

TMD evolution as a double-scale evolution

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in collaboration with Ignazio Scimemi
based on [1803.11089]

QCD
evolution

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Universität Regensburg

TMD evolution
is
a double-scale evolution

$$F(x, b; \mu, \zeta)$$

TMD evolution must be taken into account with a great care

More freedom

- Relations between anomalous dimensions
- ζ -prescription

More ambiguity

- Many scales to "tune"
- Violation of transitivity in "naive" formulation (see [1803.11089])



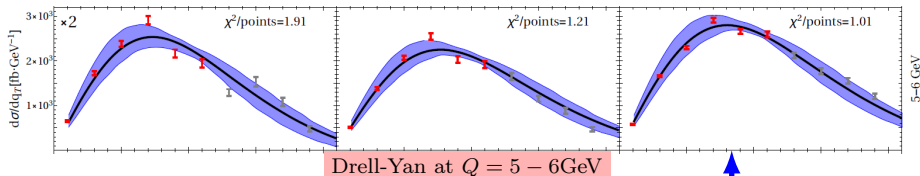
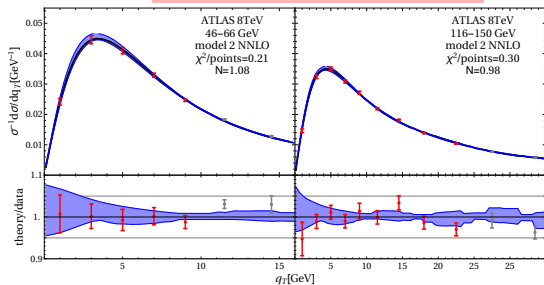
NNLO fit and extraction of (unpol.) TMDPDF

[Scimemi,AV;1706.01473]

- The largest number of data point (DY)
- The largest energy separation
- Consideration of various orders (NLO,NNLL,NNLO)
- Studies of theory error-bands

Included data (at $q_T < 0.2Q$)

	reaction	\sqrt{s}	Q	comment	points
E288	$p + Cu \rightarrow \gamma^* \rightarrow \mu\mu$	19.4 GeV	4-9 GeV	norm=0.8	35
E288	$p + Cu \rightarrow \gamma^* \rightarrow \mu\mu$	23.8 GeV	4-9 GeV	norm=0.8	45
E288	$p + Cu \rightarrow \gamma^* \rightarrow \mu\mu$	27.4 GeV	4-9 & 11-14 GeV	norm=0.8	66
CDF+D0	$p + \bar{p} \rightarrow Z \rightarrow ee$	1.8 TeV	66-116 GeV		44
CDF+D0	$p + \bar{p} \rightarrow Z \rightarrow ee$	1.96 TeV	66-116 GeV		43
ATLAS	$p + p \rightarrow Z \rightarrow \mu\mu$	7 & 8 TeV	66-116 GeV	tiny errors!	18
CMS	$p + p \rightarrow Z \rightarrow \mu\mu$	7 & 8 TeV	60-120 GeV		14
LHCb	$p + p \rightarrow Z \rightarrow \mu\mu$	7 & 8 & 13 TeV	60-120 GeV		30
ATLAS	$p + p \rightarrow Z/\gamma^* \rightarrow \mu\mu$	8 TeV	46-66 GeV		5
ATLAS	$p + p \rightarrow Z/\gamma^* \rightarrow \mu\mu$	8 TeV	116-150 GeV		9
				Total	309

Drell-Yan at $Q = 116 - 150 \text{ GeV}$ 

TMD evolution is a key element

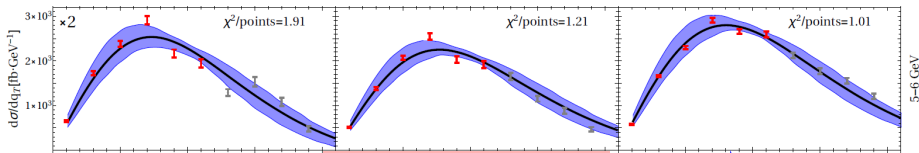
$$\frac{\chi_{\text{global}}^2}{d.o.f.} \simeq 1.25$$

Here:

- 3-loop evolution
- 2-loop coefficient function
- 2-loop matching
- ζ -prescription

plots from [1706.01473]



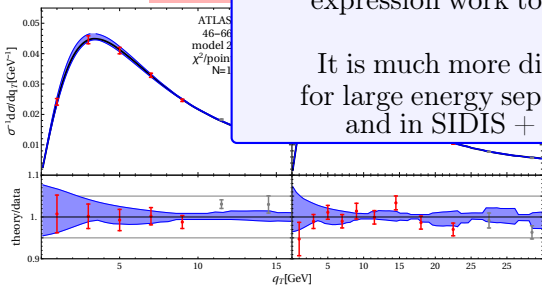


Drell-Yan at $Q = 5 - 6 \text{ GeV}$

Drell-Yan

The main difficulty is to make parts of factorized expression work together

It is much more difficult for large energy separation and in SIDIS + DY



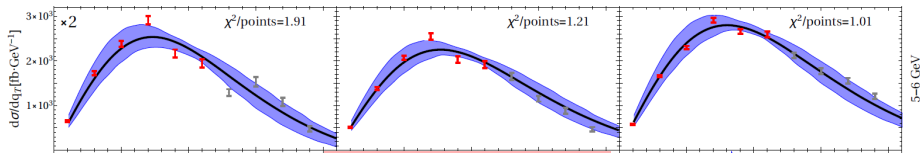
Evolution is a key element

$\chi^2_{\text{global}} / \text{d.o.f.} \simeq 1.25$

- 3-loop evolution
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Drell-Yan at $Q = 5 - 6 \text{ GeV}$

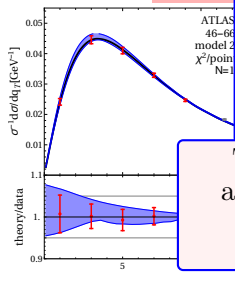
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TMD evolution is the central element, and it requires critical "reexamination"

and it is the topic of this talk

Drell-Yan



evolution is a key element

$$\frac{\chi^2_{\text{global}}}{d.o.f.} \simeq 1.25$$

3-loop evolution

2-loop coefficient

evolution

matching

description

[1706.01473]



TMD evolution is used for two practical purposes

- Compare different experiments
- Modeling TMD distribution

$$\frac{d\sigma}{dX} \sim \int d^2b e^{i(bq_T)} H_{ff'}(Q, \mu) F_{f \leftarrow h}(x_1, b; \mu, \zeta_1) F_{f' \leftarrow h}(x_2, b; \mu, \zeta_2)$$



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Minimize $\ln(Q/\mu)$
 $\mu = Q$

$\zeta_1 \zeta_2 = Q^4$
 or
 $\zeta_1 = \zeta_2 = Q^2$



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Minimize $\mathbf{L}_\mu, \mathbf{L}_{\sqrt{\zeta}}$
 $\mu \sim \sqrt{\zeta} \sim b^{-1}$

$$F(x, b; \mu, \zeta) \sim C(x, b; \mu, \zeta) \otimes \text{PDF}(x, \mu)$$

Typical model for TMD includes matching



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$$F(x, b; \mu_f, \zeta_f) = R[b, (\mu_f, \zeta_f) \rightarrow (\mu_i, \zeta_i)] F(x, b; \mu_i, \zeta_i)$$

Final
scale

TMD evolution factor

Initial
scale

Minimize $\mathbf{L}_\mu, \mathbf{L}_{\sqrt{\zeta}}$
 $\mu \sim \sqrt{\zeta} \sim b^{-1}$

$$F(x, b; \mu, \zeta) \sim C(x, b; \mu, \zeta) \otimes \text{PDF}(x, \mu)$$

Typical model for TMD includes matching



TMD evolution equations

$$\mu^2 \frac{d}{d\mu^2} F_{f\leftarrow h}(x, b; \mu, \zeta) = \frac{\gamma_F^f(\mu, \zeta)}{2} F_{f\leftarrow h}(x, b; \mu, \zeta), \quad (1)$$

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

Solution: $F(x, \mathbf{b}; \mu_f, \zeta_f) = R[\mathbf{b}; (\mu_f, \zeta_f) \rightarrow (\mu_i, \zeta_i)] F(x, \mathbf{b}; \mu_i, \zeta_i)$

- γ_F – TMD anomalous dimension
- \mathcal{D} – rapidity anomalous dimension ($= -\frac{\tilde{K}}{2}$ [Collins' book], $= K$ [Bacchetta, [at,1703.10157](#)])
- Anomalous dimensions are *universal*, i.e. depend only on flavor (gluon/quark).



TMD evolution is two-dimensional

$$\begin{pmatrix} \mu^2 \frac{d}{d\mu^2} \\ \zeta \frac{d}{d\zeta} \end{pmatrix} F = \begin{pmatrix} \frac{\gamma_F}{2} \\ -\mathcal{D} \end{pmatrix} F$$



$$\vec{\nabla} F = \vec{\mathbf{E}} F$$

$\vec{\mathbf{E}}$ is 2D evolution field
in $\vec{\nu} = (\ln \mu^2, \ln \zeta)$
coordinates

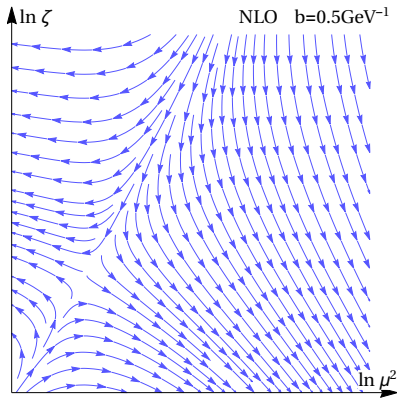


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Solution

$$R[(\mu_f, \zeta_f) \rightarrow (\mu_i, \zeta_i)] = \exp\left(\int_P d\vec{\nu} \cdot \vec{\mathbf{E}}\right)$$

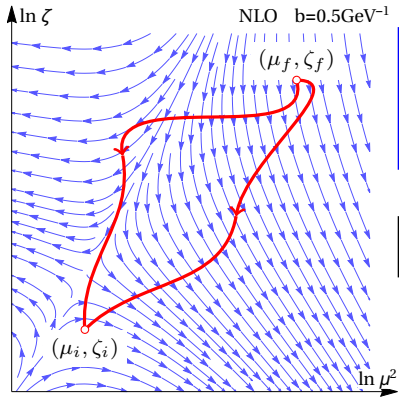


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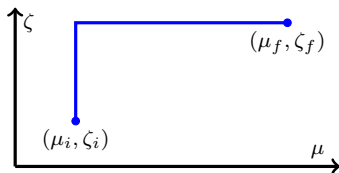
Solution

$$R[(\mu_f, \zeta_f) \rightarrow (\mu_i, \zeta_i)] = \exp \left(\int_P d\vec{\nu} \cdot \vec{E} \right)$$

The integration path
is unimportant!



Examples



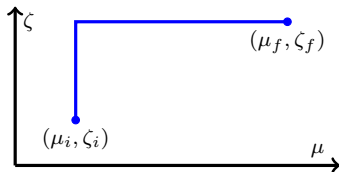
Solution 1

$$\ln R = \int_{\mu_i}^{\mu_f} \frac{d\mu}{\mu} \gamma_F(\mu, \zeta_f) - \mathcal{D}(\mu_i, b) \ln \left(\frac{\zeta_f}{\zeta_i} \right)$$

[Collins' textbook],[Aybat,Rogers,1101.5057],...
99% popular



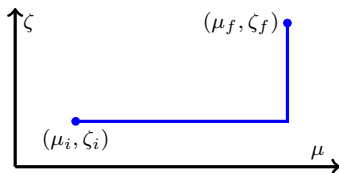
Examples



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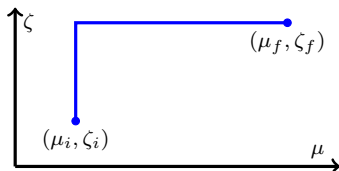


Solution 2

$$\ln R = \int_{\mu_i}^{\mu_f} \frac{d\mu}{\mu} \gamma_F(\mu, \zeta_i) - \mathcal{D}(\mu_f, b) \ln \left(\frac{\zeta_f}{\zeta_i} \right)$$



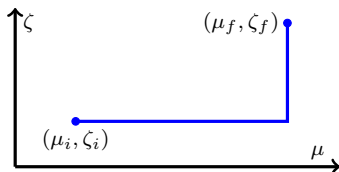
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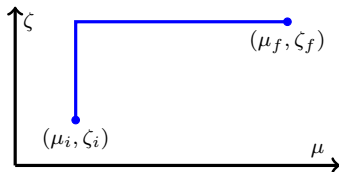


Solution 2

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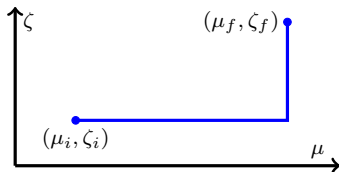
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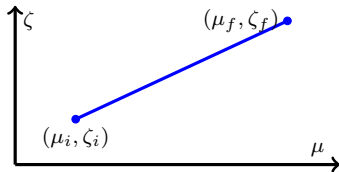
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Solution 2

$$\ln R = \int_{\mu_i}^{\mu_f} \frac{d\mu}{\mu} \gamma_F(\mu, \zeta_i) - \mathcal{D}(\mu_f, b) \ln \left(\frac{\zeta_f}{\zeta_i} \right)$$



Solution 3

$$\ln R = \int_0^1 \left(\gamma_F(\mu(t), \zeta(t)) \frac{\mu_f - \mu_i}{(\mu_f - \mu_i)t + \mu_i} - \mathcal{D}(\mu(t), b) \frac{\zeta_f - \zeta_i}{(\zeta_f - \zeta_i)t + \zeta_i} \right) dt$$

Unique solution

Solution exist only if
integrability condition holds

$$\zeta \frac{d\gamma_F}{d\zeta} = -\mu^2 \frac{d\mathcal{D}}{d\mu^2}$$



$$\vec{\nabla} \times \vec{\mathbf{E}} = 0$$

$\vec{\mathbf{E}}$ is conservative field



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Integrability condition is trivially satisfied due to
collinear overlap of divergences

$$\zeta \frac{d}{d\zeta} \gamma_F(\mu, \zeta) = -\Gamma(\mu), \quad \mu^2 \frac{d}{d\mu^2} \mathcal{D}(\mu, b) = \Gamma(\mu)$$

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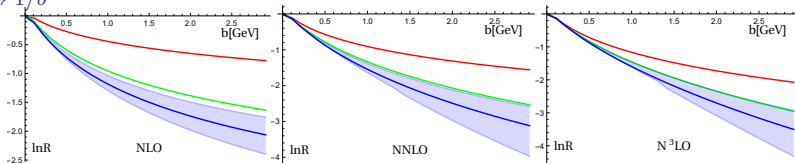
\vec{E} is conservative field

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$$\zeta \frac{d}{d\zeta} \gamma_F(\mu, \zeta) = -\Gamma(\mu), \quad \mu^2 \frac{d}{d\mu^2} \mathcal{D}(\mu, b) \neq \Gamma(\mu)$$

In fixed order PT integrability condition is violated.
The restoration procedure is ambiguous
(large impact at large- b)
See extended dicussion in [[Scimemi,AV;1803.11089](#)]

$M_Z \rightarrow 1/b^*$



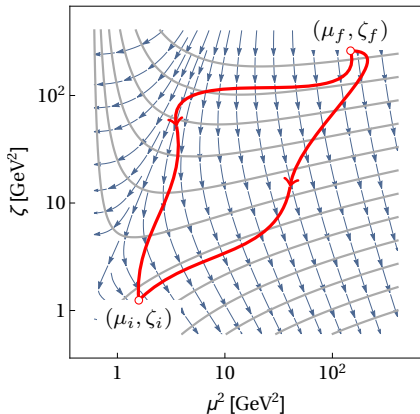
Evolution potential

Solution exist only if
integrability condition holds

$$\zeta \frac{d\gamma_F}{d\zeta} = -\mu^2 \frac{dD}{d\mu^2}$$

$$\vec{\nabla} \times \vec{E} = 0$$

\vec{E} is conservative field



Conservative field is determined
by a potential

$$\vec{E} = \vec{\nabla} U$$

Evolution is a difference
between potentials

$$R[(\mu_f, \zeta_f) \rightarrow (\mu_i, \zeta_i)] = \exp(U_f - U_i)$$



Evolution potential

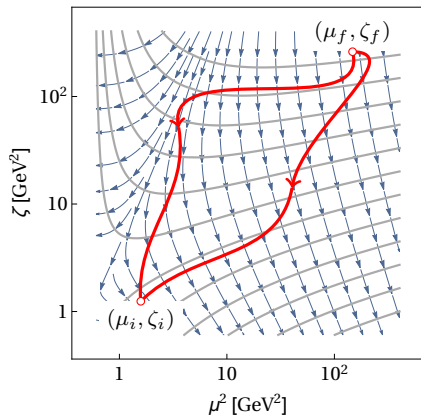
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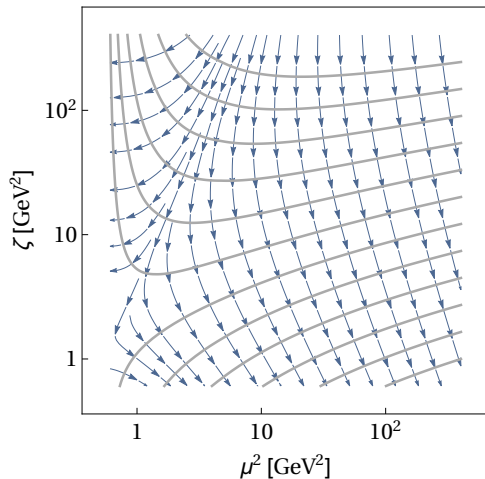
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**This absolutely standard picture
contains an important message.**

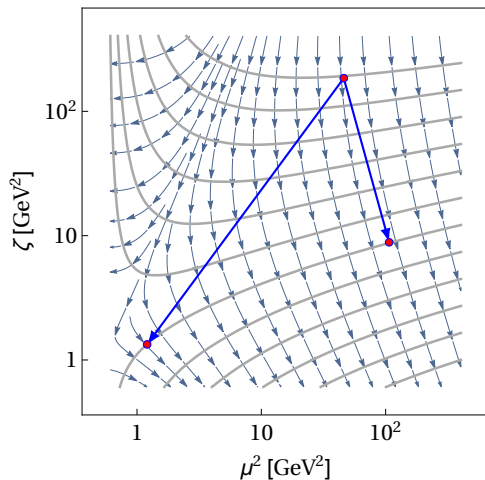


The TMD distribution is not defined by a scale (μ, ζ)
 It is defined by an equipotential line.



The scaling is defined by
~~a difference between scales~~
 a difference between potentials

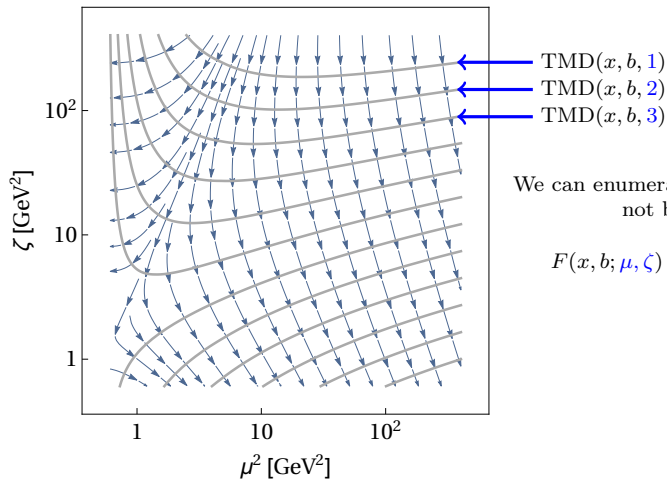
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The scaling is defined by
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Evolution factor to both points
 is the same
 although the scales are
 different by 10^2GeV^2

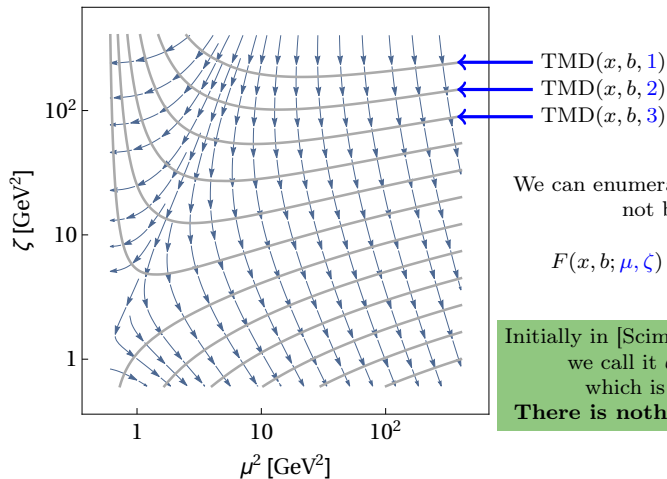
TMD distributions on the same equipotential line are equivalent.



We can enumerate them by a lines
not by (μ, ζ)

$$F(x, b; \mu, \zeta) \rightarrow F(z, b; \text{line})$$

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Initially in [Scimemi,AV,1706.01473]
we call it ζ -prescription
which is misleading.
There is nothing to prescribe.



In ζ -prescription we set
 $\zeta \rightarrow \zeta_\mu(\boldsymbol{\nu})$

- TMDs are "enumerated" by $\boldsymbol{\nu}$ (the number of line)
- TMDs are "naive" scale-independent

$$\mu \frac{d}{d\mu} F(x, b; \mu, \zeta_\mu) = 0 \quad \Rightarrow \text{No double-logs in the matching.}$$



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TMD distribution depends only on the "number" of equipotential line

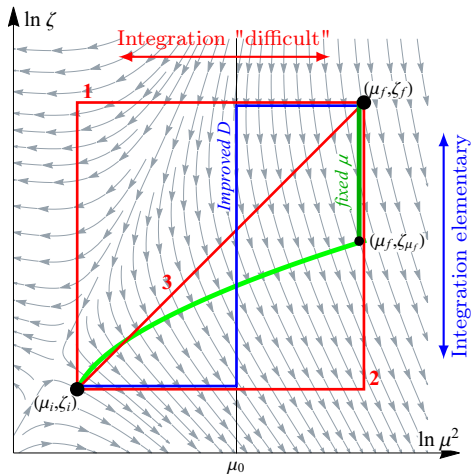
$$F(x, \mathbf{b}; \mu, \zeta) \rightarrow F(x, \mathbf{b}; \nu)$$

$$\frac{dF(x, \mathbf{b}; \nu)}{d\nu} = \frac{dU(\mathbf{b}; \nu)}{d\nu} F(x, \mathbf{b}; \nu)$$

$$\Updownarrow$$

$$F(x, \mathbf{b}; \nu) = e^{U(\mathbf{b}; \nu) - U(\mathbf{b}; \nu_0)} F(x, \mathbf{b}; \nu_0)$$

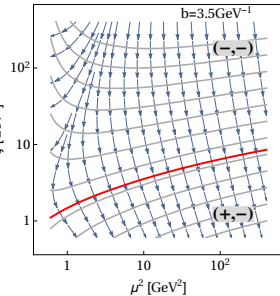
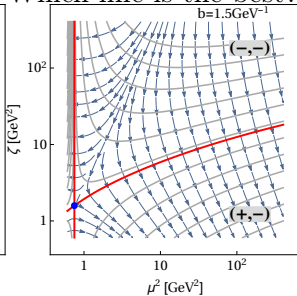
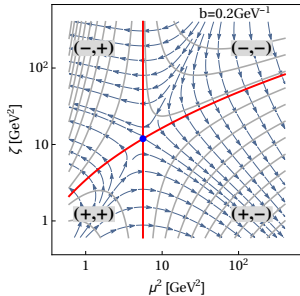
Best way to measure the difference between potentials



$$R = \left(\frac{\zeta_f}{\zeta_{\mu_f}} \right)^{-\mathcal{D}(\mu_f, b)}$$

- Numerically simple (and fast)
- $\mu_f = Q$ thus a_s is small
- Does just the same job as the Sudakov exponent

Which line is the best?



- Some non-interesting singularities at $\mu, \zeta \rightarrow \infty$
- Landau pole at $\mu = \Lambda$
- Saddle point (blue dot)

$$\mathcal{D}(\mu_{\text{saddle}}, b) = 0, \quad \gamma_M(\mu_{\text{saddle}}, \zeta_{\text{saddle}}, b) = 0$$

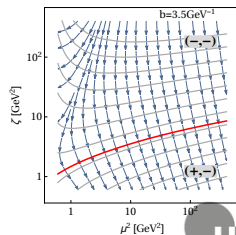
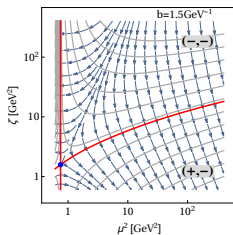
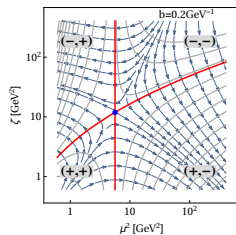
Universal scale-independent TMD

There is a unique line which passes through all μ 's

The optimal TMD distribution

$$F(x, b) = F(x, b; \mu, \zeta_\mu)$$

where ζ_μ is the special line.



TMD cross-section

$$\frac{d\sigma}{dX} = \sigma_0 \sum_f \int \frac{d^2b}{4\pi} e^{i(b \cdot q_T)} H_{ff'}(Q) \{\tilde{R}^f[b; Q]\}^2 \tilde{F}_{f \leftarrow h}(x_1, b) \tilde{F}_{f' \leftarrow h}(x_2, b),$$

with $\zeta_f = \mu_f^2 = Q^2$

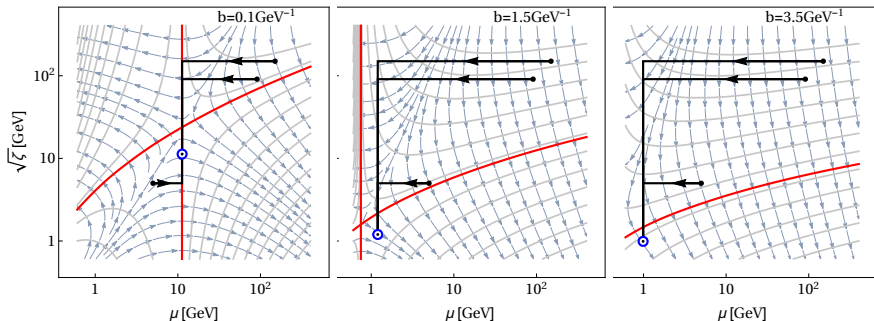
$$\tilde{R}^f[b; Q] = (Qb)^{-D_{\text{NP}}^f(Q, b)} \exp\{-D_{\text{NP}}^f(Q, b)v^f(Q, b)\}$$

- v is given perturbative series, $v = \frac{3}{2} + a_s \dots$
- \tilde{F} is TMD in the "naive" ζ -prescription

- There are no approximations (*ala* high energy expansion of integrals).
- There are only (μ_f, ζ_f) scales and no solution dependence.
- **Clear separation of TMD evolution from the model for TMD distribution.**

Evolution with b -dependent scale (CSS-like)

$(Q, Q^2) \rightarrow (\mu_b, \mu_b^2)$

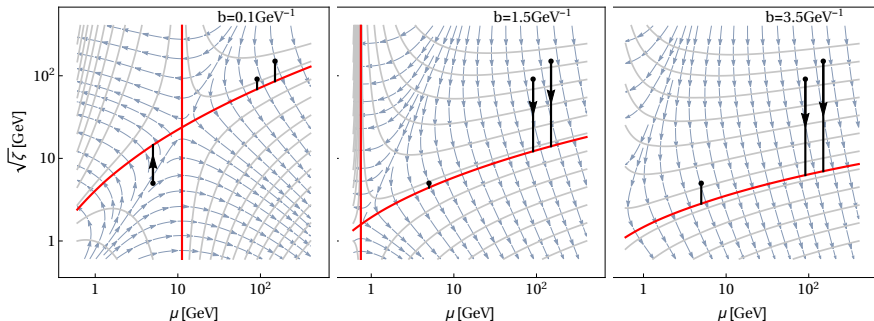


Here $\mu_b = \frac{C_0}{b^*}$ with $b_{\max} = 1.2 \text{ GeV}^{-1}$

Analogy in DIS

Scale depends on parameter $\leftrightarrow d\sigma = C(Q)R[Q \rightarrow \text{ch}(x)]f(x, \text{ch}(x))$
 PDF $f(x, \text{ch}(x))$ has no interpretation, no sense, and
 depends on the order of evolution in use.

Optimal version
 $(Q, Q^2) \rightarrow (Q, \zeta_Q)$



Analogy in DIS
 Scale (potential) is fixed $\leftrightarrow d\sigma = C(Q)R[Q \rightarrow 2\text{GeV}]f(x, 2\text{GeV})$
 PDF $f(x, 2\text{GeV})$ is just a model and
 is dependent on the order of evolution in use.

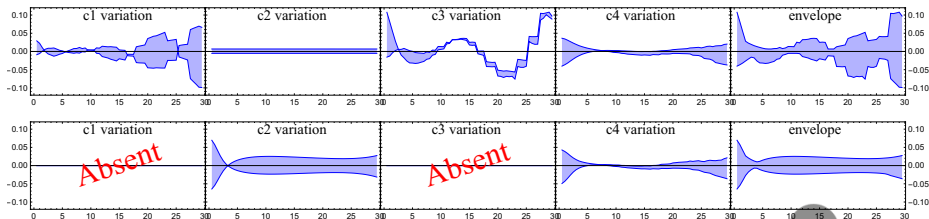
Uncertainties of TMD cross-section

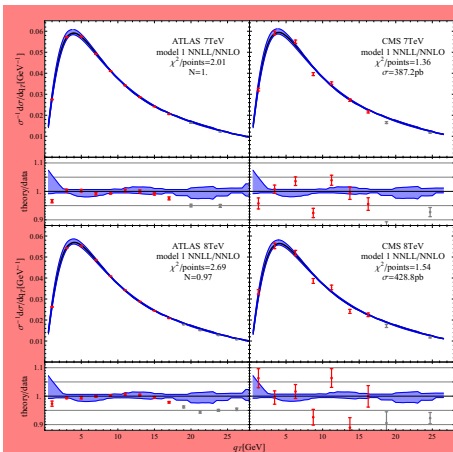
Less number of scales = less theory error

- The variation of c_1 is the variation of the evolution path (only).
- The variation of c_3 is the variation of the evolution path (almost).

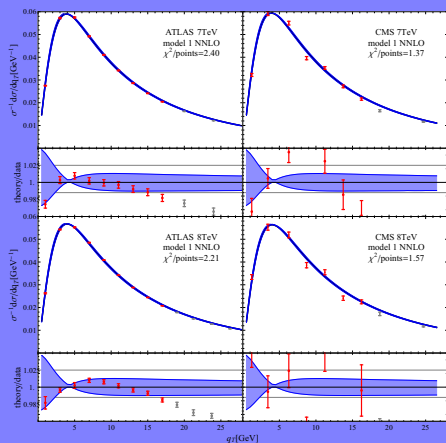
We found significant reduction of theory error band.

Z-boson production at CDF run 2



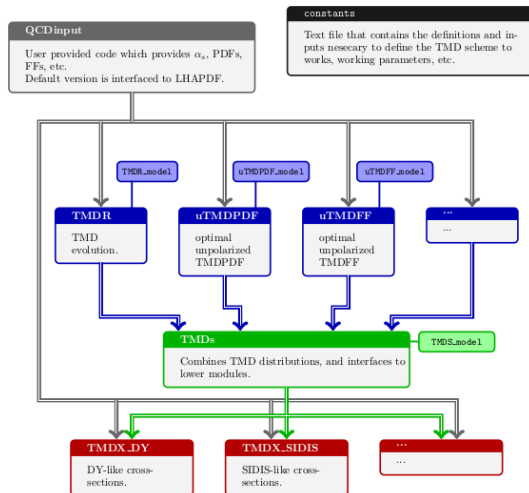
CSS-like evolution (ζ -prescription)

Optimal evolution



Update of the NNLO DY fit,
 χ^2 -values practically the same (a bit better), parameters within (previous) error-bars
 significant reduction of theory uncertainties.

arTeMiDe v1.3



- Variety of evolutions
- LO, NLO, NNLO
- No restriction for NP models
- Fast code
- DY cross-sections
- SIDIS cross-sections (not tuned yet)
- Theory uncertainty bands

<https://teorica.fis.ucm.es/artemide/>



Conclusion

Main message:

TMD evolution is a double scale evolution.

Therefore, it should be considered with care, and then it grants many simplifications.

Message 1:

TMD distributions on a same equipotential line are equivalent. Enumerate them with lines!

- Universal for all quantum numbers
- Very simple practical formula (no integrations!)
- Guaranteed absence of (large) logarithms in the matching coefficient
- TMD model is independent on evolution order.

E.g You can use NNLO unpolarized and LO Sivers together, without theory tensions

Message 2 (unpresented):

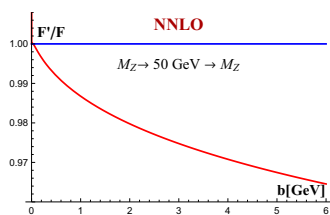
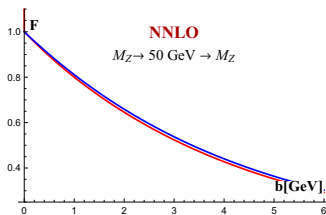
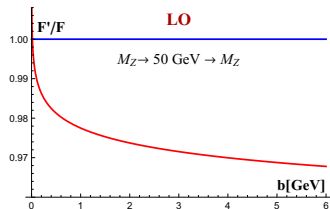
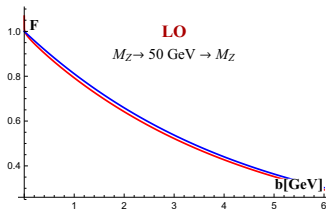
In truncated PT there is the solution-dependence of evolution (see [\[1803.11089\]](#))

- It could be strong.
- There is no unique way to fix it.

Double-scale evolution is not unique for TMD case. It also appears in k_T -resummation, joint resummation, DPDs, etc.

Problem 1: Violation of transitivity

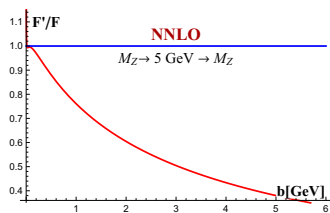
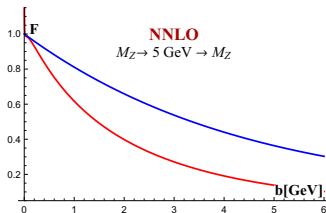
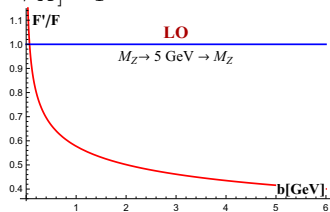
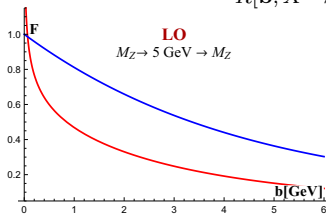
$$R[\mathbf{b}; X \rightarrow Y] R[\mathbf{b}; Y \rightarrow X] = 1$$



There is a violation of transitivity $\sim 2\%$ which seems better at NNLO

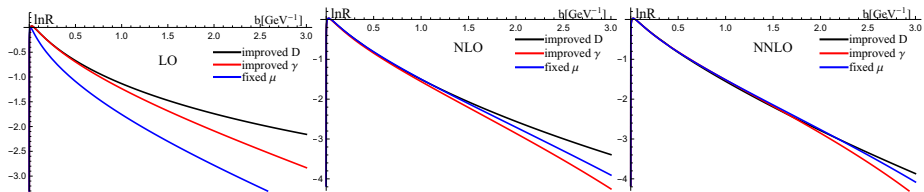
Problem 1: Violation of transitivity

$$R[\mathbf{b}; X \rightarrow Y] R[\mathbf{b}; Y \rightarrow X] \neq 1$$



There is a **VERY strong** violation of transitivity, which seems worse at NNLO

The difference between solutions is $\underbrace{\sim a^{N+1} \mathbf{L}_\mu}_{\text{main b}}$ or $\underbrace{\sim a^{N+1} \mathbf{L}_\mu \mathbf{L}_{\mu_0}^N}_{\text{large b}}$



How strong is modification of the field?

