Extraction of TMD distributions from data

Alexey Vladimirov

(Regensburg University)

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$\operatorname{Extraction}$ of TMD with <code>artemide</code>

- ▶ Main features
 - Position space
 - ζ-prescription
 - ▶ (maximum) Perturbative matching
 - Strict data cuts
- ▶ Code
 - ▶ artemide
 - ▶ Fortran 95
 - https://github.com/VladimirovAlexey/artemide-public
 - artemide-DataProcessor
 - ▶ Python
 - https://github.com/VladimirovAlexey/artemide-DataProcessor



















Universality & the chain of extractions



In my talk I am going to focus on particular aspects of TMD phenomenology

Data selection for TMD factorization

- ▶ Theoretical justification
- ▶ Experimental evidence
- Impact on present and future data-sets

PDF-bias and problem of TMD modeling

- ▶ Structure of (modern) TMD models
- ▶ PDF-bias
- ► Flavor dependence (work in progress)

Problem with determination of CS-kernel

- ▶ Correlation between TMDs and CS-kernel
- ζ-prescription





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Part I Data cuts



TMD factorization is a systematic expansion of hadronic tensor in power of $\frac{q_T}{Q}$



- Leading-power term is well-investigated
- ▶ Something is known about NLP W_1 (factorization is not proven)
- ▶ Nothing is known about NNLP W_2

Important: TMD factorization is $Q \to \infty$, q_T =fixed. Could be corrections $\sim \frac{\Lambda}{Q}$ even at $q_T \to 0$.

Factorization regions

$$q_T \lesssim \delta Q$$
 TMD factorization $= \begin{cases} q_T \lesssim \Lambda & \text{nonpertrubative regime} \\ q_T \gg \Lambda & \text{"resummation" regime} \end{cases}$

 $q_T \sim Q \gg \Lambda$ collinear factorization







Here I draw $\delta = 0.25$, how to justify this choice?

$$\delta^2 = \frac{q_T^2}{Q^2} \sim 0.06 \ll 1$$

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q_T for SIDIS

The factorization for SIDIS is done in the Breit frame



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Value of δ from the data

- Fit data at some small $\delta~(\chi/N_{pt}\sim 1)$
- **2** Increase δ and fit again, starting from the previous minimum (repeat)

3 At some moment the χ^2/N_{pt} blows up



[Scimemi, AV, 1706.01473]



Data indicates that $\delta \sim 0.2 - 0.25$

$\mathbf{But...}$

- ▶ For asymmetries (ratios of cross-section) δ could be larger
- > The method outcome strongly dependents on precision of the data
 - ▶ Precise data \rightarrow more sensitivity to small effects, e.g. power corrections
 - \blacktriangleright E.g. for ATLAS (~ 0.5% accuracy) at $\delta \sim 0.2$ deviation is $\sim 2-3\%$:(
 - ▶ E.g. for CDF (~5% accuracy) at $\delta \sim 0.2$ deviation is $\sim 2 3\%$:)
- ▶ There could be models which **incorporates** power corrections to factorization into NP-behavior of TMDs
 - ▶ The result of extraction is not a TMD distribution (although it could perfectly describe the data), e.g. it violates universality
 - \blacktriangleright Anyway, at some moment TMD factorization fails (\rightarrow next slide)





The cross-section with LP TMD factorization eventually became negative. It happens at large q_T



- \blacktriangleright The position of node depends also on process and x
- ▶ For large-Q bins the node can go down to $q_T/Q \sim 0.35 \; (\pi \text{DY at COMPASS!})$

The negative cross-section is a small-b problem

$$d\sigma \simeq \int d^2 b e^{-ibq_T} W(b) \simeq \int db b J_0(bq_T) W(b)$$

When 2D Fourier is positive definite?

- ▶ 1D cos-transformation \rightarrow Bochner's theorem \rightarrow "non-growing function"
- ▶ Generally, it is a complicated question see e.g.[Giraud,Peschanski,1405.3155]
- ▶ The first requirement(but not sufficient): W(b) has maximum at b = 0.



This fall down is (mainly) due to TMD perturbative evolution !

$$W(b,Q;x,z) \simeq R(b,Q)^2 F(x,b) D(z,b)$$

▶ In ζ -prescription F and D are (almost) monotonous functions

$$\blacktriangleright R(b,Q) = \exp(-\mathcal{D}(b,Q)\ln(Q^2/\zeta_Q(b)))$$



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Perturbation theory = predictive power



Difference between NNLO and N^3LO is not that important Difference between NLO and NNLO is important (especially at low-energy!)



Perturbation theory = predictive power

B.Bilin, DIS2021



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Principal problem for asymmetries

 ${\rm asymmetry} \simeq \frac{{\rm something}}{F_{UU}}$

• Eventually $F_{UU} = 0$ (in TMD factorization)





Cutting present data : Sivers asymmetry

Dataset name	Ref.	Reaction	# Points	
		$d^{\uparrow} + \gamma^* \rightarrow \pi^+$	1 / 9	
Compass08	[36]	$d^{\uparrow} + \gamma^* \rightarrow \pi^-$	1/9	
		$d^{\uparrow} + \gamma^* \rightarrow K^+$	1/9	
		$d^{\uparrow} + \gamma^* \to K^-$	1 / 9	
Compage16	[30]	$p^{\uparrow} + \gamma^* \rightarrow h^+$	5 / 40	
Compussio	[00]	$p^{\uparrow} + \gamma^* \rightarrow h^-$	5 / 40	
Hermes		$p^{\uparrow} + \gamma^* \rightarrow \pi^+$	11 / 64	
	[35]	$p^{\uparrow} + \gamma^* \rightarrow \pi^-$	11 / 64	
		$p^{\uparrow} + \gamma^* \rightarrow K^+$	12 / 64	
		$p^{\uparrow} + \gamma^* \rightarrow K^-$	12 / 64	
JLab	[41, 42]	$p^{\uparrow} + \gamma^* \rightarrow \pi^+$	1/4	
		$p^{\uparrow} + \gamma^* \rightarrow \pi^-$	1/4	
		$p^{\uparrow} + \gamma^* \rightarrow K^+$	1/4	
		$p^{\uparrow} + \gamma^* \rightarrow K^-$	0 / 4	
SIDIS total			63	
CompassDY	[40]	$\pi^- + d^\uparrow \to \gamma^*$	2 / 3	
Star.W+		$p^{\uparrow} + p \rightarrow W^+$	5 / 5	
Star.W-	[43]	$p^{\uparrow} + p \rightarrow W^{-}$	5 / 5	
Star.Z		$p^{\uparrow} + p \rightarrow \gamma^*/Z$	1 / 1	
DY total			13	
Total			76	







Conclusion for part 1

▶ There is a natural limit of TMD factorization $q_T < (0.2 - 0.3)Q$

- This limit is required from by theory
- ▶ This limit is also seen in the data
- Pushing this limit higher does not help practically
 - ▶ At $q_T \sim 0.5Q$ cross-section become negative
 - It is pure perturbative effect
- ▶ Ways out:
 - ▶ Interpolate to fix order (works only at large Q)
 - ▶ Introduce b_{min}
 - Go to power corrections
 - ...

▶ A lot of stuff to explore especially at lower energy JLab \rightarrow EIC





Part II PDF-bias



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TMD distributions are independent 3D function for each flavor $\text{TMDPDF} = F(x, b), \quad \text{TMDFF} = D(z, b)$ **Too much freedom!**



Current status of the small-b matching

		Twist of	Twist-2	Twist-3	Order of	
Name	Function	leading	distributions	distributions	leading power	Ref.
		matching	in matching	in matching	coef.function	
unpolarized	$f_1(x,b)$	tw-2	$f_1(x)$	-	N ³ LO (α_s^3)	[21, 22]
Sivers	$f_{1T}^{\perp}(x,b)$	tw-3	-	T(-x, 0, x)	NLO (α_s^1)	[23]
helicity	$g_{1L}(x,b)$	tw-2	$g_1(x)$	$T_g(x)$	NLO (α_s^1)	[16, 17]
worm-gear T	$g_{1T}(x,b)$	tw-2/3	$g_1(x)$	$T_g(x)$	LO (α_s^0)	[13, 14]
transversity	$h_1(x,b)$	tw-2	$h_1(x)$	$T_h(x)$	NNLO (α_s^2)	[19]
Boer-Mulders	$h_1^{\perp}(x, b)$	tw-3	-	$\delta T_{\epsilon}(-x,0,x)$	LO (α_s^0)	[14]
worm-gear L	$h_{1L}^{\perp}(x,b)$	tw-2/3	$h_1(x)$	$T_h(x)$	LO (α_s^0)	[13, 14]
pretzelosity	h_{1T}^{\perp}	tw-3/4	-	$T_h(x)$	LO (α_s^0)	eq.(4.8)

refs. are defined in [V.Moos,AV,2008.01744]

▶ Twist-2 and twist-3 contributions at all powers of b^2 (tree)

► Typical expression (here for Sivers function):

$$f_{1T}^{\perp}(x,b) = \pm \pi \Big\{ T_q(x) + \sum_{n=1}^{\infty} \left(\frac{x^2 b^2 M^2}{4} \right)^n \int_0^1 du \int dy \frac{\delta(x-uy)}{(n+1)!(n-1)!} \left(\frac{\bar{u}}{u} \right)^n \frac{1+(n-1)u+u^2}{1-u} T_q(y) \Big\}$$

▶ T(x) = T(-x, 0, x) is Qiu-Sterman function

 \blacktriangleright TMDs_{proton} ~ TMDs_{nuclei}

Non-trivial matching for pretzelosity

• Leading term:
$$h_{1T}^{\perp}(x,b) = -x^2 \int_x^1 \frac{du}{u} \frac{1-u^2}{u} \mathcal{T}_h\left(\frac{x}{u}\right) + \text{tw-4}$$



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Matching is essential part on nowadays TMD phenomenology

- ▶ The region 5GeV $\lesssim q_T < 0.25Q$ is accurately described by $f_{NP} \sim 1$ ▶ LHC, Tevatron, RHIC, (→ EIC)
- ▶ It is observed that q(x) carries the most part of x-dependence. I.e. $f_{NP}(x,b) \sim f_{NP}(b)$
 - Greatly reduces the parametric freedom
- ▶ It is observed that $q_f(x)$ carries the most part of the flavor dependence, i.e. $f_{NP}(x, b) \sim$ flavor-independent
- ▶ In fact, the simplest model $F_f(x,b) \sim C \otimes q_f(x) f_{NP}(b)$ capable to describe the most part of the data rather accurately

Matching to PDF leads to high predictive power, but is also a pitfall \rightarrow





Result of a TMD fit is 100% dependent on PDF in use!

- ▶ Different PDF set are different
 - ▶ Especially in a "TMD-important" region x ~ 0.1 - 0.5
 - Different flavor decomposition

► As the result:

PDF & FF sets	χ^2/N_{pt}	
HERA20 & DSS	0.76	
HERA20 & JAM19	0.93	
NNPDF31 & DSS	1.00	
NNPDF31 & JAM19	1.65	
HERA20 & DSS (N^3LO)	0.88	
NNPDF31 & DSS $(N^{3}LO)$	1.31	

SIDIS+DY fit [SV19]



Obviously, one must include PDF uncertainty into the fit

Including PDF uncertainty "straightforwardly"

▶ PDF uncertainty is **larger** than the experimental precision

- ▶ LHC 5-7% vs. 1%
- ▶ Low energy DY 10-50% vs. 10%
- SIDIS 10-50% vs.

 \blacktriangleright TMD physics (in comparison to DIS) is sensitive to different x-domain

▶ Strongly depends on the set



 $\begin{array}{l} \leftarrow \text{SV19 fit} \\ \text{NNPDF+DSS} \\ \text{SIDIS+DY} \\ N_{pt} = 1039 \\ \text{fit made for central replica} \\ \leftarrow \text{ distribution of } \chi^2 \text{ for} \\ 1000 \text{ replicas of NNPDF} \end{array}$



PDF essentially changes behavior of TMD = PDF-bias



SV19 fit, 40 random replicas of NNPDF3.1

In fact, each PDF replica must be equipped by its own f_{NP} It will partially compensate PDF-bias So, together they form a TMD distribution



Inclusion of PDF uncertainty into TMD fit (work in progress)

Computationally intensive work

- ▶ Represent PDF uncertainty as MC replicas (1000 replicas)
- ▶ Make a fit of TMD distribution, based on each replica

One could expect that result would be less dependent on PDF



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Computationally intensive work

- ▶ Represent PDF uncertainty as MC replicas (1000 replicas)
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One could expect that result would be less dependent on PDF

NO



We have observed that "simple" (5 params!) model from SV19 does not fit all PDF sets equally well **Reason:** absence of flavor dependence **Solution:** add flavor dependence $f_{NP} \rightarrow f_{NP}^{u,d,\bar{u},\bar{d},rest}$



Now, TMDs based on different PDFs are in agreement



Meanwhile the NP parameters of model strongly distributed λ_1 λ_2 λ_3 λ_4



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Conclusion for part 2

▶ Matching of TMD to PDF is important!

- ▶ TMD totally dependent on the PDF in use
 - There is no agreement between PDF sets
 - \blacktriangleright Uncertainty in PDF lead to crazy TMDs \rightarrow fit each PDF replica
- ▶ To compensate PDF-bias one needs flavor dependence
 - Results for different PDFs are in agreement
 - ▶ TMDs are in agreement
 - Uncertainty on TMD is much larger

▶ Work in progress

▶ Future: one needs joint fit of PDF + TMD



Part III Decorrelation of TMD evolution



TMD evolution depends on non-perturbative CS-kernel

$$\mathcal{D} = -\frac{1}{2}K = \frac{1}{2}F_{q\bar{q}} = -\frac{1}{2}\gamma_{\nu}^{f_{\perp}}$$

$$\mu^2 \frac{d}{d\mu^2} F_{f \leftarrow h}(x, b; \mu, \zeta) = \frac{\gamma_F^f(\mu, \zeta)}{2} F_{f \leftarrow h}(x, b; \mu, \zeta)$$

$$\zeta \frac{d}{d\zeta} F_{f \leftarrow h}(x, b; \mu, \zeta) = -\mathcal{D}^f(b, \mu) F_{f \leftarrow h}(x, b; \mu, \zeta)$$

CS is fundamental QCD function

$$\mathcal{D}(b,\mu) = \lambda_{-} \frac{ig}{2} \frac{\mathrm{Tr} \int_{0}^{1} d\beta \langle 0|F_{b+}(-\lambda_{-}n+b\beta)W_{C'}|0\rangle}{\mathrm{Tr} \langle 0|W_{C'}|0\rangle} + Z_{\mathcal{D}}(\mu)$$

- ▶ Independent observable
- ▶ Measures QCD-vacuum



Thus, TMD distributions are functionals of CS-kernel $f_1[\mathcal{D}](x,b;\mu,\zeta)$

- \blacktriangleright Is it a problem? YES, because we extract simultaneously \mathcal{D} and TMDs
 - Extraction is not universal!
 - ▶ E.g. one cannot use D from lattice, together with TMDs from pheno.
 - ▶ In principle, very large/broad pull of data will reduce correlation
 - Problem of comparison/interpretation of result
- ▶ One makes situation worse by splitting to perturbative and NP parts
 - ▶ Keep CS-kernel a whole function



There are <u>several solutions</u> for this problem my-preferred is ζ -prescription



- In a nutshell: define TMD at ζ(b, μ)
 ζ(b, μ) = equi-evolution line (NP!)
- ▶ Equivalent to fixed-point definition
 - TMDs on the same equi-evolution line are the same (by definition!)
- Generally: does not matter which line use as reference
- But there is one very special line = which passes though the saddle point
 - TMD on this line = optimal TMD



Why optimal TMD is optimal?

1 At saddle point $\mathcal{D} = 0$

$$(\text{optimal})f_1(x,b) = f_1[\mathcal{D}=0](x,b;(\mu,\zeta)_{\text{saddle}})$$

Optimal equi-potential line is continuous (important for small-b matching)Greatly simplifies all equations





Everything is nonperturbative! Position of saddle-point, ζ -line,.... Solve all equations in terms of \mathcal{D} !

▶ At large-b saddle-point goes below Λ_{QCD} .

Not possible to build perturbative-like solution

▶ But there is an exact solution! (see [Scimemi,AV,1912.06532,app.C])

$$2\mathcal{D} + 2\beta(a_s)\frac{\partial g(a_s,\mathcal{D})}{\partial a_s} - \Gamma_{\text{cusp}}(a_s)\frac{\partial g(a_s,\mathcal{D})}{\partial \mathcal{D}} + \gamma_V(a_s) = 0.$$
, $g(a_s,0) = 0$





CS-kernel still correlated with the TMDs





Conclusion for part 3

- ▶ Independent extraction of evolution and TMDs is cumbersome task
- ▶ Keep CS-kernel as a whole function!
- ▶ Fixed-scale schemes are preferable
 - \triangleright ζ -prescription





Conclusion

- ▶ Extraction of TMDs is a very peculiar task
 - Involves several NP functions
 - Requires strict data-cuts
 - Perturbative input is important
 - Many open theoretical questions
- ▶ JLab
 - ▶ There will be not much **pure** TMD-factorizable data
 - ▶ Large-x
 - Paradise to study power corrections
 - ▶ Higher-twist TMDs
 - Mass/kinematic corrections
 - Interesting and weakly studied field
 - current/next frontier of QCD
 - Going to be the challenge for theoreticians
- ► EIC
 - ▶ Will happen in 10+ years (I doubt that our understanding will remain the same)





Always take into account uncertainties!

