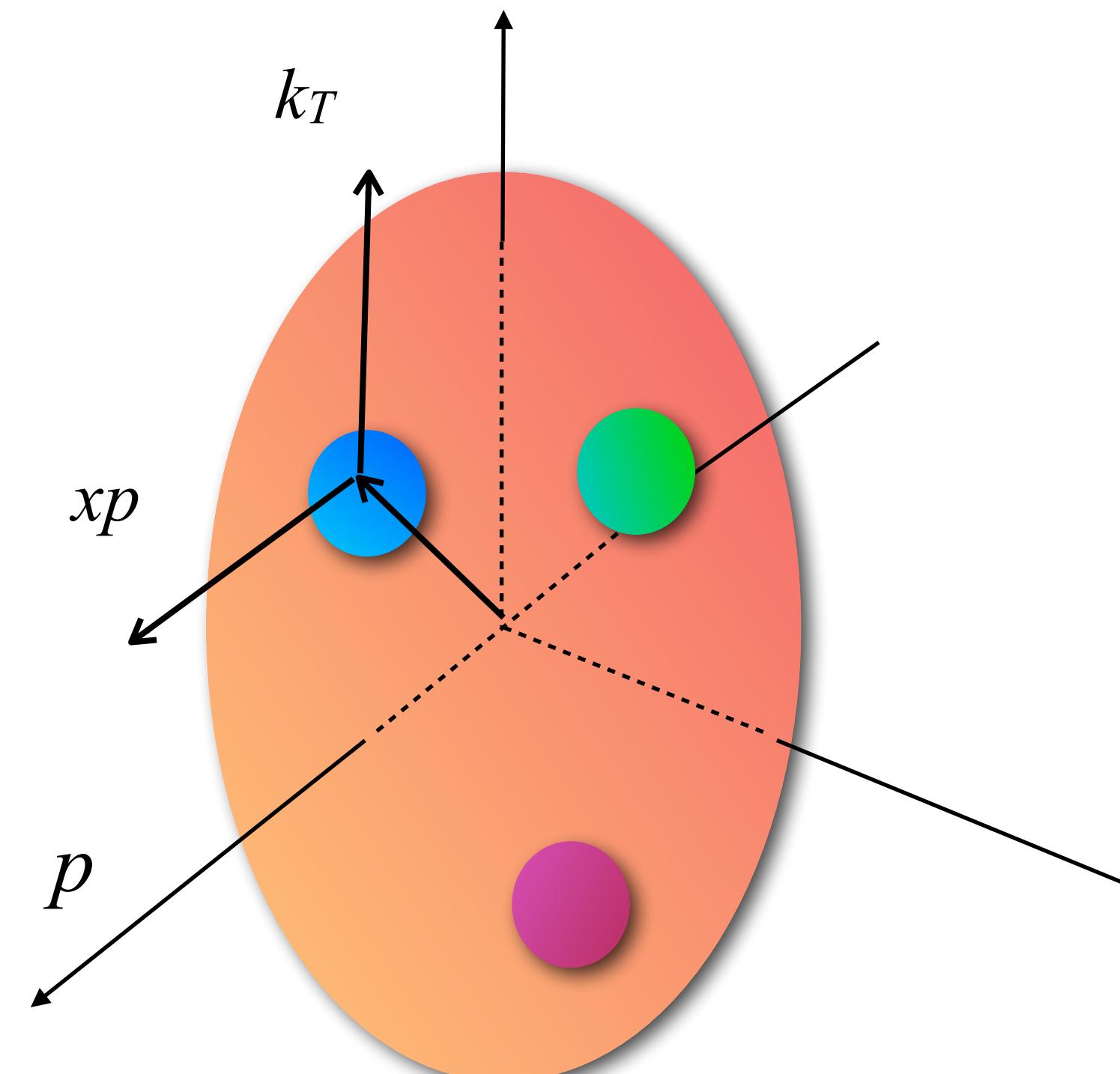
Flavor dependence of unpolarized quark Transverse Momentum Distributions from a global fit

A. Bacchetta, V. Bertone, C. Bissolotti, G. Bozzi, M. Cerutti, FD, M. Radici, L. Rossi, A. Signori

MAP Collaboration [JHEP08\(2024\)232](https://doi.org/10.1007/JHEP08(2024)232)

Transverse Momentum Distributions

3-dimensional map of the internal structure of the nucleon



Universality: same function, multiple processes

(x, k_T) dependence

Q2 energy scale evolution

flavor?

unpolarized TMD PDF
unpolarized TMD FF

nucleon pol.

quark pol.

	U	L	T
U	f_1		h_1^\perp
L		g_{1L}	h_{1L}^\perp
T	f_{1T}^\perp	g_{1T}	h_1, h_{1T}^\perp

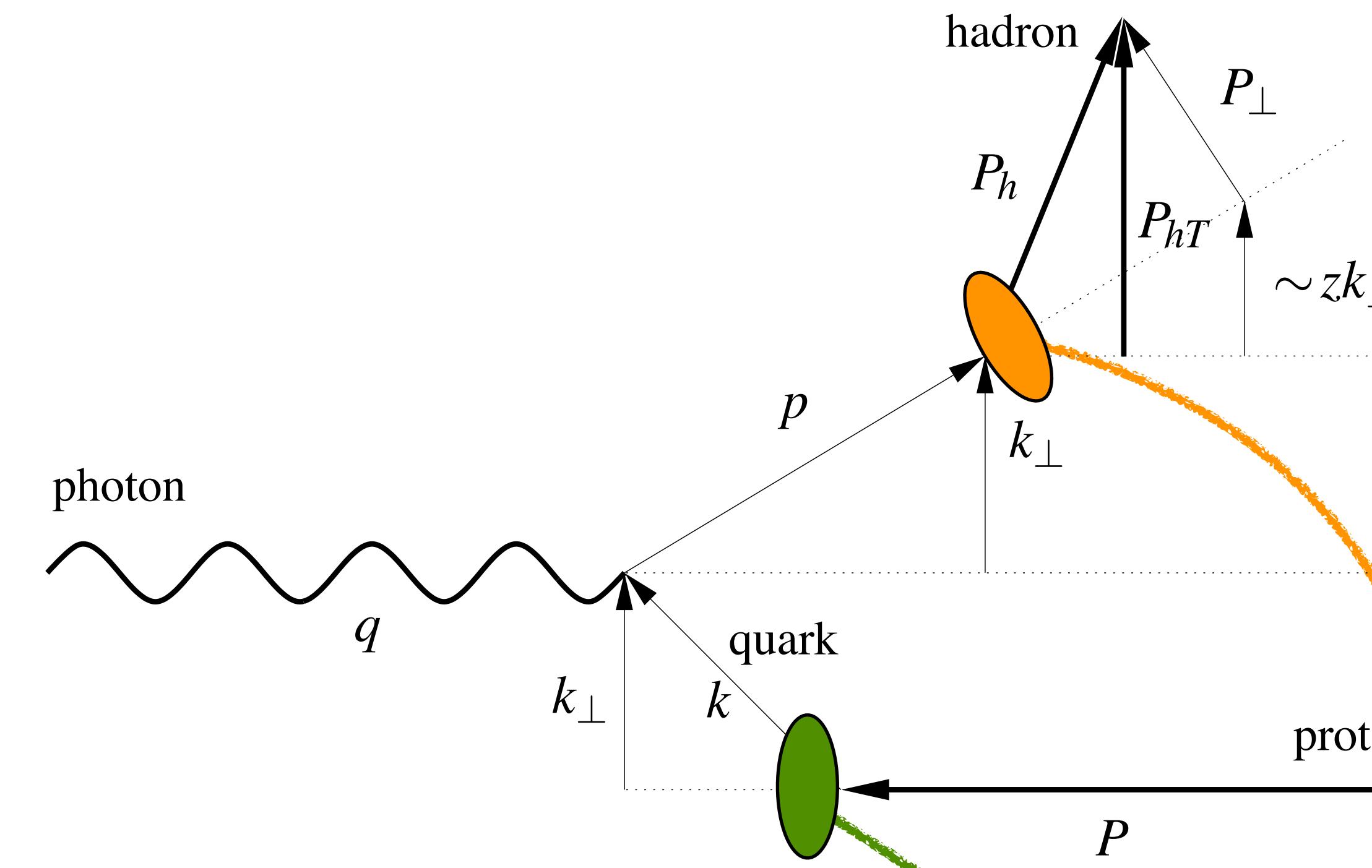
t-odd t-even

TMD formalism: factorization

SIDIS

TMD FF

TMD PDF



Bacchetta, Diehl, et al., JHEP 02 (2007)

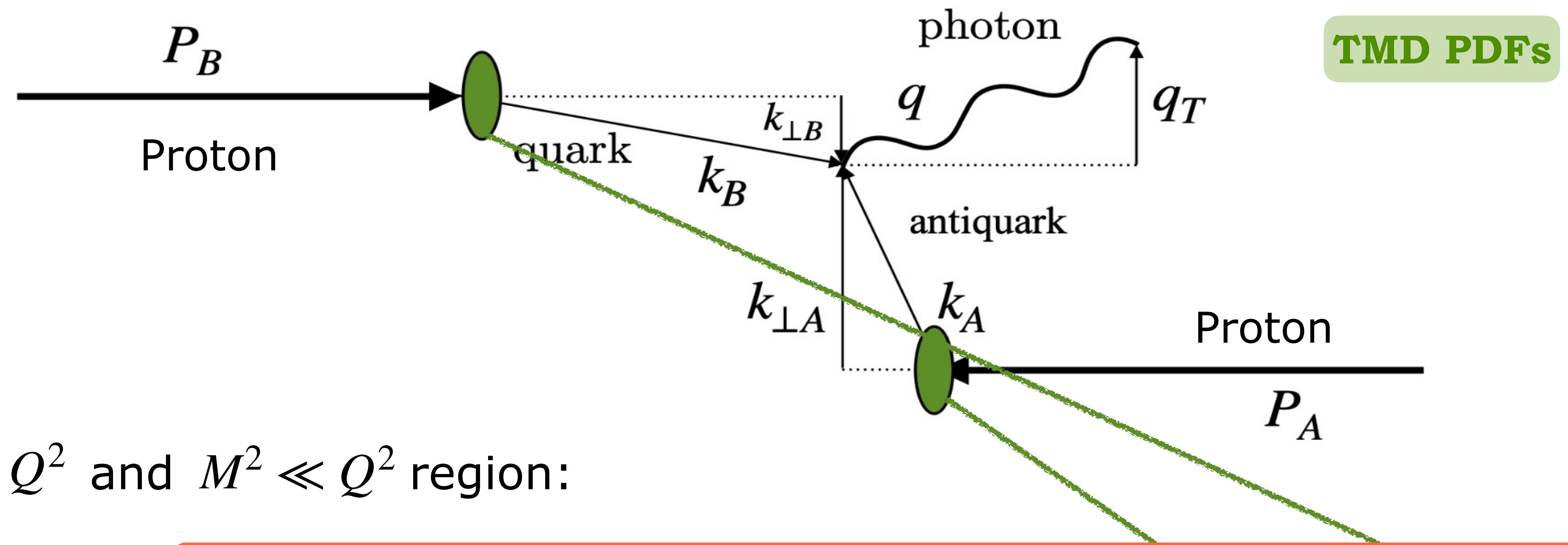
$$F_{UU,T}(x \cdot z; \mu_F, \mathbf{P}_{hT}^2, Q^2) = \boxed{x \sum_a H_{UU,T}^a(Q^2, \mu^2) \int d^2 \mathbf{k}_\perp d^2 \mathbf{P}_\perp f_1^a(x, \mathbf{k}_\perp^2; \mu^2) D_1^{a \rightarrow h}(z, \mathbf{P}_\perp^2; \mu^2) \delta^{(2)}(z \mathbf{k}_\perp - \mathbf{P}_{hT} + \mathbf{P}_\perp)} \\ + Y_{UU,T}(Q^2, \mathbf{P}_{hT}^2) + \mathcal{O}(M^2/Q^2)$$

- The **W term** dominates in the region where $q_T \ll Q$
- The Y term has been excluded in the MAP analysis

W Term

TMD formalism: factorization

DY



In $q_T^2 \ll Q^2$ and $M^2 \ll Q^2$ region:

$$F_{UU}^1(x_A, x_B, \mathbf{q}_T, Q) = x_A x_B \mathcal{H}^{DY}(Q; \mu) \sum_a c_a(Q^2) \int d|\mathbf{b}_T| |\mathbf{b}_T| J_0(|\mathbf{q}_T| |\mathbf{b}_T|) \hat{f}_1^a(x_A, \mathbf{b}_T^2; \mu, \zeta_A) \hat{f}_1^b(x_B, \mathbf{b}_T^2; \mu, \zeta_B)$$

W term

TMD factorization: TMD components

TMD in Fourier space

$$\hat{F}(x, b_T^2; \mu, \zeta) = \int \frac{d^2 \mathbf{k}_\perp}{(2\pi)^2} e^{i \mathbf{b}_T \cdot \mathbf{k}_\perp} F(x, k_\perp^2; \mu, \zeta)$$

$$\hat{f}_1^q(x, b_T^2; \mu, \zeta) = \sum_j C_{q/j}(x, b_*; \mu_{b_*}, \mu_{b_*}^2) \otimes f_1^j(x, \mu_{b_*})$$

Collinear extractions

Perturbative : A

$$\times \exp \left\{ K(b_*; \mu_{b_*}) \ln \frac{\sqrt{\zeta}}{\mu_{b_*}} + \int_{\mu_{b_*}}^\mu \frac{d\mu'}{\mu'} \left[\gamma_F - \gamma_K \ln \frac{\sqrt{\zeta}}{\mu'} \right] \right\}$$

Evolution to final energy scale of the process

: B

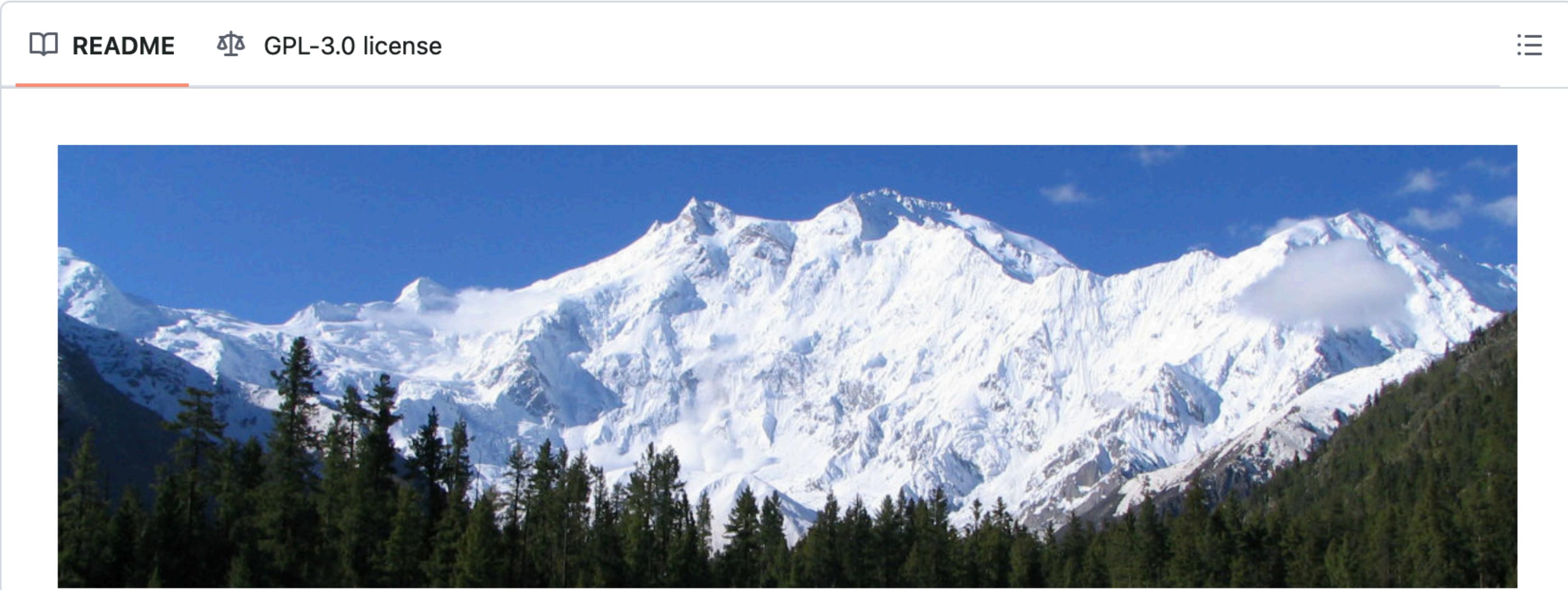
$$\times f_{NP}(x, b_T^2) \exp \left\{ g_K(b_T^2) \ln \frac{\sqrt{\zeta}}{\sqrt{\zeta_0}} \right\}$$

Non-perturbative part of the TMD

Parametrization

TO BE FITTED

Nanga Parbat: a MAP fit framework



Nanga Parbat: a TMD fitting framework

Nanga Parbat is a fitting framework aimed at the determination of the non-perturbative component of TMD distributions.

Download

You can obtain NangaParbat directly from the github repository:

<https://github.com/MapCollaboration/NangaParbat>

Available global fits

DY + SIDIS

	Accuracy	Ndata	$\chi^2/Ndata$
PV 17 <small>Bacchetta, Delcarro et al. JHEP 06 (2017)</small>	NLL	8059	1.55
SV 2019 <small>Scimemi, Vladimirov JHEP 06 (2020)</small>	N^3LL^-	1039	1.06
MAPTMD22 <small>Bacchetta, Bertone et al. JHEP 10 (2022)</small>	N^3LL^-	2031	1.06

Flavor independent

Our starting point: MAPTMD22 FI global fit

Flavor independent

Global fit: DY + SIDIS

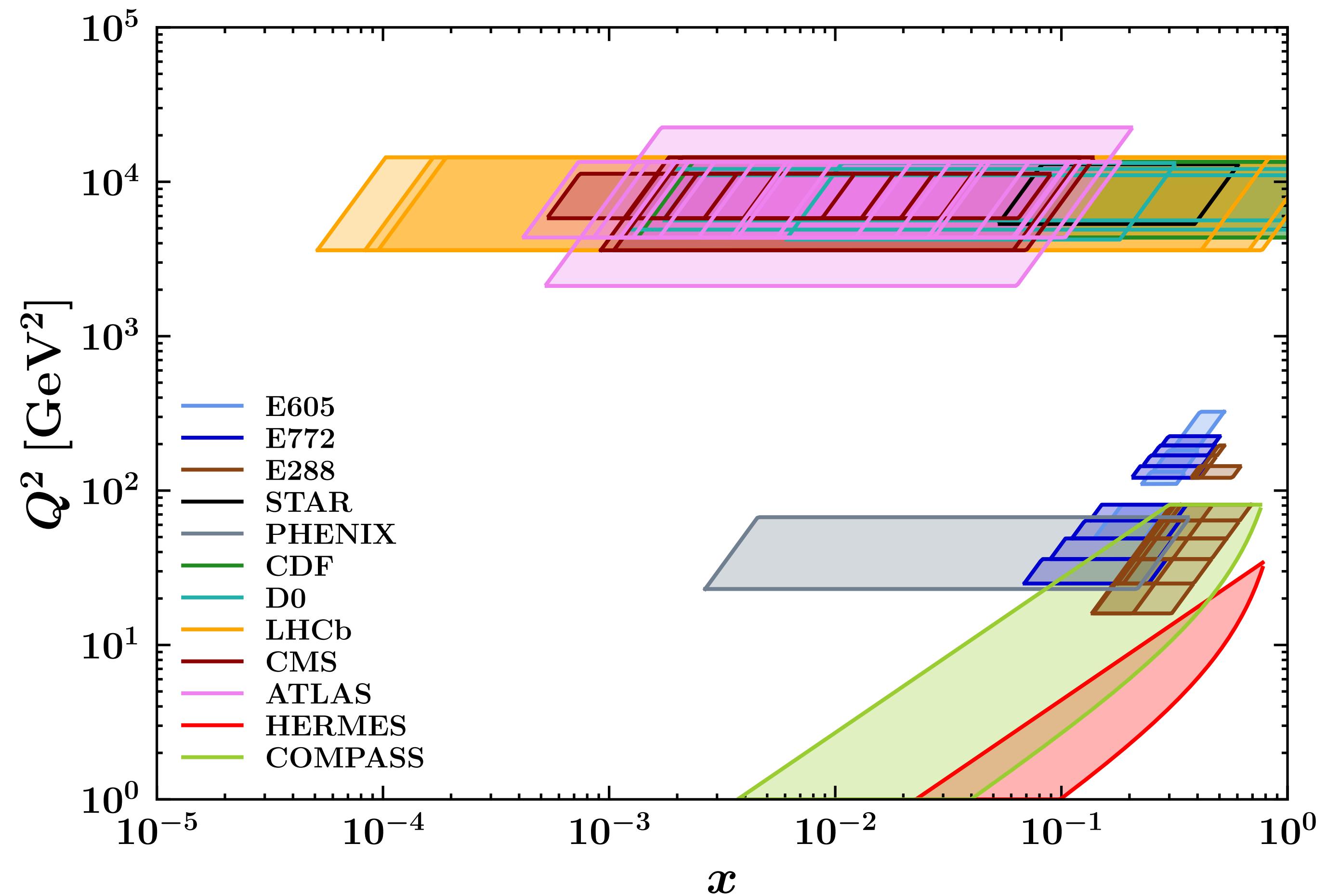
2031 data points

DY data

484

SIDIS data

1547



MAPTMD22 FI global fit

2031 data points

Perturbative accuracy: N³LL-

$$L = \ln \left(\frac{\mu^2}{\mu_b^2} \right)$$

Accuracy	H and C	K and γ_F	γ_K	PDFs/FFs and as evol.
LL	0	-	1	-
NLL	0	1	2	LO
NLL'	1	1	2	NLO
NNLL	1	2	3	NLO
NNLL'	2	2	3	NNLO
N ³ LL-	2	3	4	NNLO + NLO
N ³ LL	2	3	4	NNLO
N ³ LL'	3	3	4	N ³ LO

[Bacchetta, Bertone, Bissolotti, et al., JHEP 07 (2020)]

TMD handbook, Boussarie, et al., 2023

PDF: MMHT2014nnlo
FF: DSS14-17 NLO

MAPTMD22 parametrisation

$$f_{NP}(x, b_T^2) \exp \left\{ g_K(b_T^2) \ln \frac{\sqrt{\zeta}}{\sqrt{\zeta_0}} \right\}$$

$$f_{1NP}(x, b_T^2) \propto \text{F.T. of} \left(e^{-\frac{k_\perp^2}{g^{1A}}} + \lambda_B k_\perp^2 e^{-\frac{k_\perp^2}{g^{1B}}} + \lambda_C e^{-\frac{k_\perp^2}{g^{1C}}} \right)$$

$$D_{1NP}(x, b_T^2) \propto \text{F.T. of} \left(e^{-\frac{P_\perp^2}{g_{3A}}} + \lambda_{FB} k_\perp^2 e^{-\frac{P_\perp^2}{g_{3B}}} \right)$$

$$g_K(b_T^2) = -g_2^2 \frac{b_T^2}{4}$$

$$g_1(x) = N_1 \frac{(1-x)^\alpha}{(1-\hat{x})^\alpha} \frac{x^\sigma}{\hat{x}^\sigma}$$

$$g_3(z) = N_3 \frac{(z^\beta + \delta)(1-z)^\gamma}{(\hat{z}^\beta + \delta)(1-\hat{z})^\gamma}$$

11 parameters for TMD PDF
 + 1 for NP evolution + 9 for TMD FF
 = 21 free parameters

MAPTMD22 summary

- Global fit of DY and SIDIS data: **2031** data points
- ***Normalization*** of SIDIS multiplicities beyond NLL
- Number of fitted parameters: **21**
- Perturbative accuracy: **N^3LL-**

Really good description:

$$\chi^2/N_{\text{data}} = 1.06$$

MAPTMD22 improvement: MAPTMD24

- Global fit of DY and SIDIS data: **2031** data points → Same data sets
- **Normalization** of SIDIS multiplicities beyond NLL → Same approach
- Number of fitted parameters: **21** → Same parametrisation
(Still flavour independent)

MAPTMD22 improvement: MAPTMD24

- Global fit of DY and SIDIS data: **2031** data points → Same data sets
- **Normalization** of SIDIS multiplicities beyond NLL → Same approach
- Number of fitted parameters: **21** → Same parametrisation
- Perturbative accuracy: **full N^3LL** $\xrightarrow{\text{PDF: MMHT2014nnlo}} \text{NNPDF31NNLO}$
 $\xrightarrow{\text{FF: DSS14-17}} \text{MAPFF10NNLO}$

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MAPFF10NNLO**
- Not as good description: $\chi^2/N_{\text{data}} = 1.40$

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*PDF: MMHT2014nnlo
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MAPFF10NNLO**
- Not as good description: $\chi^2/N_{\text{data}} = 1.40$
Improved framework but worse description. WHY?

MAPTMD* χ^2 comparison

TMD extraction	collinear configuration	χ^2 / N_{data}		
		DY	SIDIS	Total
MAPTMD22	MMHT+DSS	1.66	0.87	1.06
MAPTMD24 FI	NNPDF+MAPFF	1.58	1.34	1.40

MAPTMD* x^2 comparison

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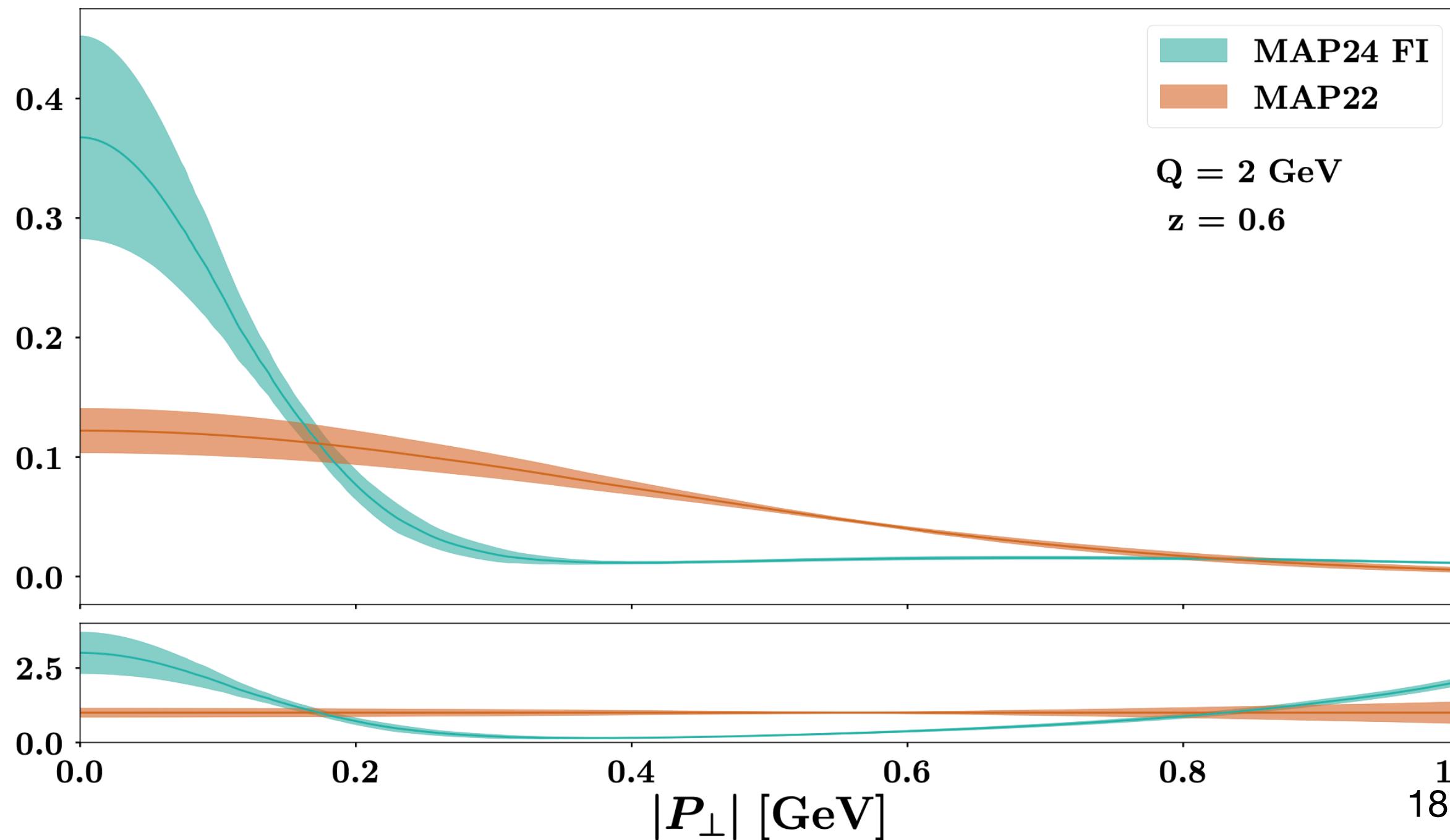
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SIDIS: convolution of TMD PDF x **FF**

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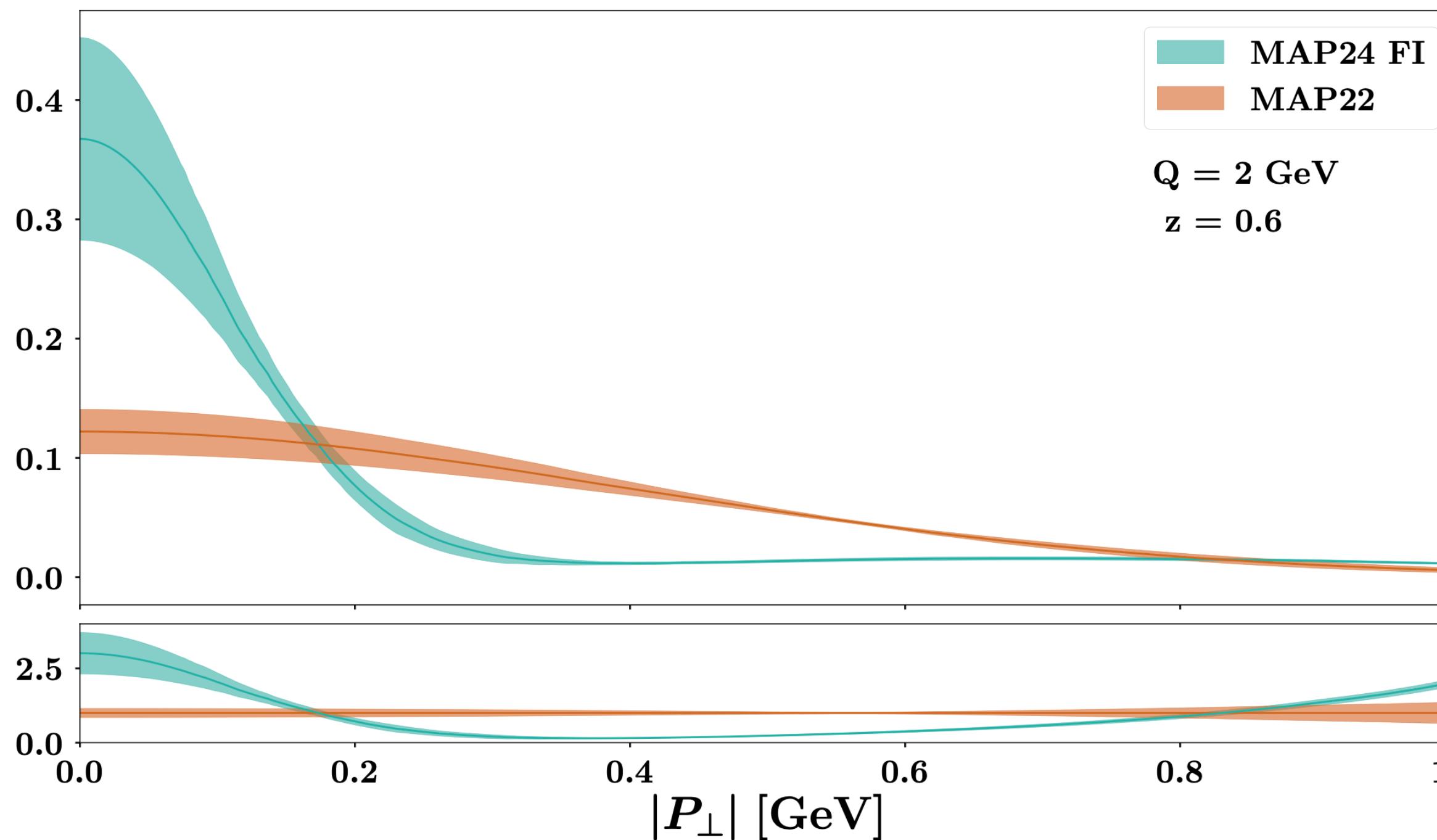
MAPFF1.0nnlo

- NNLO
- NN approach
- new behavior
- smaller uncertainties

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MAPTMD24 collinear configurations

Data set	N_{dat}	χ^2_0/N_{dat}
DY collider total	251	2.14
Dy fixed target total	233	0.68
HERMES total	344	2.72
COMPASS total	1203	0.99
SIDIS total	1547	1.38
Total	2031	1.40

NNPDF + MAPFF

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NNPDF + MAPFF

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Similar
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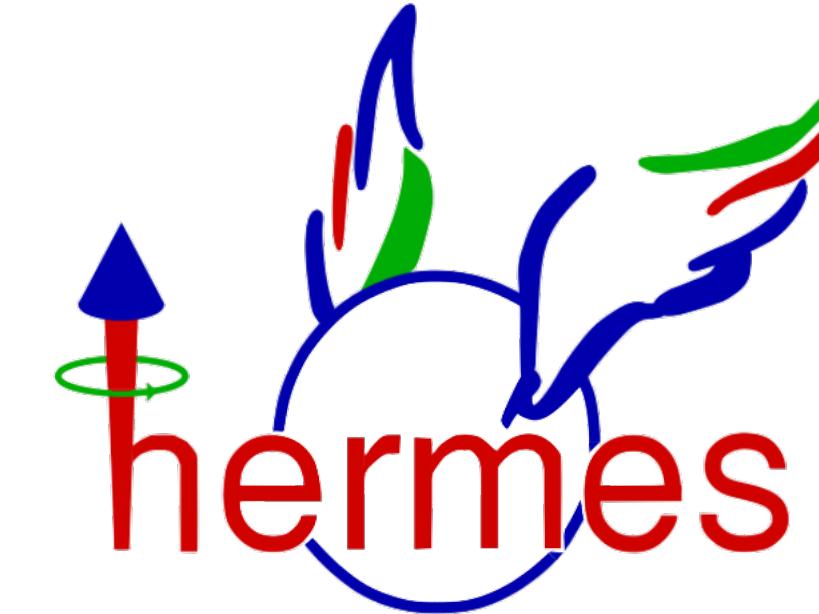
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DY collider total	251	2.14
Dy fixed target total	233	0.68
HERMES total	344	2.72
COMPASS total	1203	0.99
SIDIS total	1547	1.38
Total	2031	1.39

NNPDF + MAPFF



Data set	N_{dat}	χ^2_0/N_{dat}
DY collider total	251	2.40
Dy fixed target total	233	0.75
HERMES total	344	0.95
COMPASS total	1203	0.88
SIDIS total	1547	0.90
Total	2031	1.07

NNPDF + DSS

Data set	N_{dat}	χ^2_0/N_{dat}
DY collider total	251	2.01
Dy fixed target total	233	1.11
HERMES total	344	2.51
COMPASS total	1203	0.99
SIDIS total	1547	1.33
Total	2031	1.39

MMHT + MAPFF

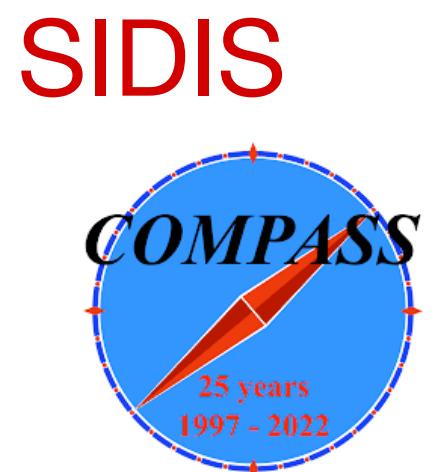
Data set	N_{dat}	χ^2_0/N_{dat}
DY collider total	251	2.06
Dy fixed target total	233	1.24
HERMES total	344	0.71
COMPASS total	1203	0.92
SIDIS total	1547	0.87
Total	2031	1.06

MMHT + DSS (MAP22)

MAPTMD24: included processes



$$\begin{aligned} e + p &\rightarrow e' + \pi^+ + X \\ e + p &\rightarrow e' + \pi^- + X \\ e + p &\rightarrow e' + K^+ + X \\ e + p &\rightarrow e' + K^- + X \end{aligned} \quad + D \text{ target}$$

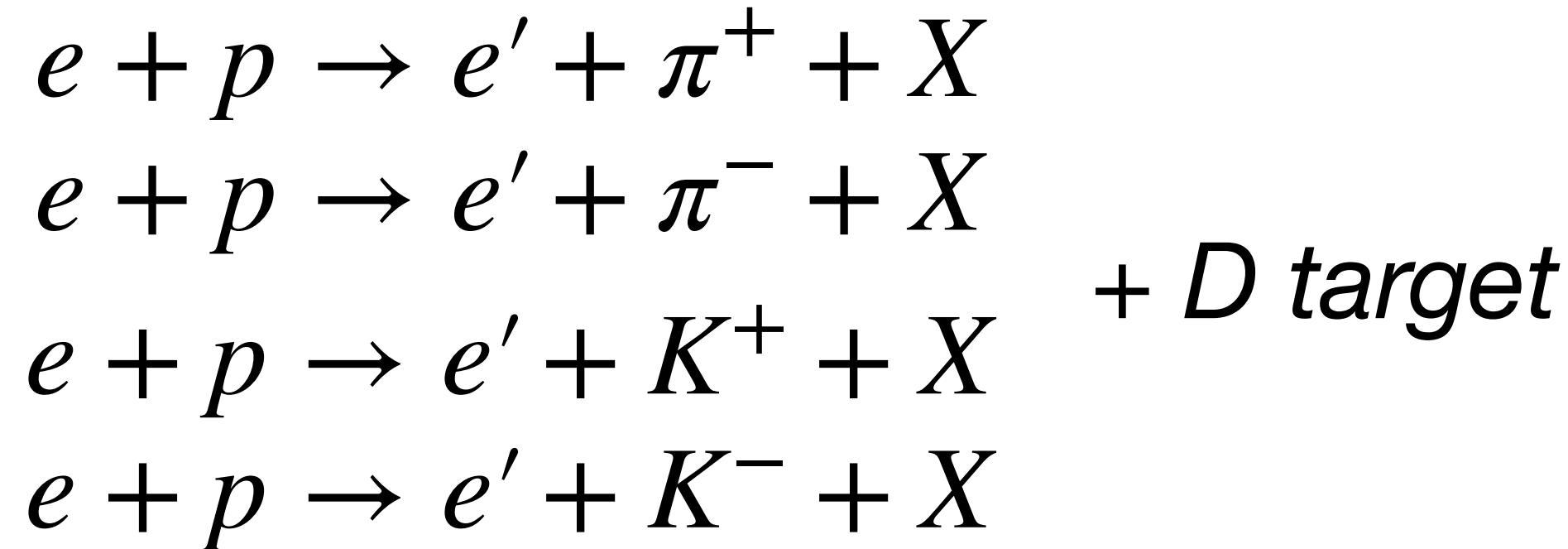


D target, unidentified final state h

Drell-Yan

$q\bar{q}$ in the initial state

MAPTMD24: included processes



HIGH sensitivity to
flavor dependence



D target, unidentified final state h

LOW sensitivity to
flavor dependence

Drell-Yan

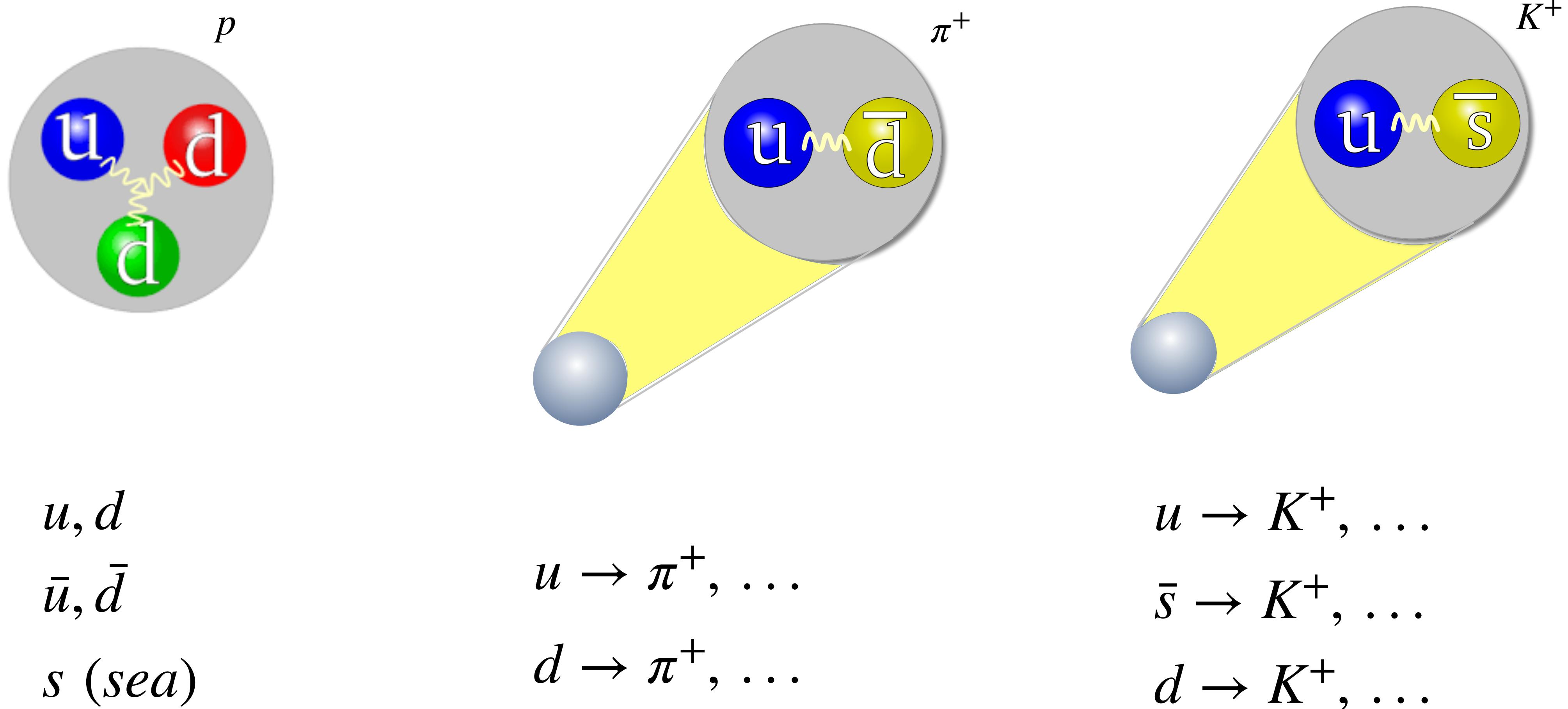
q̄q in the initial state

MAPTMD24: need for flavor

Statement: introducing new FFs highlighted a tension between theory and SIDIS data.

A possible solution is to introduce flavor dependence

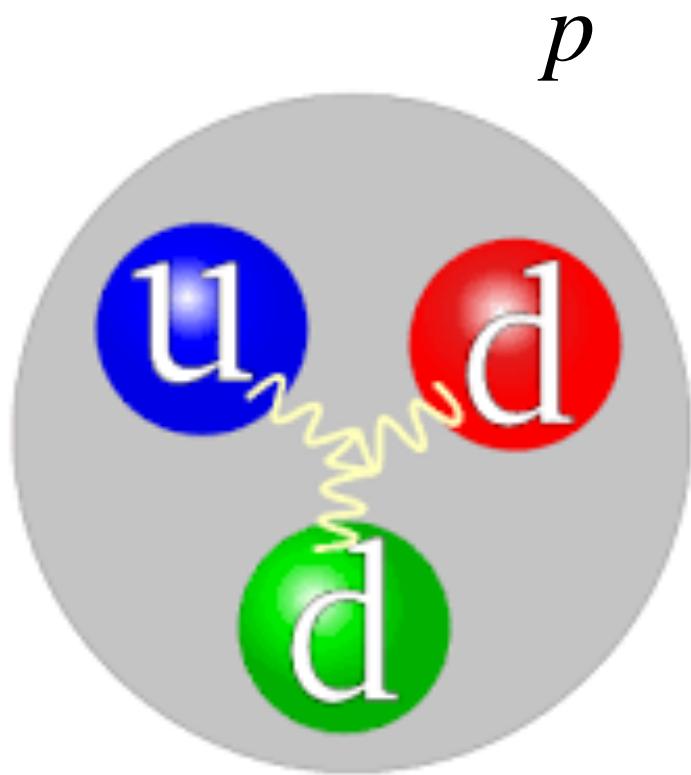
MAPTMD24 flavor parametrization



Negative fragmenting mesons: charge conjugation

MAPTMD24 flavor parametrization

Same parametrisation as MAPTMD22 but...



PDF: 5 sets of parameters

$u, d, \bar{u}, \bar{d}, sea$

FF: 5 sets of parameters

favored and unfavored pion fragmentation;

favored, unfavored and s-quark kaon fragmentation

(10x5) parameters for TMD PDF
+ (9x5) for TMD FF
+ 1 for NP evolution
= **96** free parameters

MAPTMD24 extraction - Results

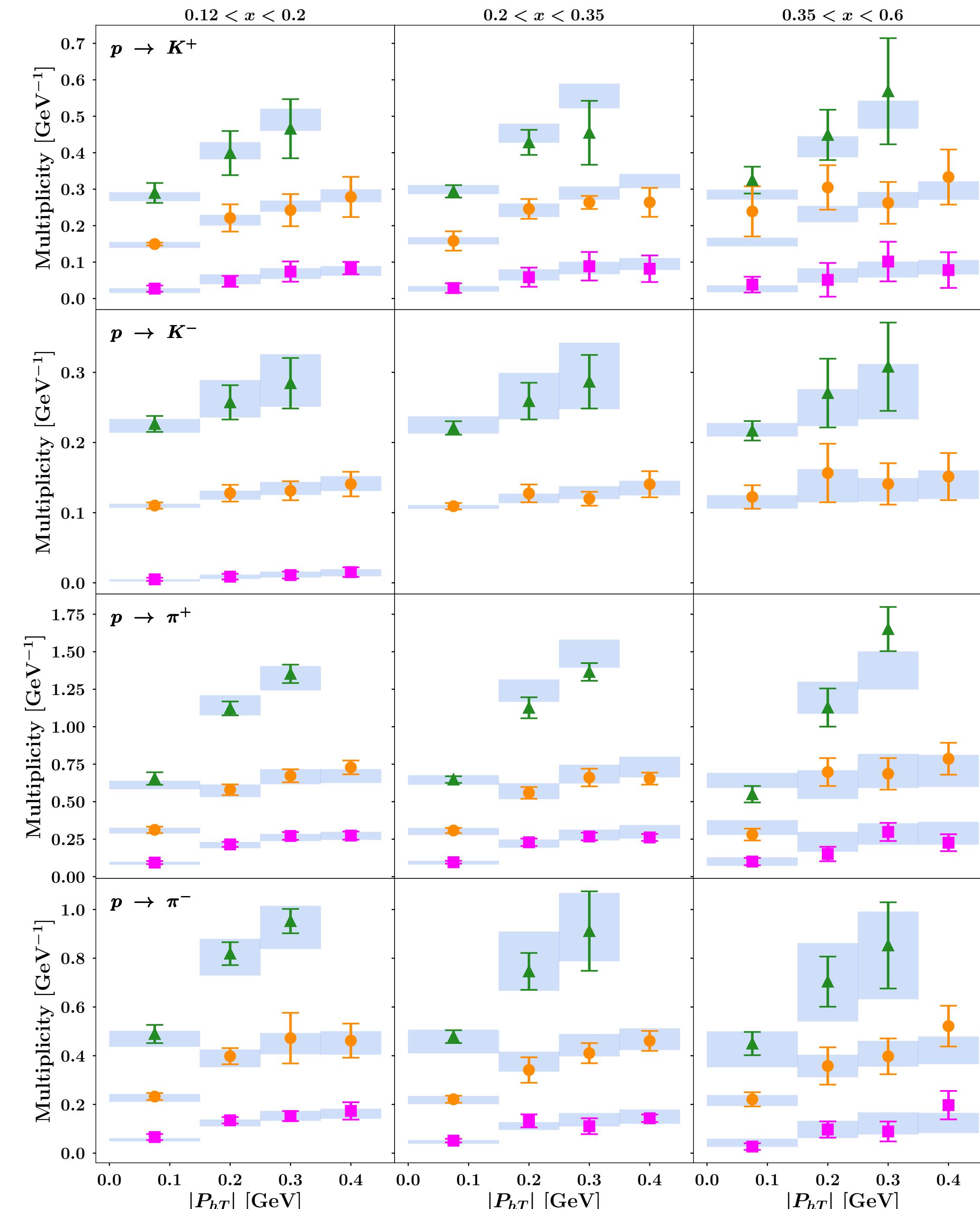
Data set	N ³ LL			
	N_{dat}	χ^2_D	χ^2_λ	χ^2_0
DY collider total	251	1.37	0.28	1.65
DY fixed-target total	233	0.63	0.31	0.94
<i>HERMES total</i>	344	0.81	0.24	1.05
<i>COMPASS total</i>	1203	0.67	0.27	0.94
SIDIS total	1547	0.70	0.26	0.96
Total	2031	0.81	0.27	1.08

$$\chi^2/N_{\text{data}} = 1.08$$

$$\chi^2_0 = \chi^2_D + \chi^2_\lambda$$

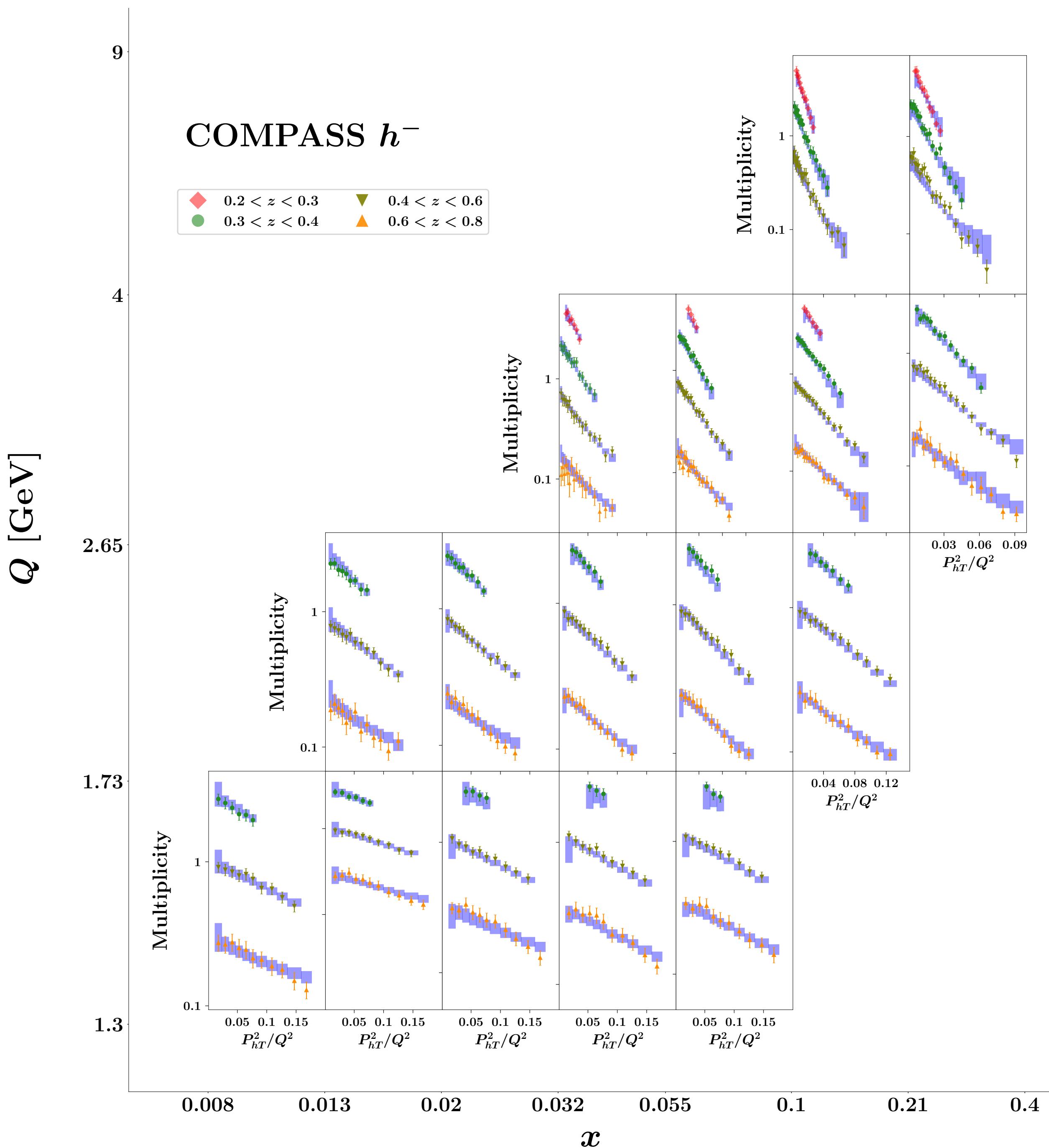
MAPTMD24 - Results

	N ³ LL			
Data set	N_{dat}	χ^2_D	χ^2_λ	χ^2_0
DY collider total	251	1.37	0.28	1.65
DY fixed-target total	233	0.63	0.31	0.94
<i>HERMES total</i>	344	0.81	0.24	1.05
COMPASS total	1203	0.67	0.27	0.94
SIDIS total	1547	0.70	0.26	0.96
Total	2031	0.81	0.27	1.08



MAPTMD24 - Results

	N ³ LL			
Data set	N_{dat}	χ^2_D	χ^2_λ	χ^2_0
DY collider total	251	1.37	0.28	1.65
DY fixed-target total	233	0.63	0.31	0.94
<i>HERMES total</i>	344	0.81	0.24	1.05
<i>COMPASS total</i>	1203	0.67	0.27	0.94
SIDIS total	1547	0.70	0.26	0.96
Total	2031	0.81	0.27	1.08



MAPTMD24 - Results

	N ³ LL			
Data set	N_{dat}	χ^2_D	χ^2_λ	χ^2_0
DY collider total	251	1.37	0.28	1.65
DY fixed-target total	233	0.63	0.31	0.94
<i>HERMES total</i>	344	0.81	0.24	1.05
<i>COMPASS total</i>	1203	0.67	0.27	0.94
SIDIS total	1547	0.70	0.26	0.96
Total	2031	0.81	0.27	1.08



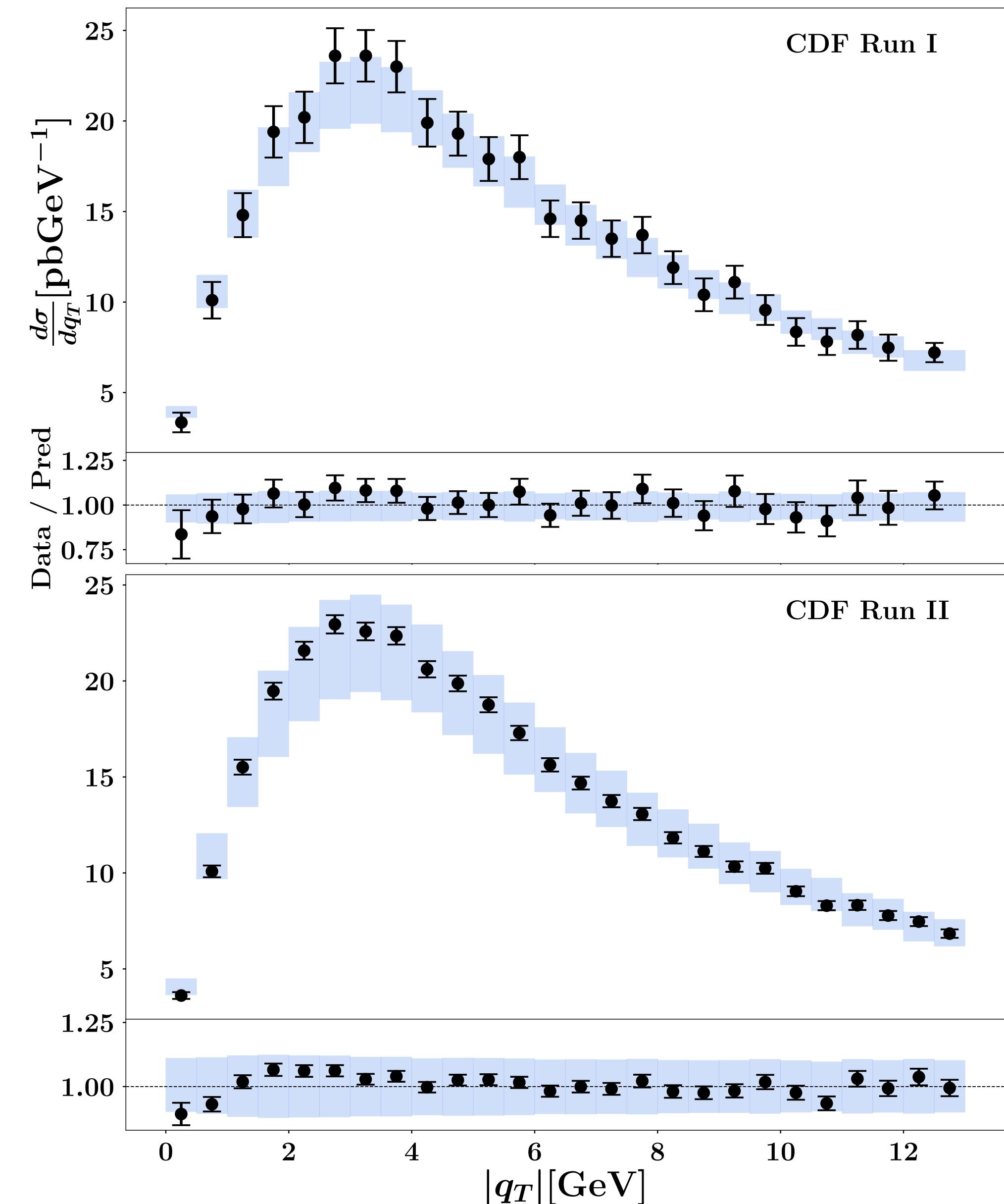
SIDIS data: really good agreement

MAPTMD24 - Results

N ³ LL				
Data set	N_{dat}	χ^2_D	χ^2_λ	χ^2_0
DY collider total	251	1.37	0.28	1.65
DY fixed-target total	233	0.63	0.31	0.94
<i>HERMES total</i>	344	0.81	0.24	1.05
<i>COMPASS total</i>	1203	0.67	0.27	0.94
SIDIS total	1547	0.70	0.26	0.96
Total	2031	0.81	0.27	1.08

E288 E772 E605

DY fixed: still really good agreement



MAPTMD24 - Results

	N ³ LL			
Data set	N_{dat}	χ^2_D	χ^2_λ	χ^2_0
DY collider total	251	1.37	0.28	1.65
DY fixed-target total	233	0.63	0.31	0.94
<i>HERMES total</i>	344	0.81	0.24	1.05
<i>COMPASS total</i>	1203	0.67	0.27	0.94
SIDIS total	1547	0.70	0.26	0.96
Total	2031	0.81	0.27	1.08

Tevatron

LHCb

CMS

STAR 510

Atlas

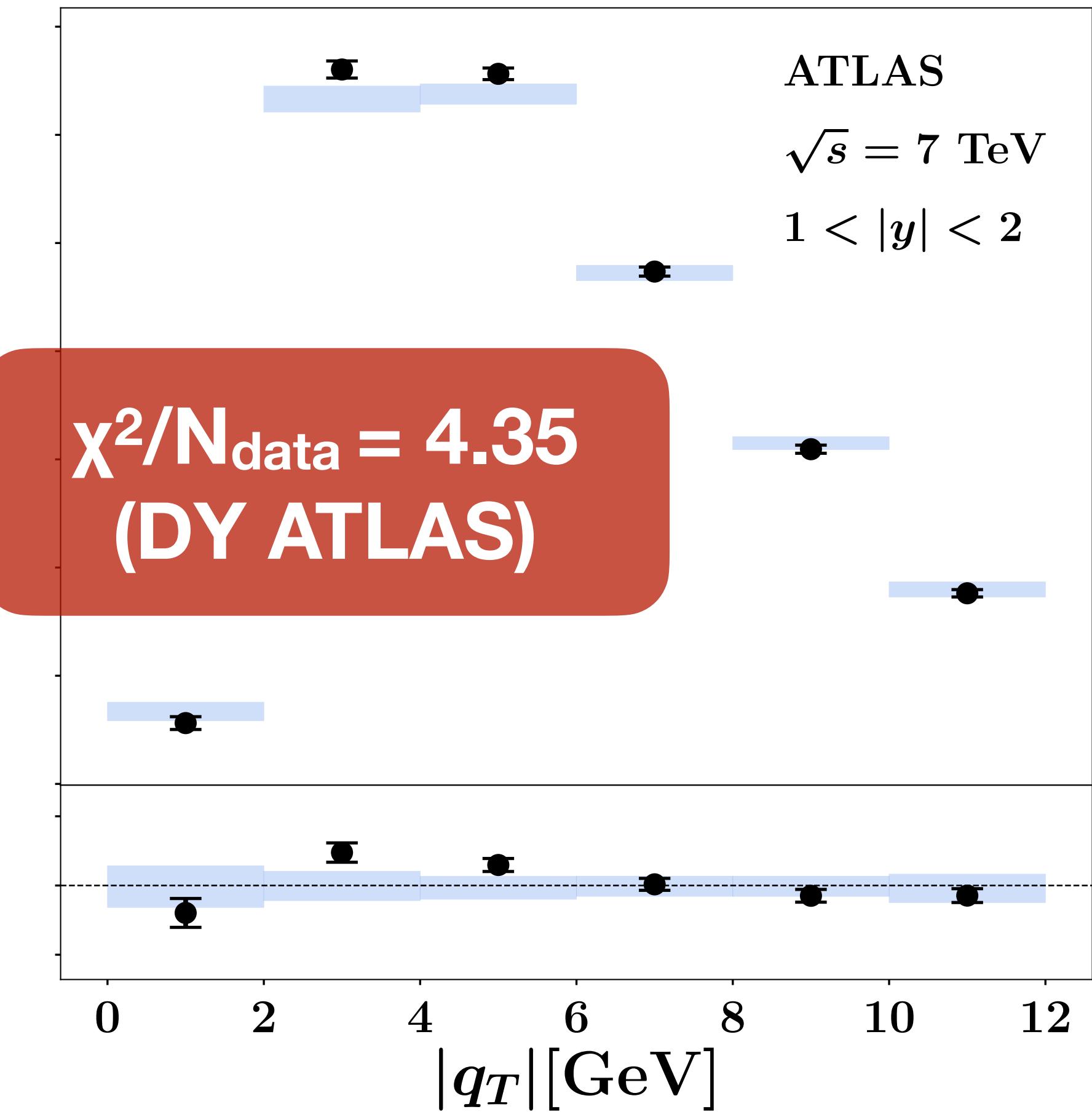
PHENIX 200

DY collider: quite good agreement

MAPTMD24 - Results

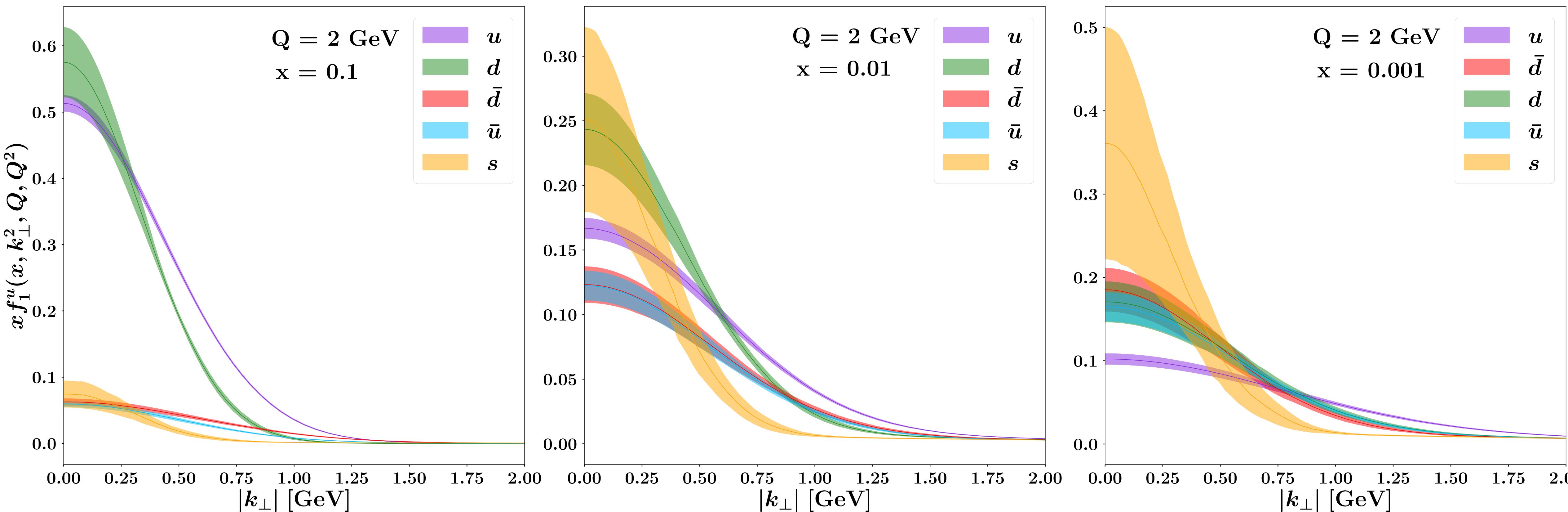
	N ³ LL			
Data set	N_{dat}	χ^2_D	χ^2_λ	χ^2_0
DY collider total	251	1.37	0.28	1.65
DY fixed-target total	233	0.63	0.31	0.94
<i>HERMES total</i>	344	0.81	0.24	1.05
<i>COMPASS total</i>	1203	0.67	0.27	0.94
SIDIS total	1547	0.70	0.26	0.96
Total	2031	0.81	0.27	1.08

DY collider: quite good agreement

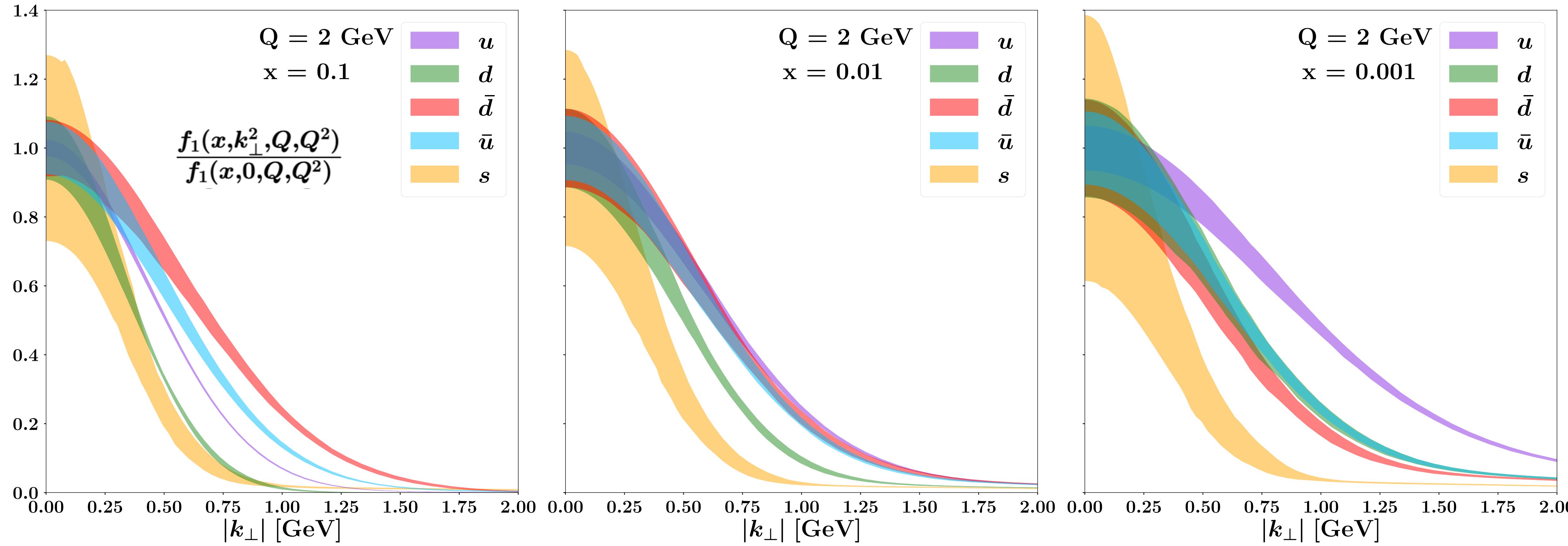


MAPTMD24 - Results

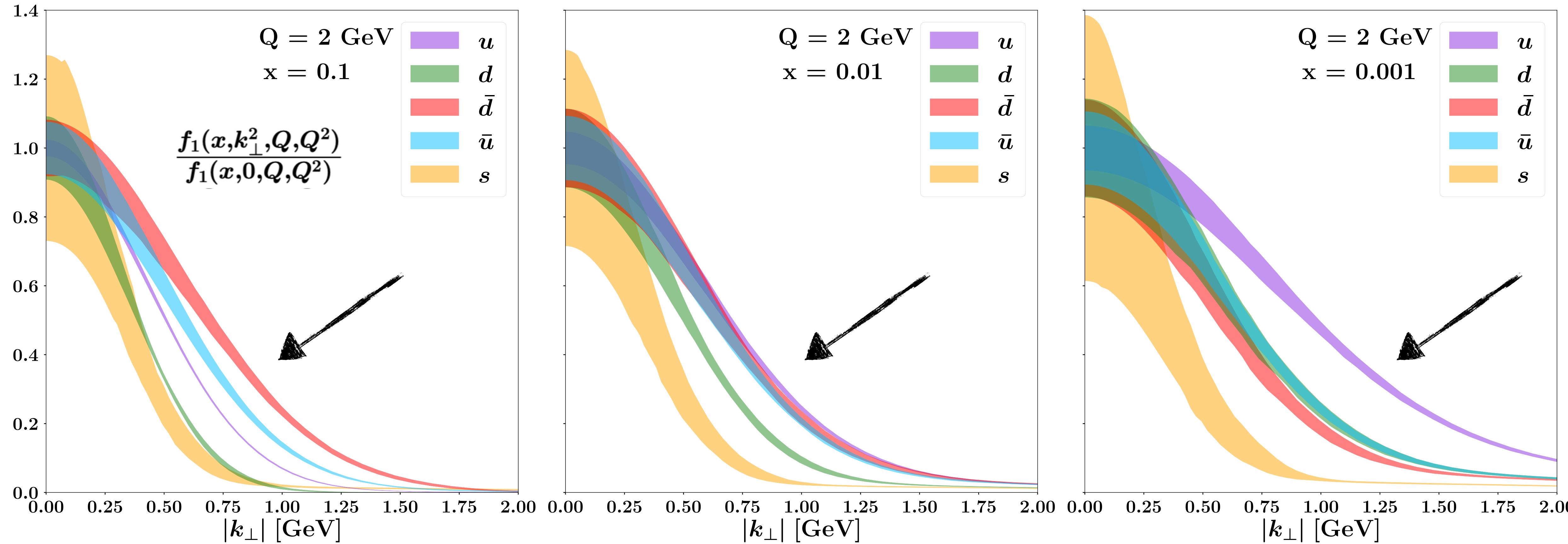
Flavor-dependent TMD PDFs



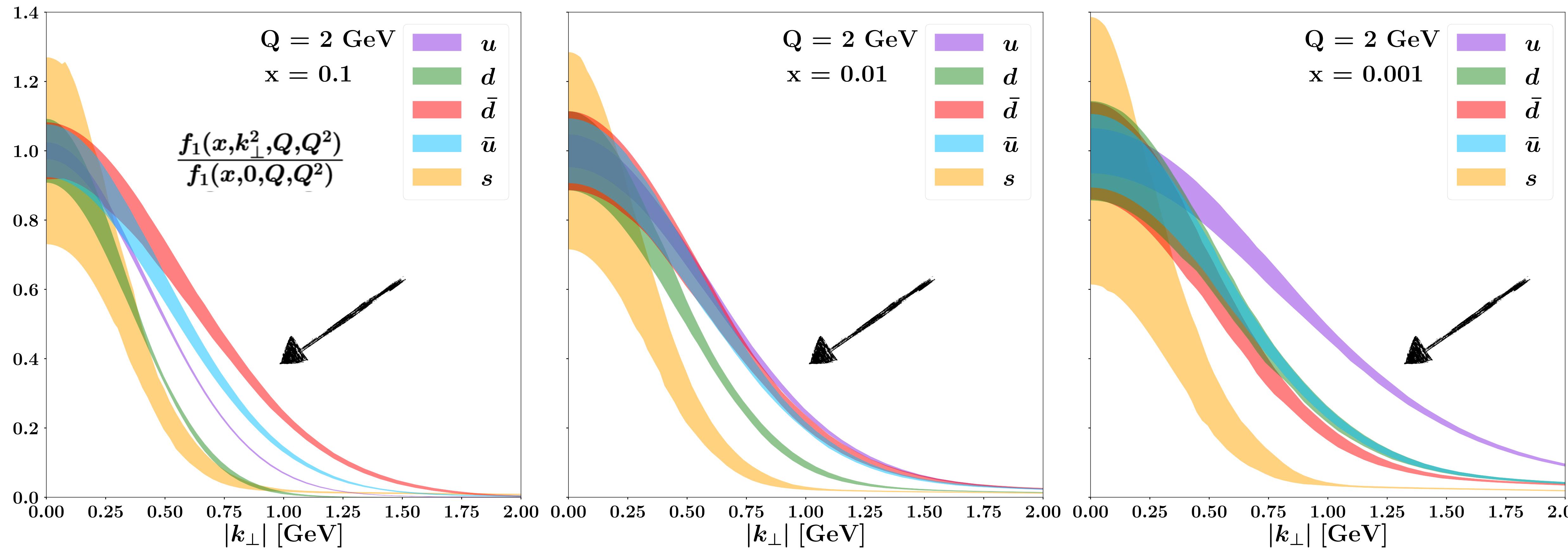
MAPTMD24 - TMD PDF



MAPTMD24 - TMD PDF

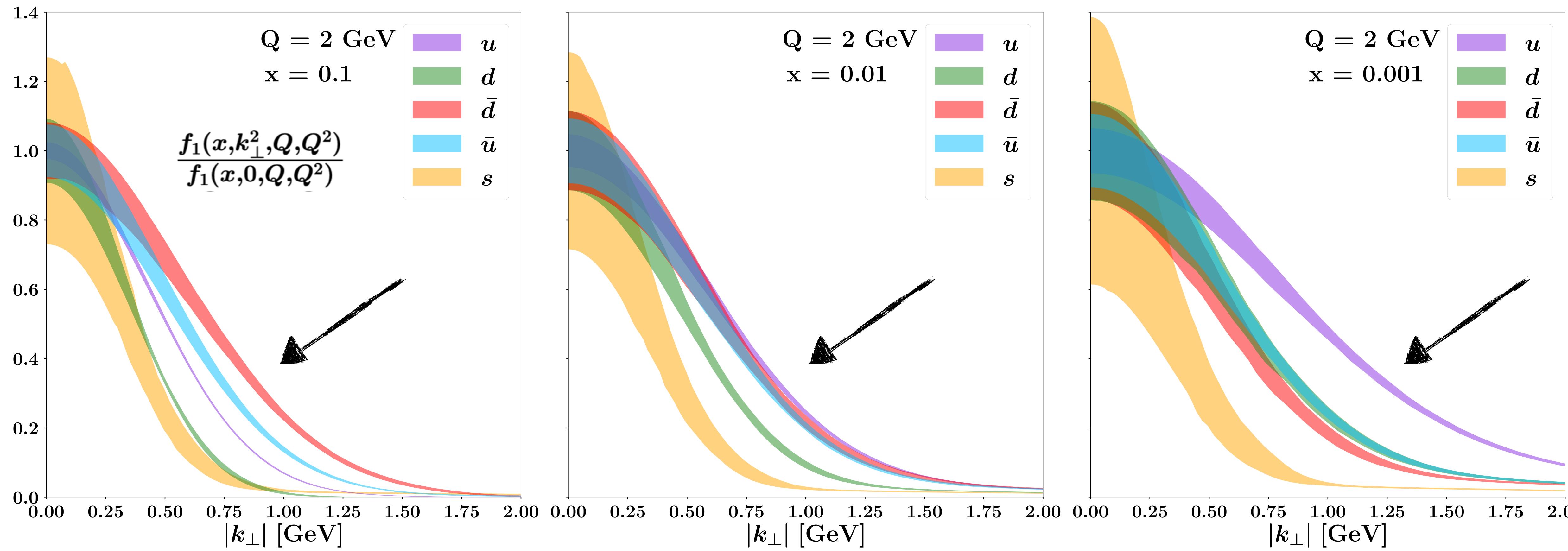


MAPTMD24 - TMD PDF



Very different k_\perp - behaviours!

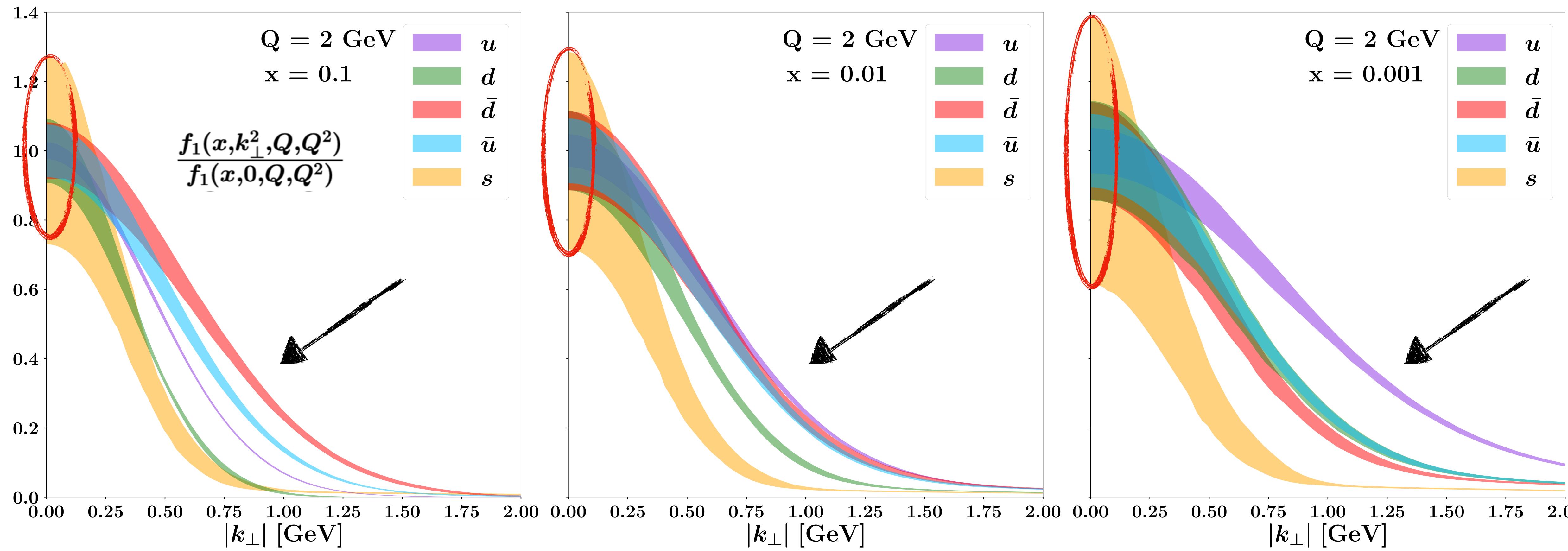
MAPTMD24 - TMD PDF



Very different k_\perp - behaviours!

It also changes with x

MAPTMD24 - TMD PDF

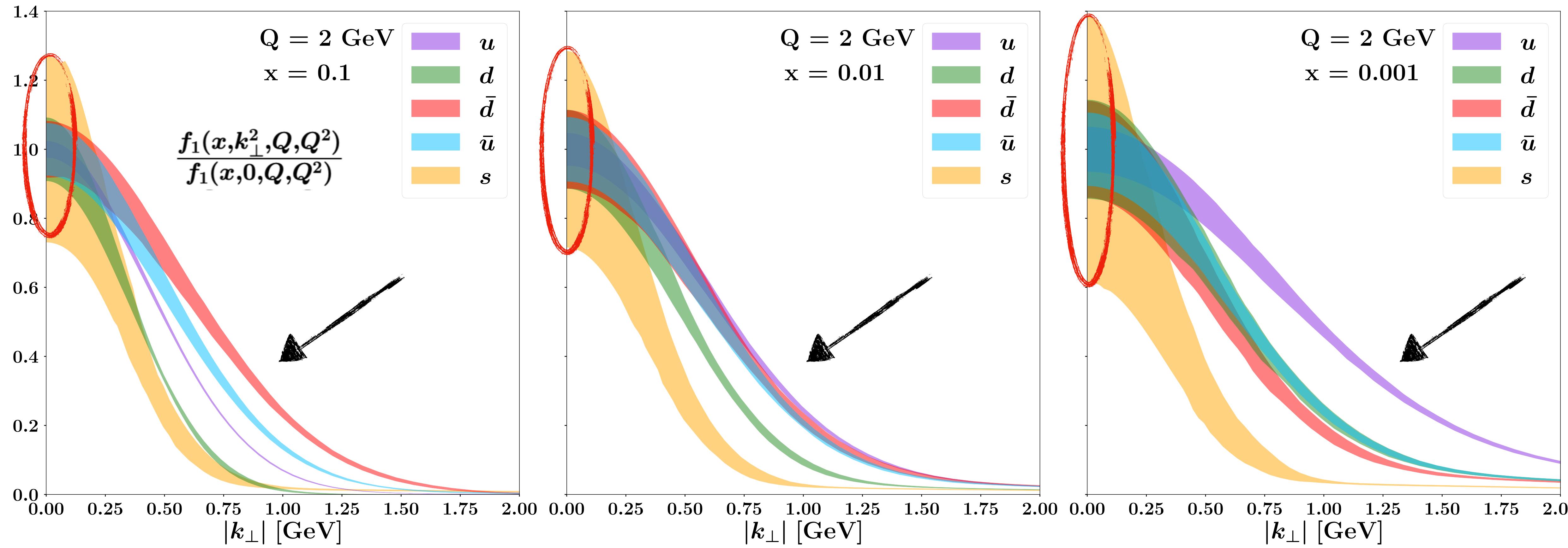


Very different k_\perp - behaviours!

It also changes with x

MAPTMD24 - TMD PDF

The sea is the least constrained

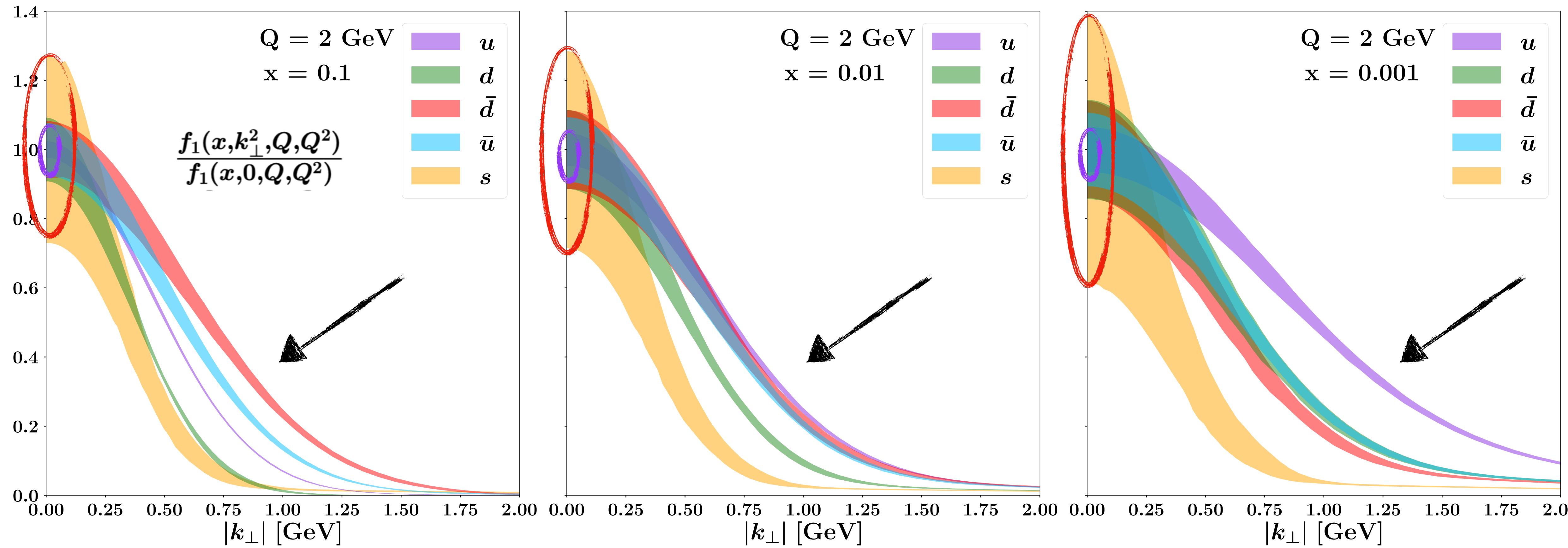


Very different k_\perp - behaviours!

It also changes with x

MAPTMD24 - TMD PDF

The sea is the least constrained



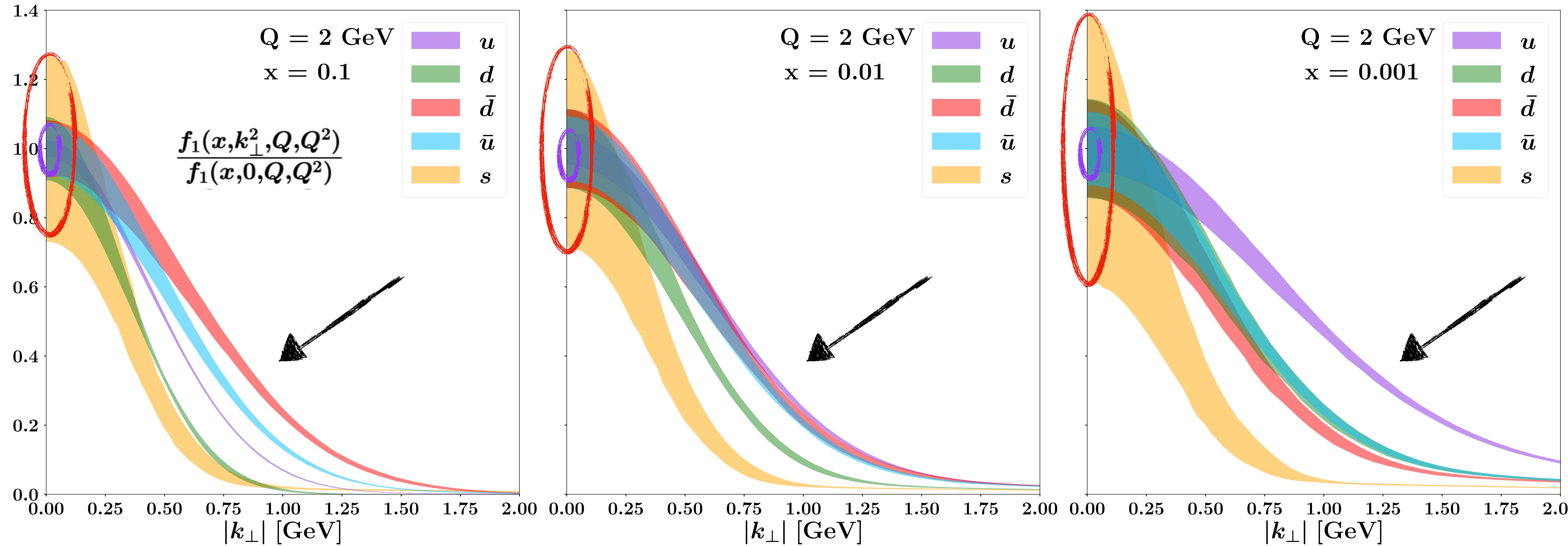
Very different k_\perp - behaviours!

It also changes with x

MAPTMD24 - TMD PDF

The sea is the least constrained

The up quark is the most constrained

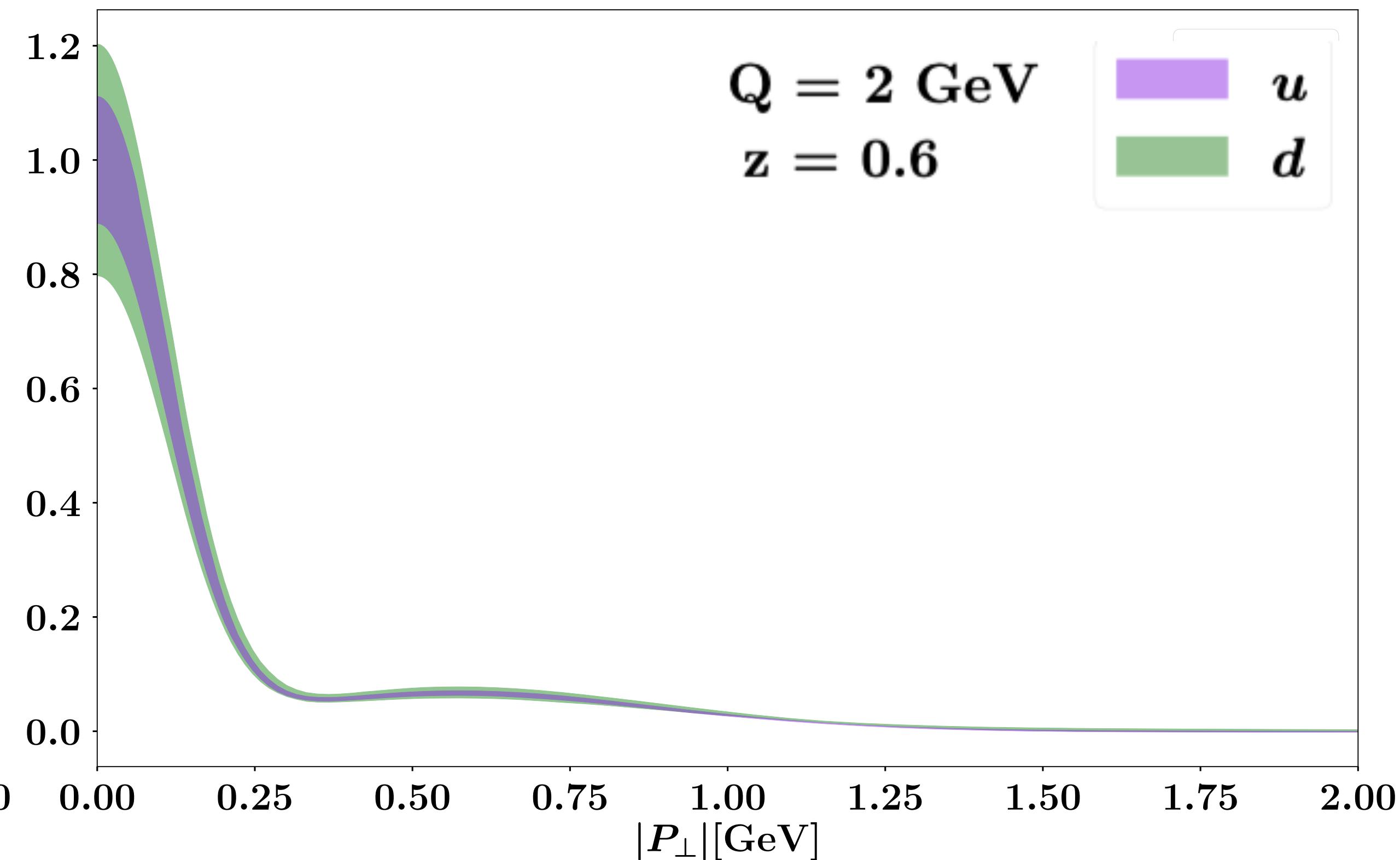
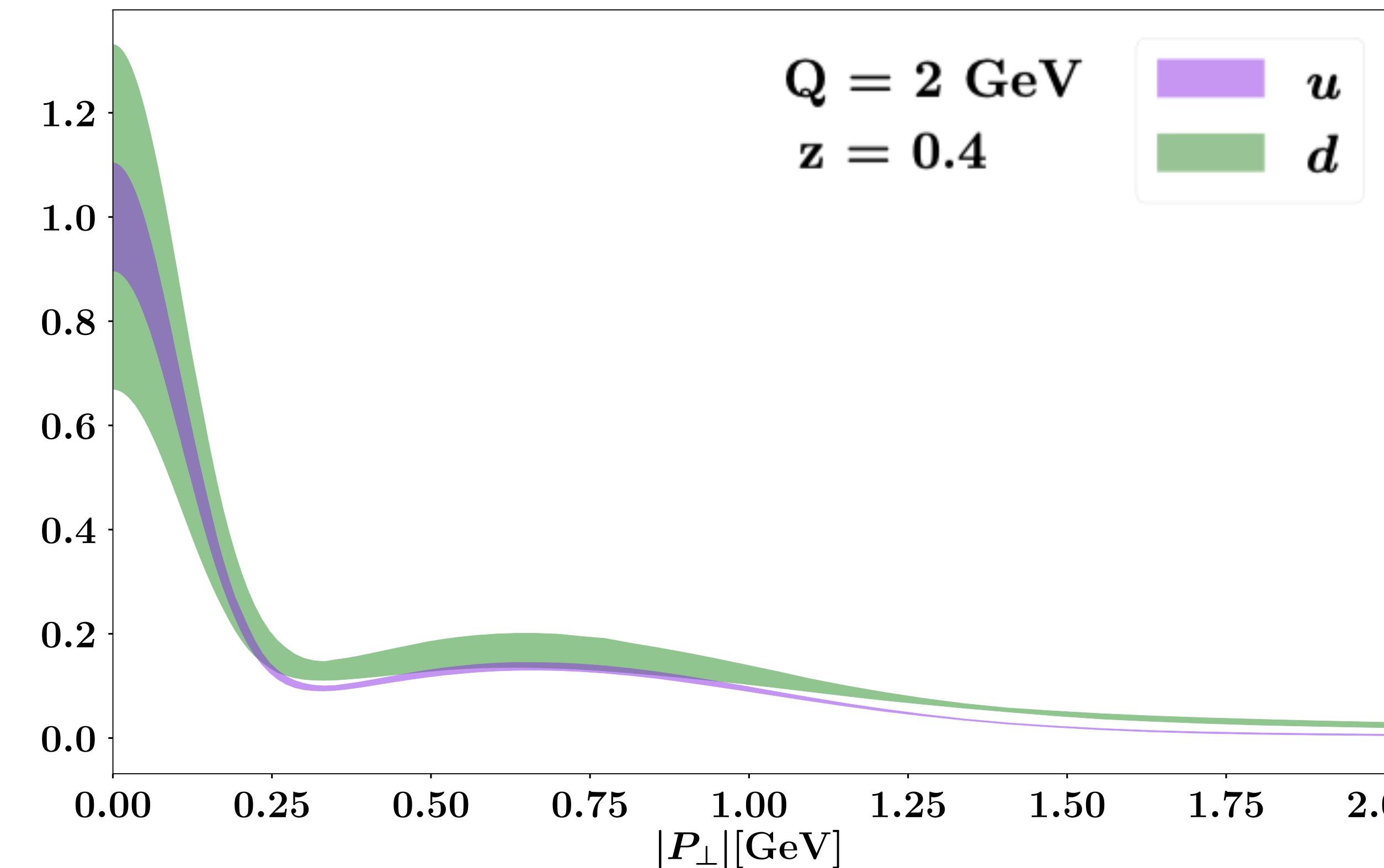


Very different k_{\perp} - behaviours!

It also changes with x

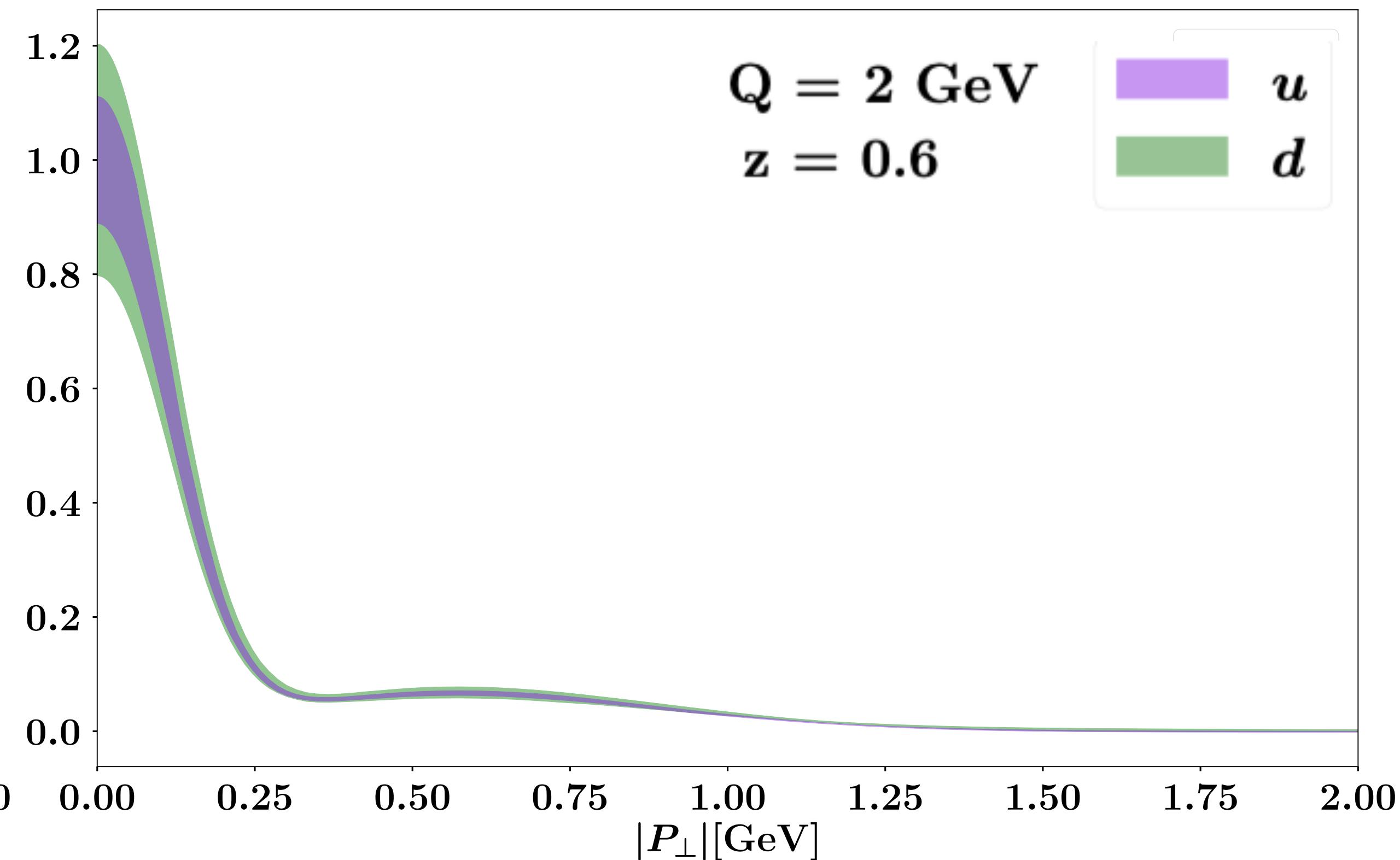
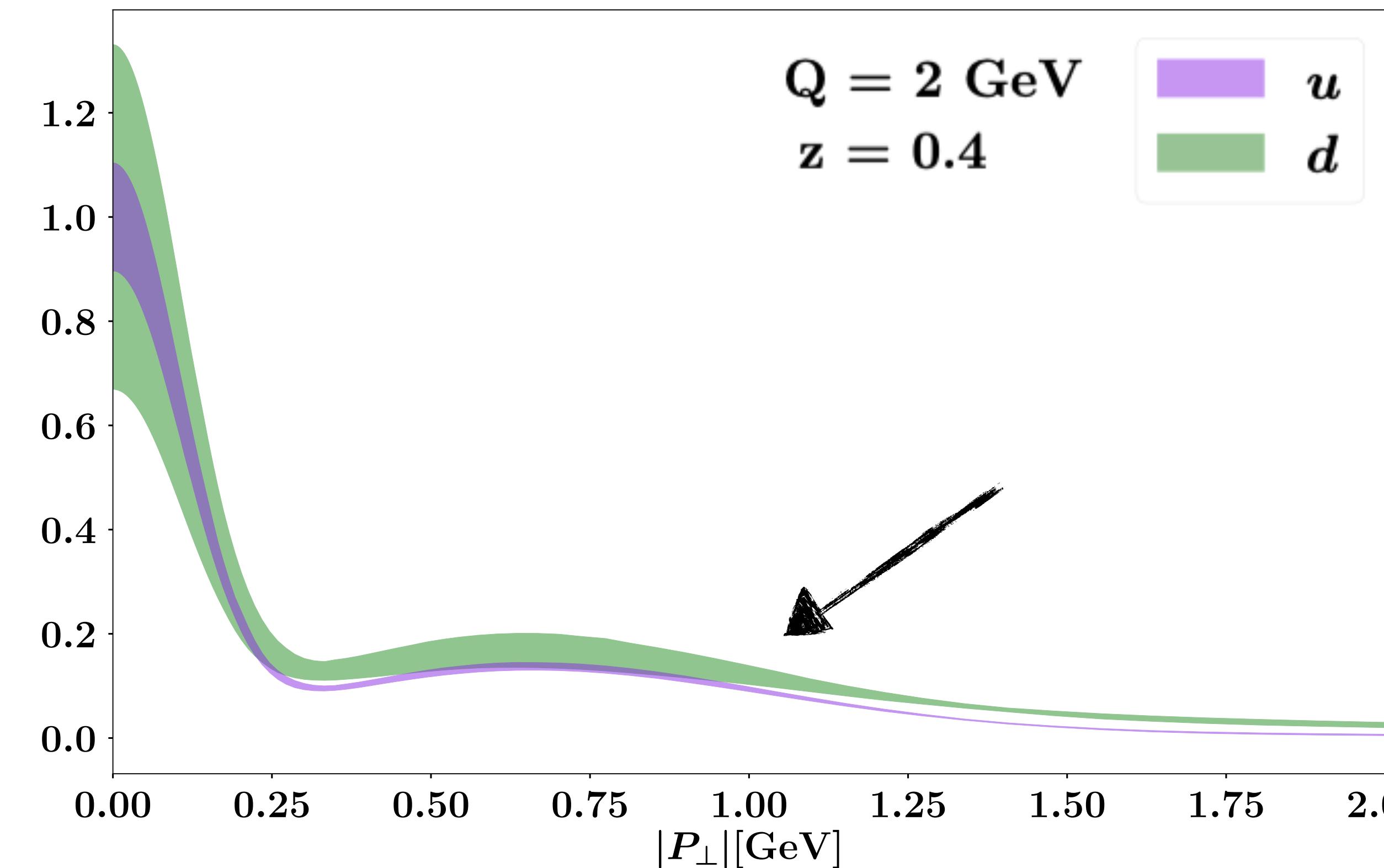
MAPTMD24 - TMD FF

$$\frac{D_1 \rightarrow \pi^+(z, P_\perp^2, Q, Q^2)}{D_1 \rightarrow \pi^+(z, 0, Q, Q^2)}$$



MAPTMD24 - TMD FF

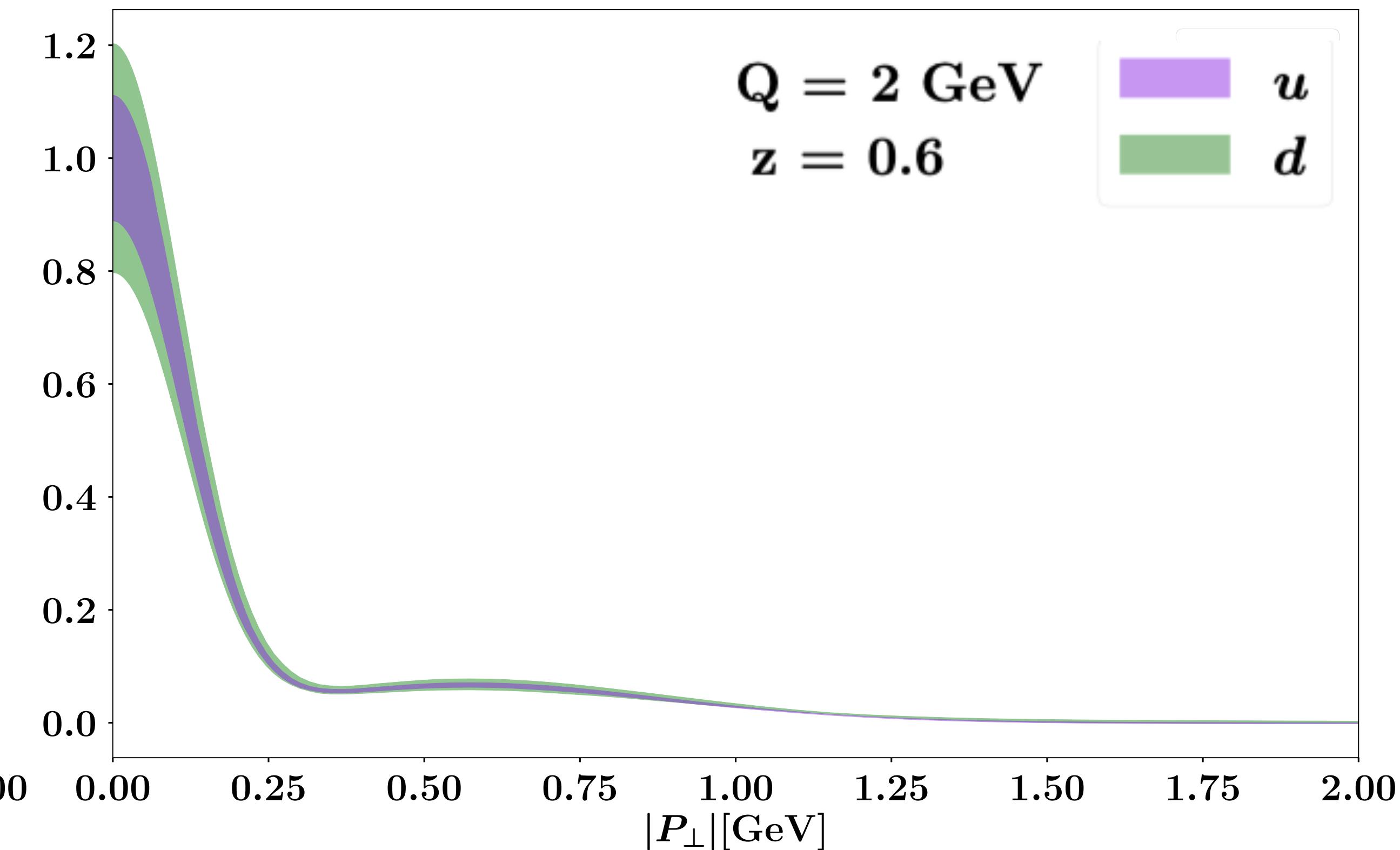
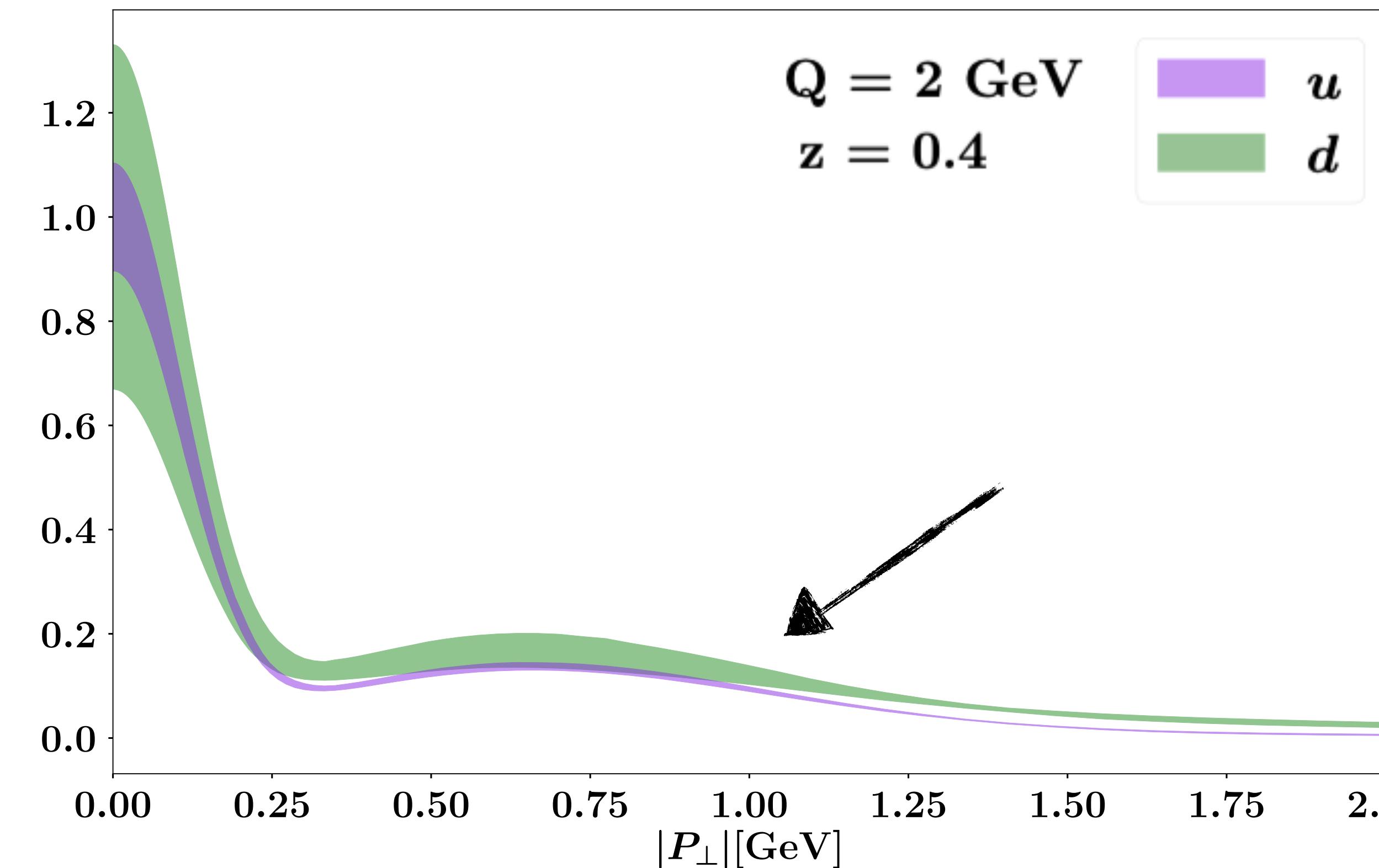
$$\frac{D_1 \rightarrow \pi^+(z, P_\perp^2, Q, Q^2)}{D_1 \rightarrow \pi^+(z, 0, Q, Q^2)}$$



MAPTMD24 - TMD FF

$$\frac{D_1^{\rightarrow\pi^+}(z, P_\perp^2, Q, Q^2)}{D_1^{\rightarrow\pi^+}(z, 0, Q, Q^2)}$$

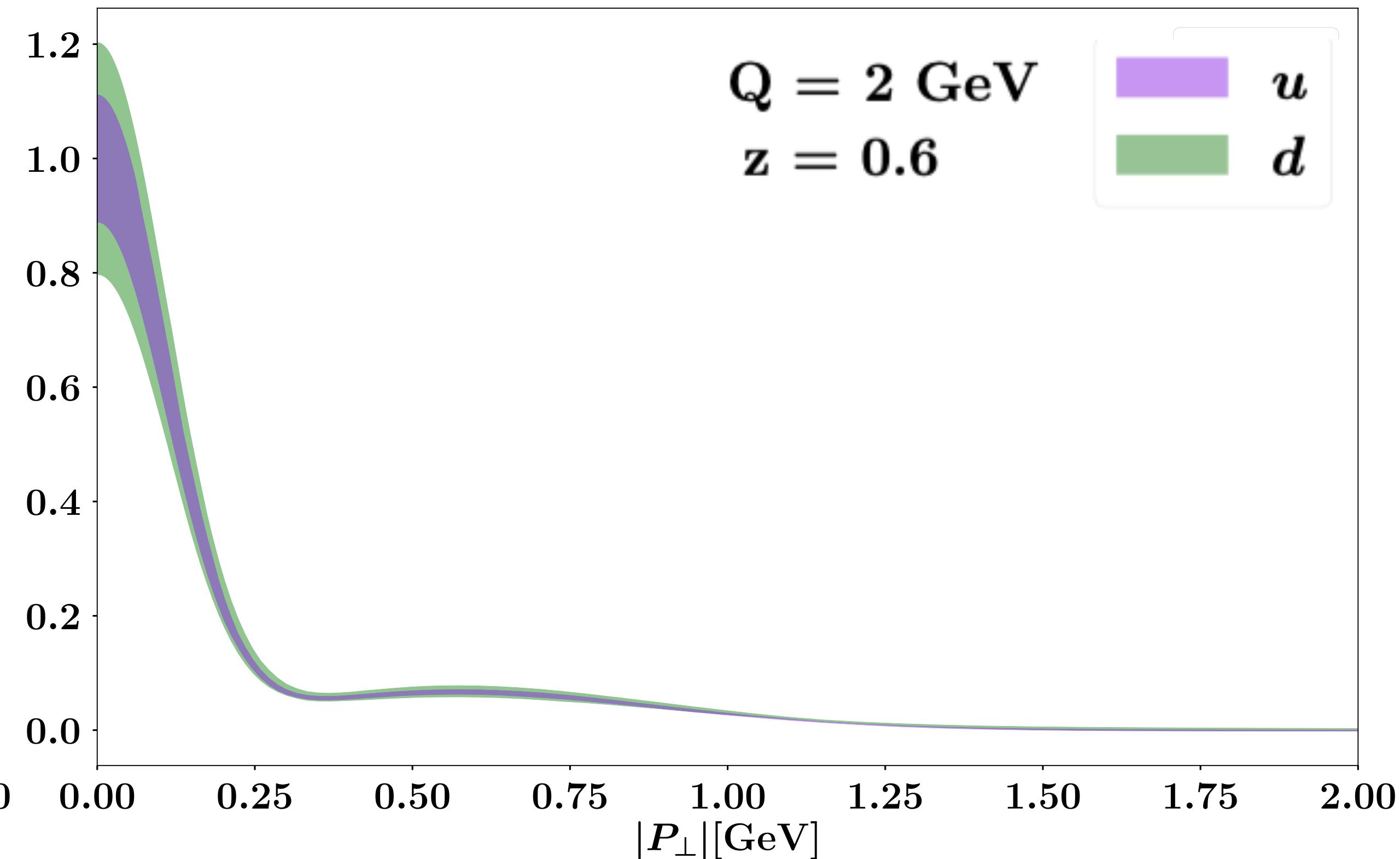
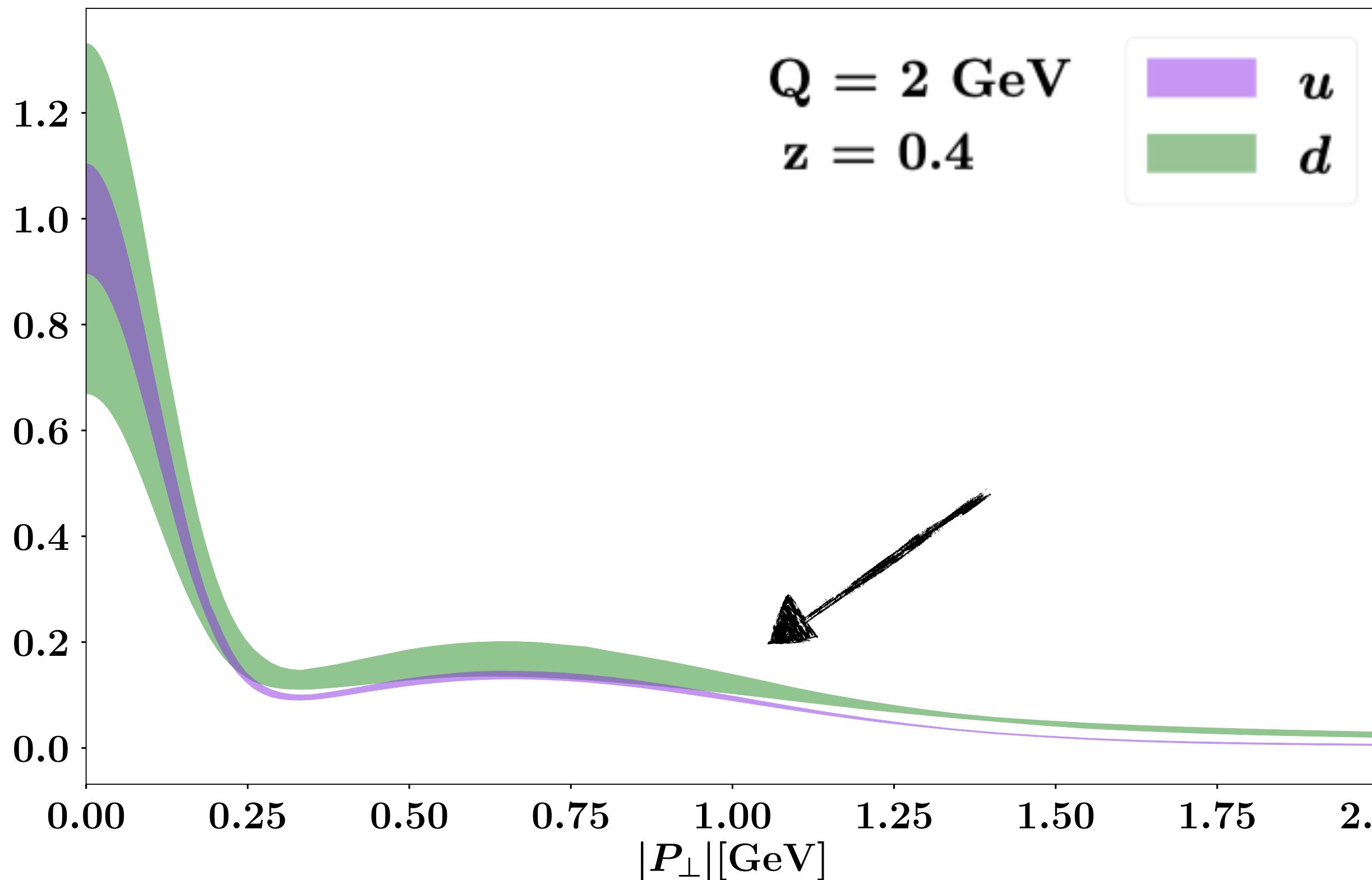
The favored is better constrained than the unfavored one



MAPTMD24 - TMD FF

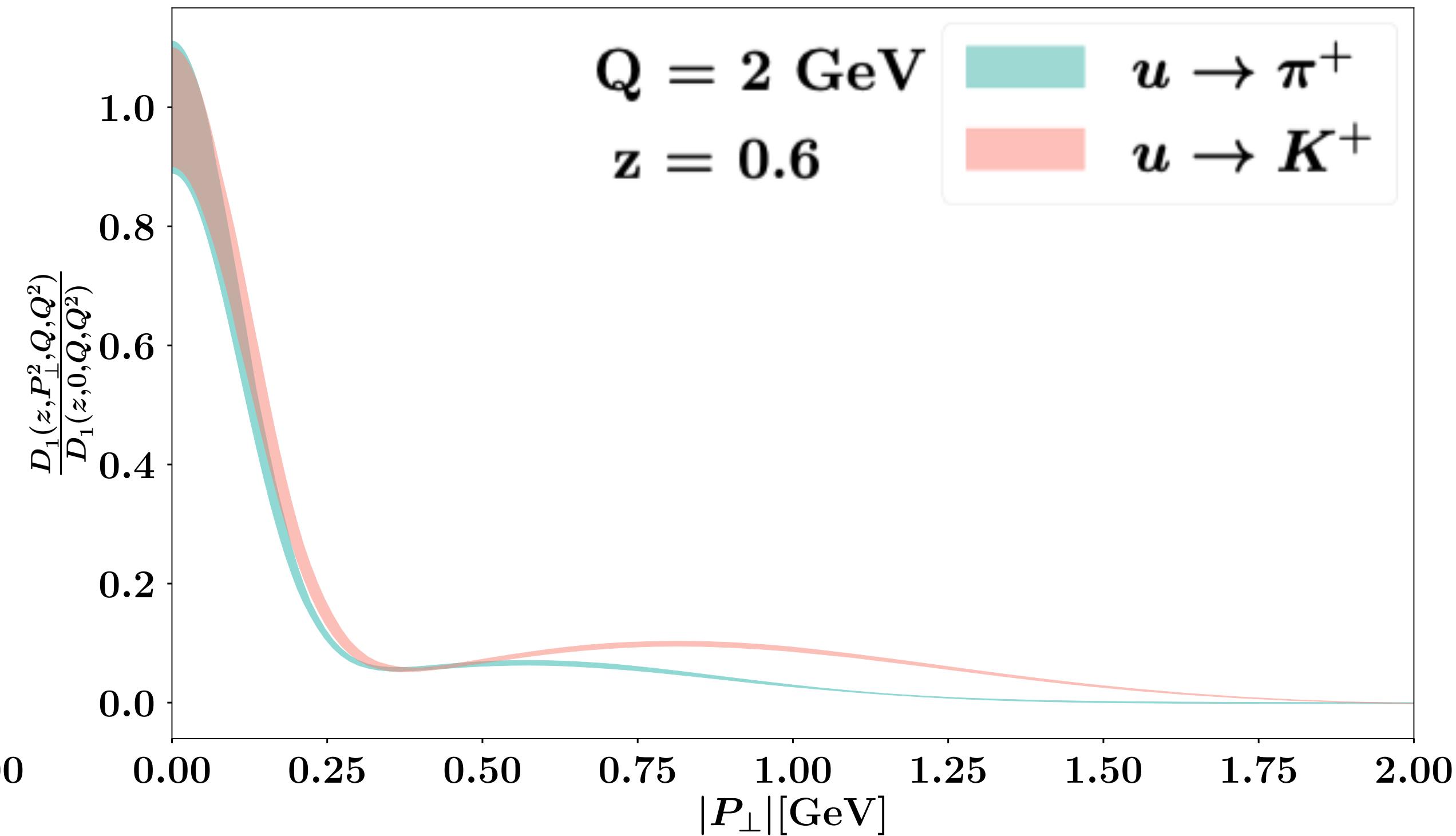
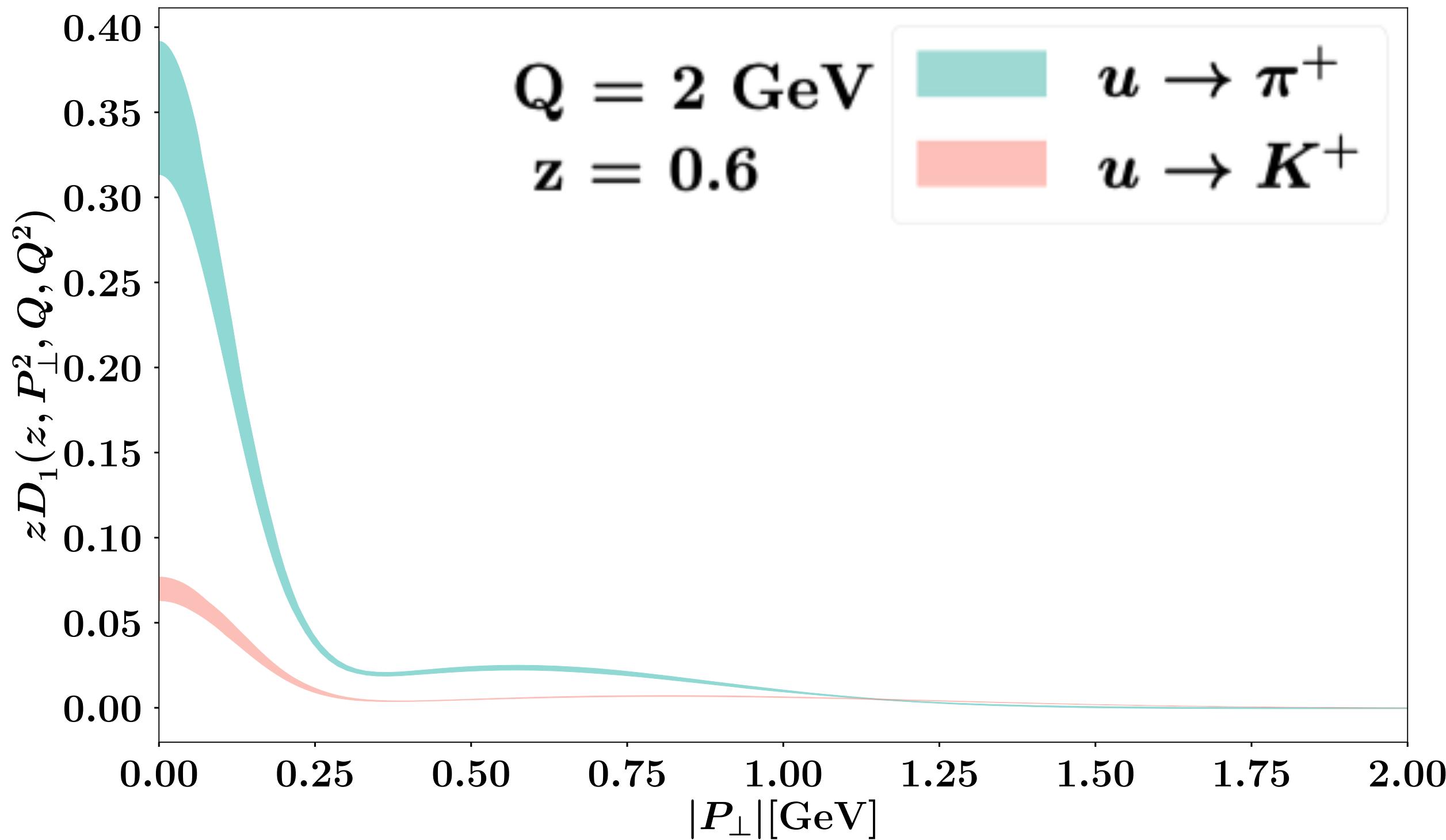
$$\frac{D_1^{\rightarrow\pi^+}(z, P_\perp^2, Q, Q^2)}{D_1^{\rightarrow\pi^+}(z, 0, Q, Q^2)}$$

The favored is better constrained than the unfavored one

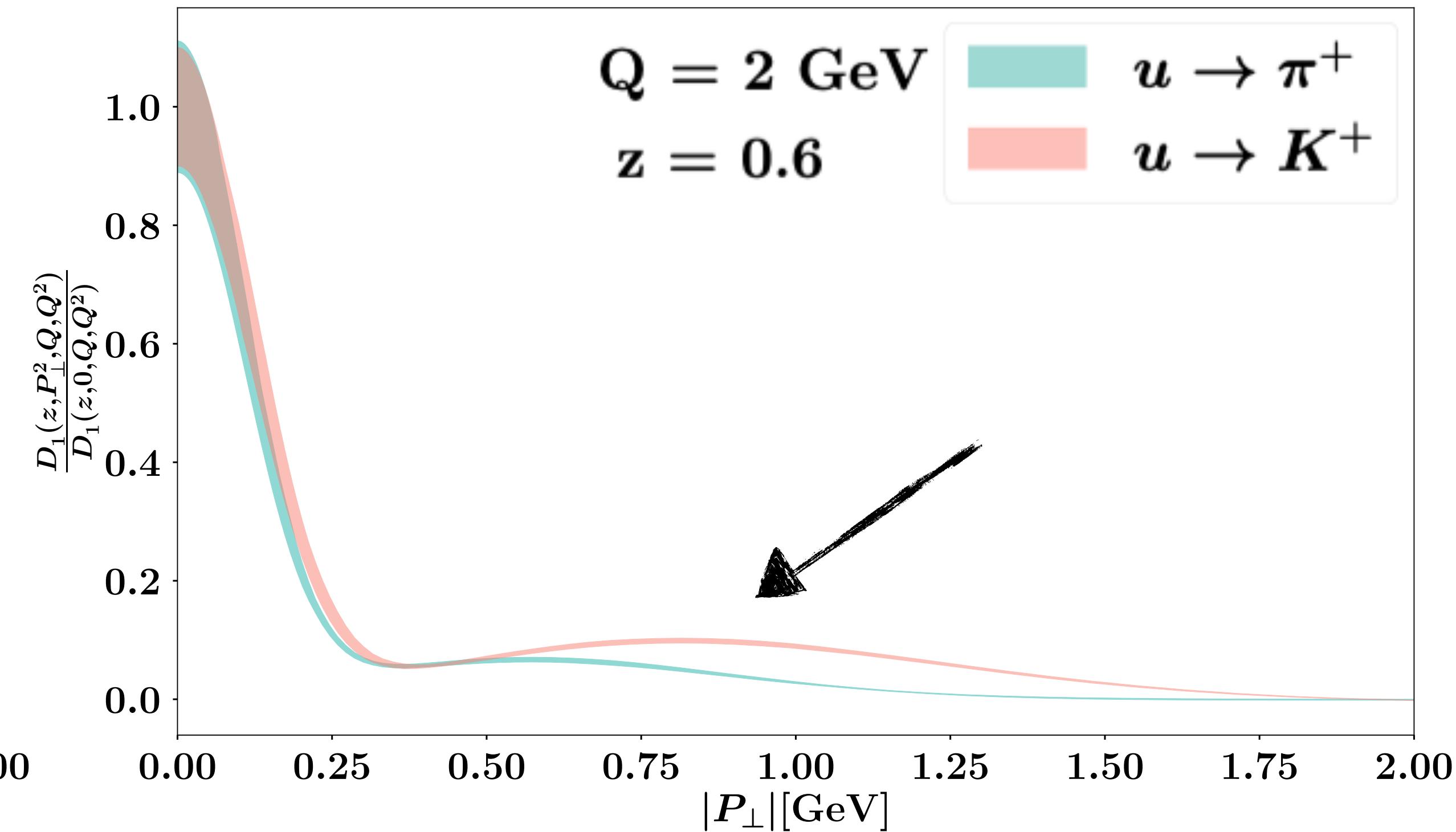
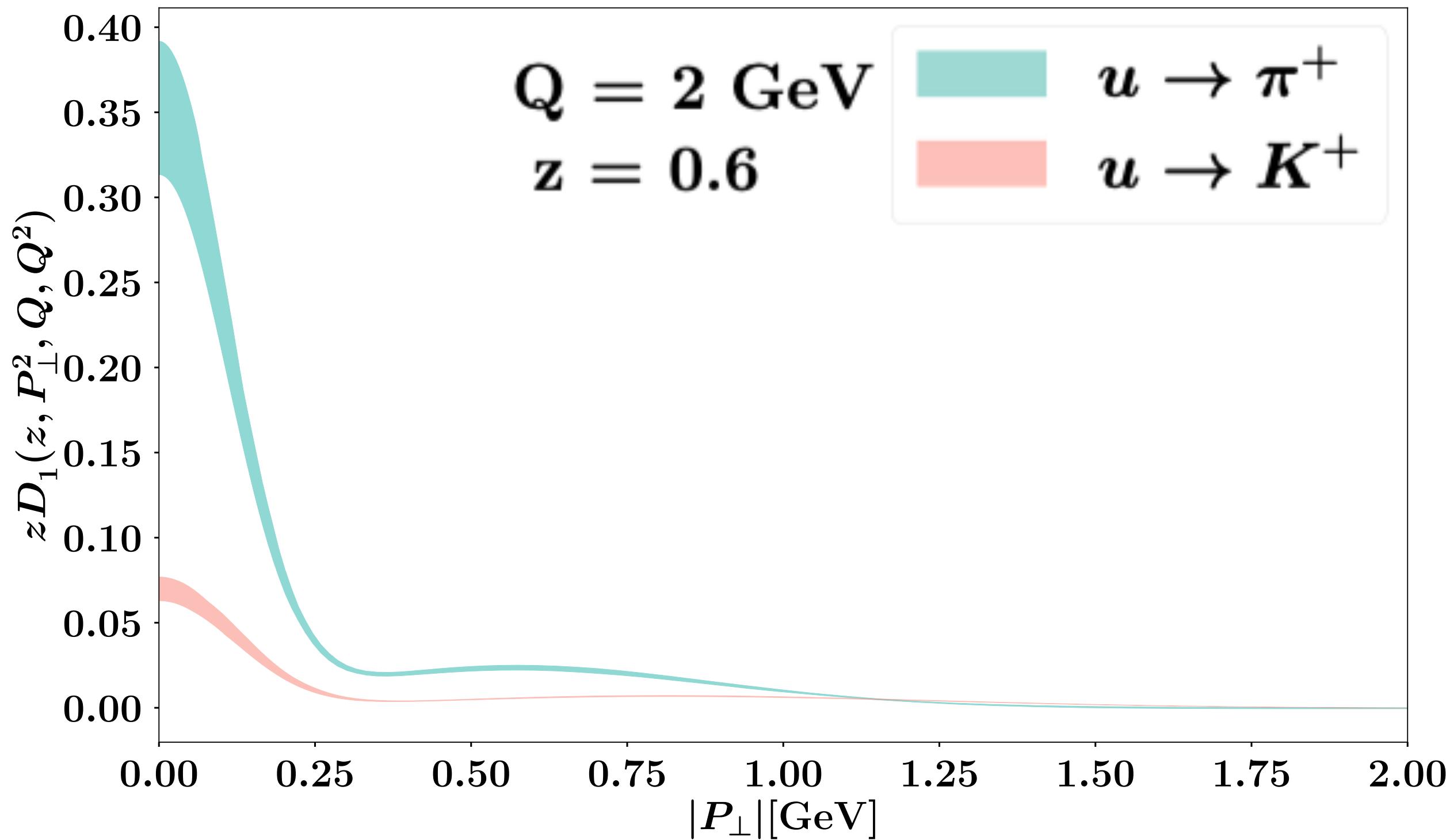


Some signals of differences between favoured and unfavored channels

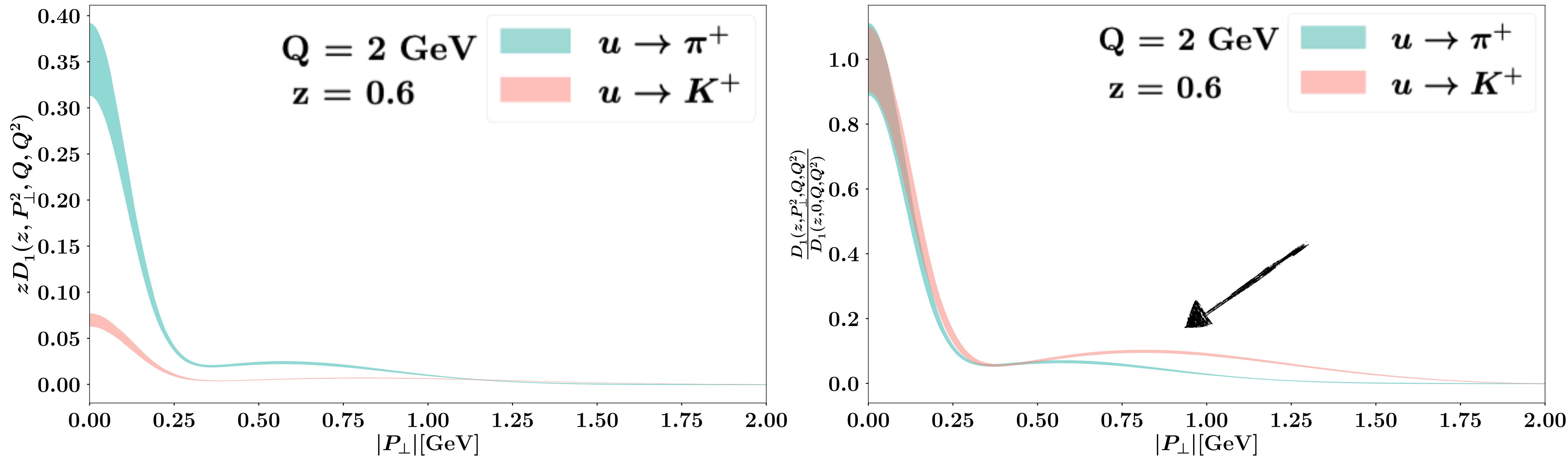
MAPTMD24 - TMD FF



MAPTMD24 - TMD FF

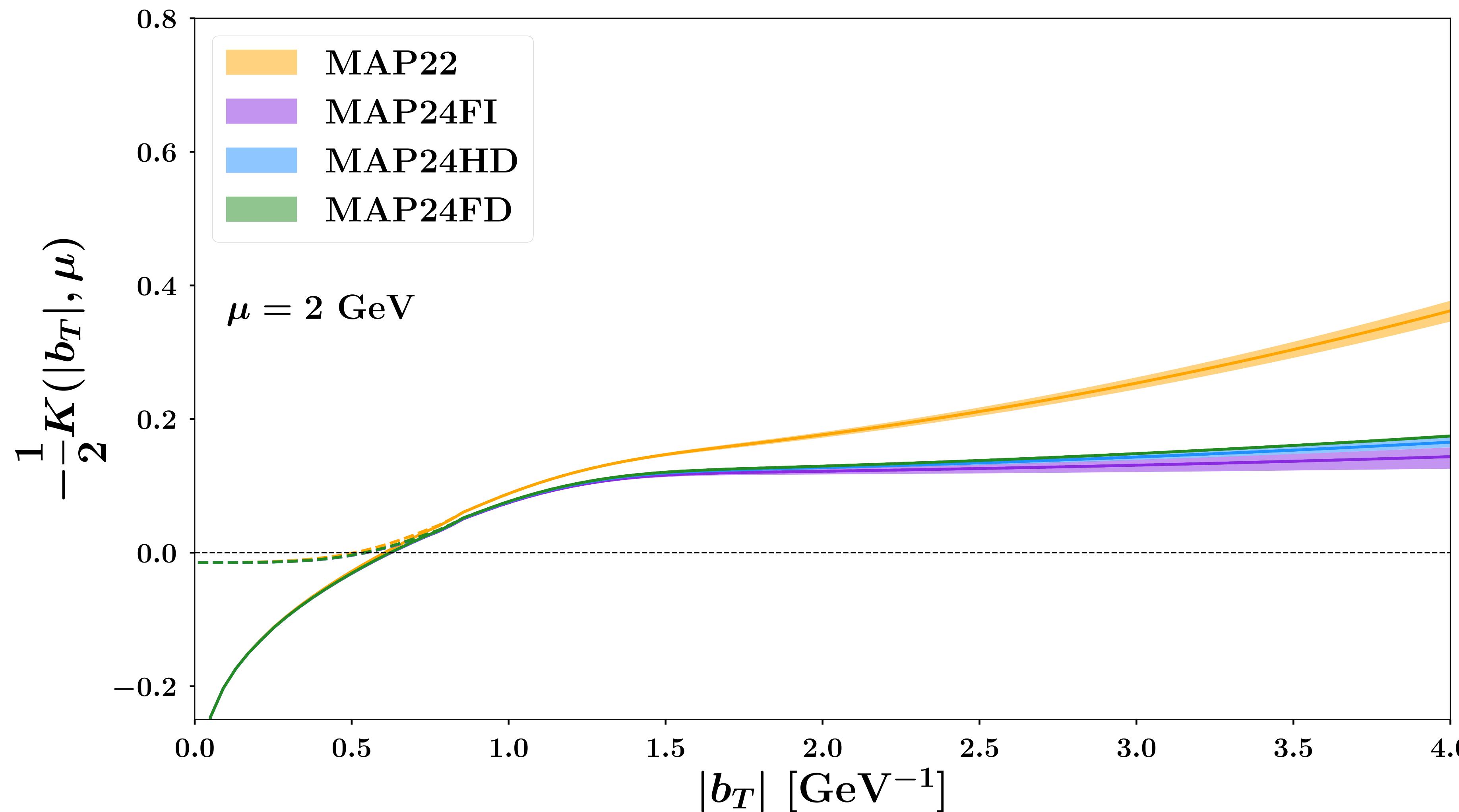


MAPTMD24 - TMD FF

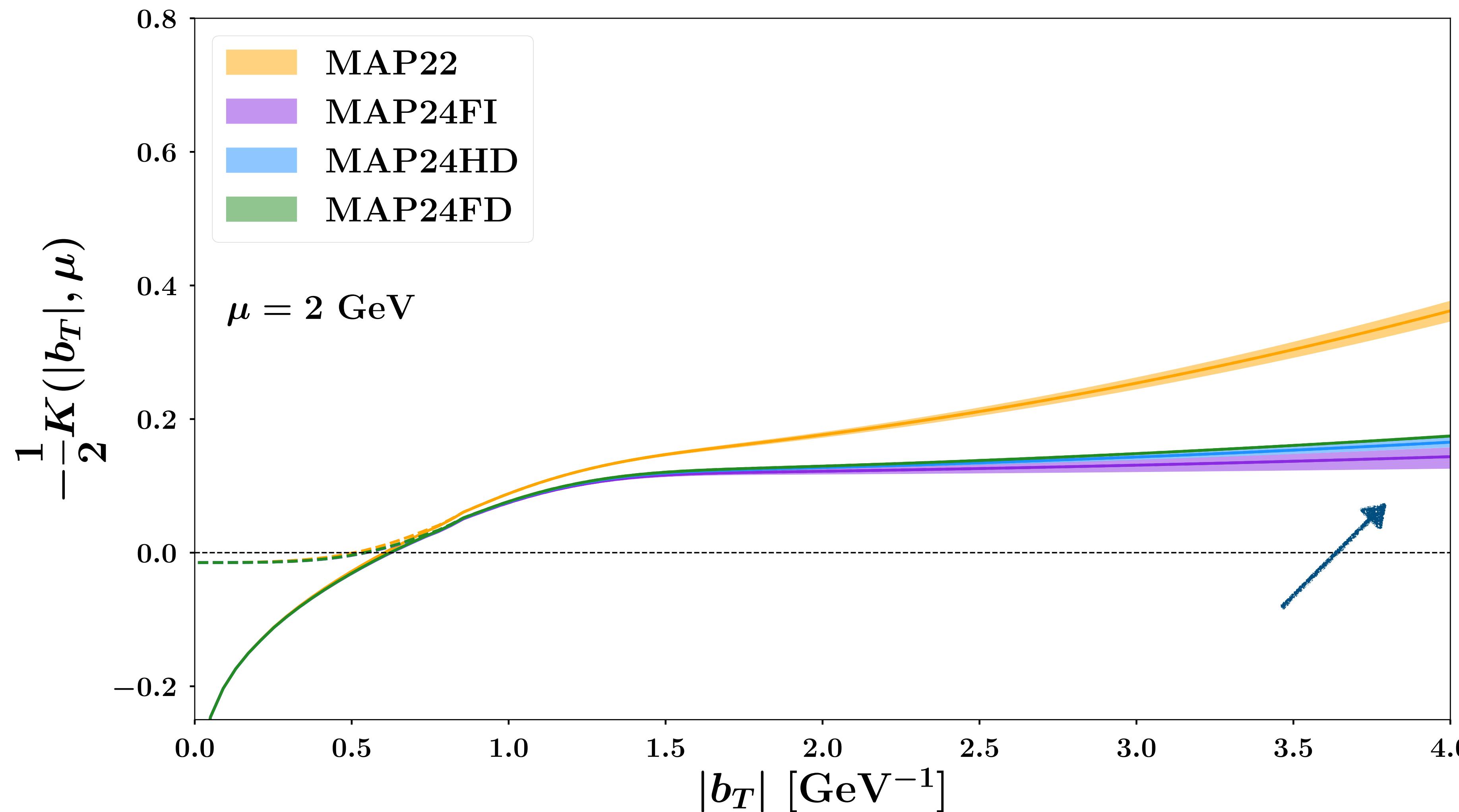


Strong differences between different hadron fragmentations!

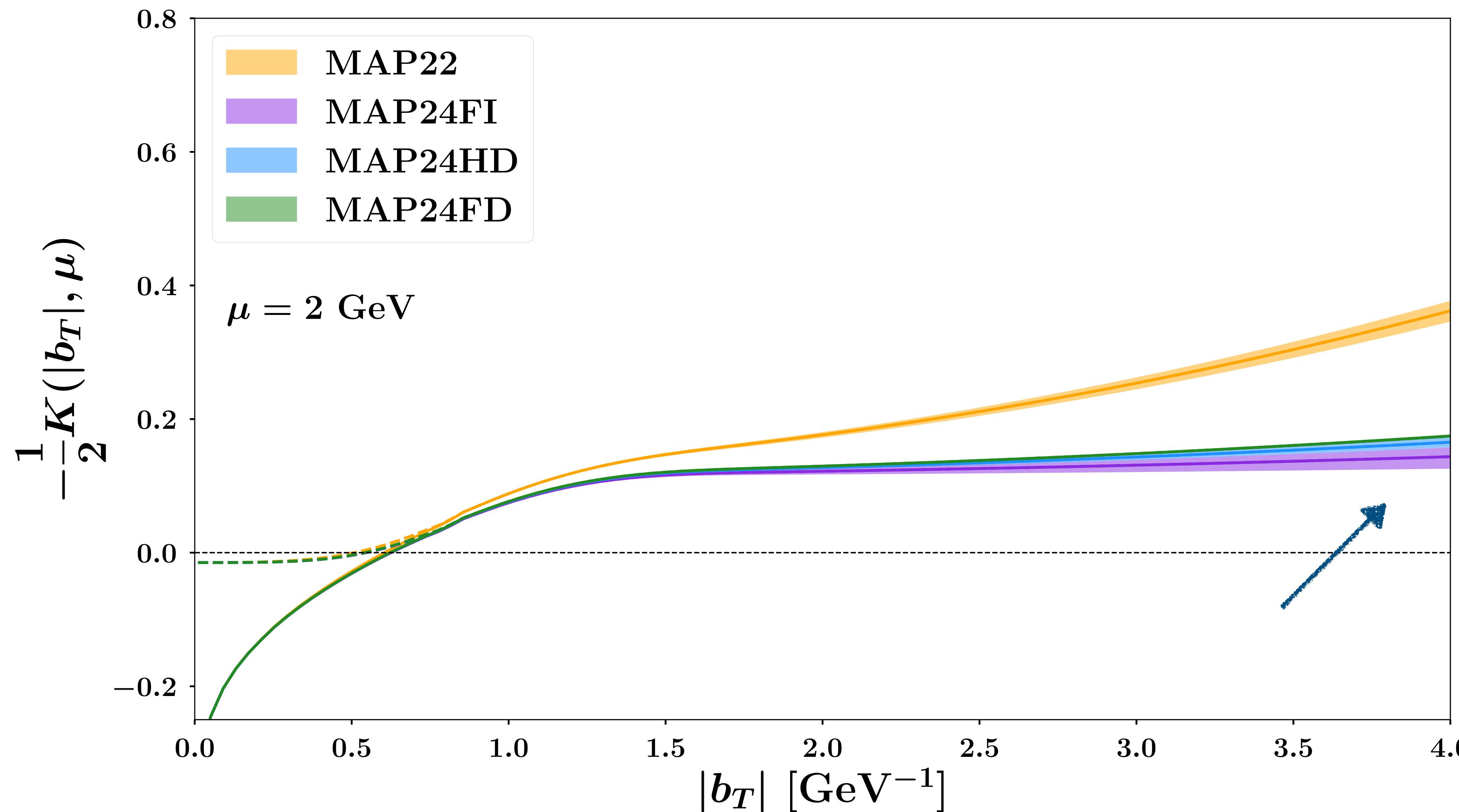
MAPTMD24 evolution - Collins-Soper kernel



MAPTMD24 evolution - Collins-Soper kernel

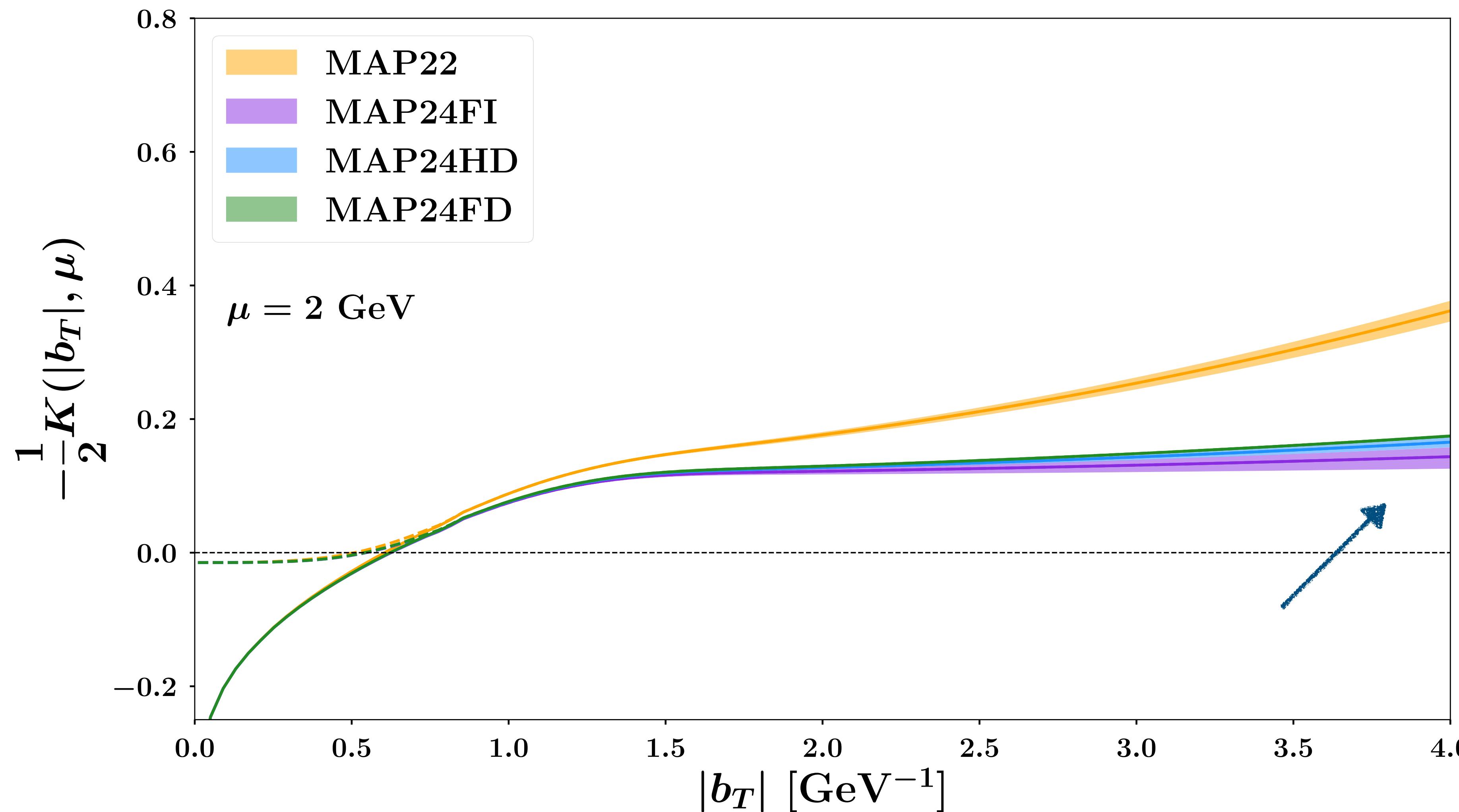


MAPTMD24 evolution - Collins-Soper kernel



*Independent of our
non-perturbative choices*

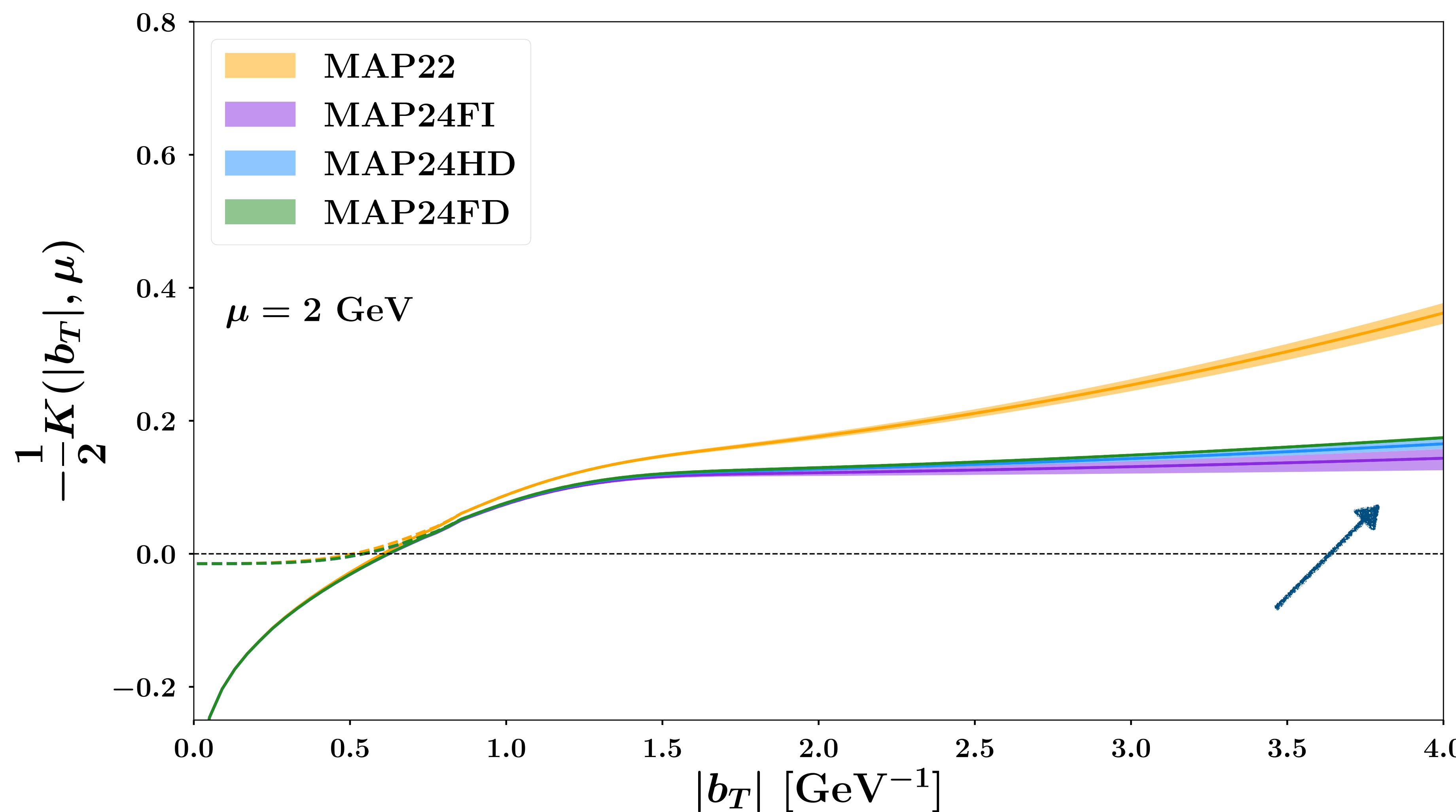
MAPTMD24 evolution - Collins-Soper kernel



*Independent of our
non-perturbative choices*

Quite flat behaviour

MAPTMD24 evolution - Collins-Soper kernel



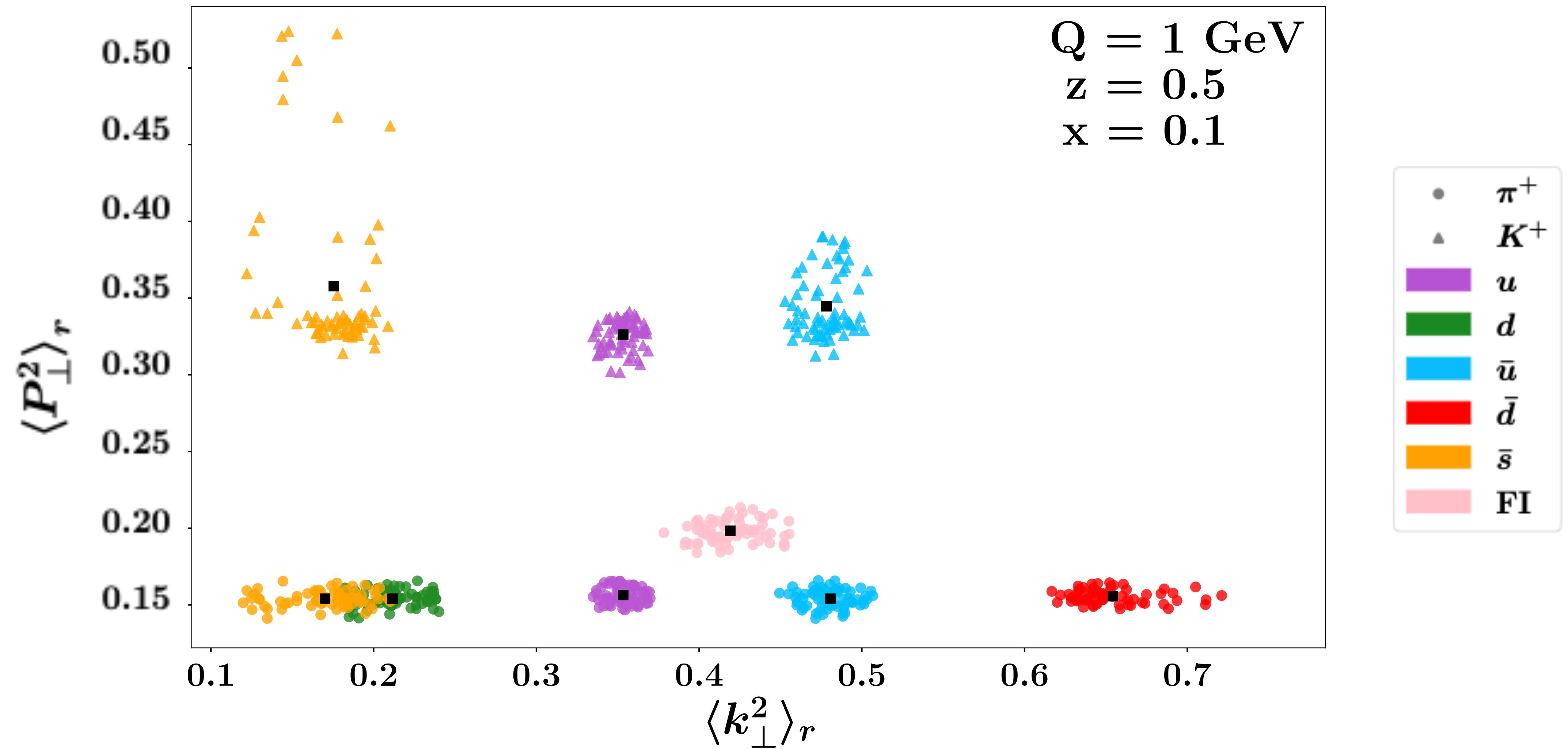
*Independent of our
non-perturbative choices*

Quite flat behaviour

*Compatible with latest
lattice calculation*

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MAPTMD24 - Scatter plots



Evidence of different behaviors for different flavors

Evidence of different behaviors for different measured hadrons

Conclusions

- The extractions of **unpolarized quark TMDs** through global fits have reached very high accuracy (NNNLL), we need to introduce **flavor dependence** to obtain good theory/data agreement
- **MAPTMD24** is the first simultaneous extraction of flavor-dependent unpolarized TMD PDFs and FF through a global fit
- We observed **significant** differences between the flavors in the **TMD PDFs**.
- We observed **significant** differences between different final hadrons in the **TMD FFs**.
- We are finding a weak signal between different flavors in the same final hadron.