Nonfactorization and the VdM scan method

Methods applied and lessons learnt at the CMS experiment

Joscha Knolle on behalf of the CMS Collaboration

LHC Lumi Days June 04-05, 2019





HELMHOLTZ RESEARCH FOR GRAND CHALLENGES



- VdM method assumes: beam proton densities factorize in *x* and *y*
- xy nonfactorization causes bias
- possible features:
 - "tilt"
 - \Rightarrow correlation parameter ρ
 - "stretch"
 - \Rightarrow double Gaussian



- VdM method assumes: beam proton densities factorize in *x* and *y*
- xy nonfactorization causes bias
- possible features:
 - "tilt"
 - \Rightarrow correlation parameter ϱ
 - "stretch"
 - \Rightarrow double Gaussian



- VdM method assumes: beam proton densities factorize in *x* and *y*
- xy nonfactorization causes bias
- possible features:
 - "tilt"
 - \Rightarrow correlation parameter ϱ
 - "stretch"
 - \Rightarrow double Gaussian



- VdM method assumes: beam proton densities factorize in *x* and *y*
- xy nonfactorization causes bias
- possible features:
 - "tilt"
 - \Rightarrow correlation parameter ϱ
 - "stretch"
 - \Rightarrow double Gaussian



- VdM method assumes: beam proton densities factorize in *x* and *y*
- xy nonfactorization causes bias
- possible features:
 - "tilt"
 - \Rightarrow correlation parameter ϱ
 - "stretch"
 - \Rightarrow double Gaussian



- VdM method assumes: beam proton densities factorize in *x* and *y*
- xy nonfactorization causes bias
- possible features:
 - "tilt"
 - \Rightarrow correlation parameter ϱ
 - "stretch"
 - \Rightarrow double Gaussian

Beam imaging method



- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V



- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V





- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V







- beam 2 fixed at x = 0, beam 1 moved in 19 Δx steps
 - repeat: beam 1 and 2, in x and y
- reconstruct primary vertices
 - limited by vertex resolution V







Beam imaging scans & analysis strategy



DESY. | Nonfactorization at CMS | Joscha Knolle, 04 Jun 2019

Page 7

Transverse proton density models and fit results

- analytical convolution ⊗ V requires Gaussian-based fit models
- simplest case: single 1D Gaussians

$$g(x) \cdot g(y) = \exp\left[-\frac{1}{2}\left(\frac{x^2}{\sigma_x^2} + \frac{y^2}{\sigma_y^2}\right)\right]$$

⇒ large angular and radial dependency of residuals







Transverse proton density models and fit results Fill 4266 (2015, 13 TeV) CMS Preliminary Scan #3X, BCID 771, fit model: g1 + g2 - g 0.4 analytical convolution $\otimes V$ requires relative residuals Gaussian-based fit models 0.2 y [mm] single Gaussian with correlations $g(x,y) = \exp\left[-\frac{1}{2(1-\varrho^2)}\left(\frac{x^2}{\sigma_y^2} + \frac{y^2}{\sigma_y^2} - \frac{2\varrho xy}{\sigma_y \sigma_y}\right)\right]$ -0.2 normalized sums of Gaussians -0.4 - add wider Gaussian g_2 -0.4 -0.2 n 0.2 0.4 \Rightarrow wider tails x [mm] subtract narrow Gaussian q₃ Fill 4266 (2015, 13 TeV residual \Rightarrow flattened centre CMS Preliminary Scan #3X, BCID 771, fit model: g1 + g2 - g3 \Rightarrow best-fit model: $g_1 + g_2 - g_3$ average relative ojectic radia for pp at 13 TeV in 2015, 2016 & 2017, but no sufficient description in 2018 01 02 0.3 04

Page 10

radius [mm]

Determination of VdM bias

Bias from VdM simulations

- unbiased overlap from direct integration of product of fitted proton densities
- biased overlap from repeated MC simulations of VdM scan pairs using fitted proton densities
- VdM results corrected for difference



Determination of VdM bias

Bias from VdM simulations

- unbiased overlap from direct integration of product of fitted proton densities
- biased overlap from repeated MC simulations of VdM scan pairs using fitted proton densities
- VdM results corrected for difference

Closure test with toy models

- randomly generate toy models, simulate set of beam imaging scans, and apply same fit procedure as for data
- select toys with similar fit properties as data fits, evaluate VdM bias
- reconstruction of VdM bias from good fits with precision $\lesssim 0.5\,\%$ possible



Effects impacting beam position

Beam-beam deflection



- electric repulsion of proton beams, depending on transverse separation
- causes movement of fixed beam



- beams drift away from nominal orbit
- changes beam position within step, and step size among different steps

Length scale doesn't affect fixed beam, doesn't introduce step size variations

Recently implemented methods & developing ideas

Offset scan analysis

Offset scans and general considerations



- normal VdM scans
- offset scans with beams at constant separation in non-scanning direction
- ⇒ differently affected by nonfactorization

 reconstruct luminous region profile (= product of beam proton densities) from fit to rate vs 2D beam separation

- sensitive to beam separation effects: length scale, beam-beam, orbit drift
- four separation configurations shared between two scans
 ⇒ consistency check
- tails limited by low statistics
- can use any luminometer
 ⇒ not limited by central DAQ
- offset scans only done in 2017 & 2018 VdM programmes

Offset scan analysis

Luminous region modelling and fit results



- ad-hoc position correction to reach rate agreement at coinciding steps
 ⇒ thus correcting for beam position effects
- no limitation on luminous region models
- found good description with single and double Gaussians
 ⇒ main result: ρ of main

Gaussian component

Offset scan analysis

Results on time evolution of nonfactorization



- in Fill 6868: two offset scan pairs taken ~12 hours apart
- separately analysed ⇒ test of time evolution of nonfactorization
- \Rightarrow correlation parameter ϱ of luminous region profile increases with time
- ⇒ consistently observed in beam imaging method



Beamspot evolution analysis

- at each scan step, fit 3D ellipsoid to vertex distribution. including vertex resolution
- extract observables: mean, width, tilt
- analytically compute predictions from models for observables
- fit predictions to rate & beamspot data
- plots: fit results with $g_1 \pm g_2 \pm g_3$ for Scan #1X. Fill 4954 (2016, 13 TeV)



Data

0.6

Data

Eit

Ei+

Diagonal scan analysis

- VdM scans in X, Y, X+Y, X−Y, analysed with standard method ⇒ obtain four widths
- fit ellipse to obtain 1σ contour line of luminous region profile
 ⇒ can extract correlation parameter ρ
- diagonal scans only done in 2018 PbPb VdM programme





Summary



- In Run 2, beam imaging is the main method of evaluating nonfactorization.
- Complementary methods are being developed:
 - offset scans,
 - beamspot evolution,
 - diagonal scans.
- Improved nonfactorization results are underway.
- · Essential next steps:
 - direct comparison of results from different methods,
 - toy simulations including all methods and effects.

References

CMS luminosity measurements

- CMS-PAS-LUM-15-001 (pp 13 TeV 2015)
- CMS-PAS-LUM-16-001 (pp 5.02 TeV 2015)
- CMS-PAS-LUM-17-001 (pp 13 TeV 2016)
- CMS-PAS-LUM-17-002 (Pbp/pPb 8.16 TeV 2016)
- CMS-PAS-LUM-17-004 (pp 13 TeV 2017)
- CMS-PAS-LUM-18-002 (pp 13 TeV 2018)

Method descriptions

- Klute, Medlock, and Salfeld-Nebgen, JINST **12** (2017) P03018 (beam imaging method)
- Chatrchyan et al. (CMS Collaboration), JINST 9 (2014) P10009 (vertex splitting)
- Aaboud et al. (ATLAS Collaboration), Eur. Phys J. C 76 (2016) 653 (offset scan & beamspot evolution analysis)
- CMS Collaboration, CMS-DP-2016-051 (beamspot fits)
- Webb, CERN-THESIS-2015-054 (beamspot evolution analysis)

Backup

Reconstructed bias for different bunch crossings



 $\Rightarrow\,$ we use average of bias from five BCIDs as correction for all BCIDs

