

Exploring machine learning methods for unbinned data analysis of Drell-Yan phase space at CMS

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Introduction

Particle detectors introduce smearing in measurements:

- Finite detector resolution
- Acceptance cuts (geometrical and physical)
- Binning effect in distributions

Goal: Use data (at detector level) and infer true desitributions (at particle level).



Only then we can reliably compare the measurements against other sources:

- Semi-Inclusive theoretical predictions
- MC generator parameter tuning
- Other experiments

Unfolding (deconvolution).



Current Unfolding approach

Detector response can be modeled as a linear application.

$$x_{detector} = R * x_{truth} + x_{backg} \tag{1}$$

Where *R* represents the detector **response** as $R_{ij} = P(truthi | measurej)$.

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Various approaches exist to unfolding:

• Naive R inversion

$$x_{truth} = R^{-1} * (x_{detector} - x_{backg})$$
(2)

• Iterative Bayesian Unfolding (IBU): Given a **response** matrix and a prior distribution.

$$x_{j}^{(n)} = \sum_{i} P_{n-1}(truthi|measurej) * P(measurej)$$
(3)

Unfolding drawbacks:

- Works for binned data
- 1-dimensional by desing (very difficult to extend to higher dimensions)
- Does not account for relations between observables

Machine Learning approach

Distribution reweight.

- Using a binary classifier to distinguish between two distributions
- Likelihood ratio can be aproximated and used as weight function

$$w_{A\to B} = \frac{p_B(x)}{p_A x} \approx \frac{f(x)}{1 - f(x)} \tag{4}$$

 Distributions are binned for representation but weights are estimated per event

Machine Learning approach



Figure 1: Transverse momentum distribution. $pp \rightarrow DY7 TeV$.

Advantages of ML reweight.

- Used for non-trivial distributions
- Easily extended to multiple dimensions
- Learns complex relations between observables

Machine Learning approach

Iterative Neural Network Reweight: OmniFold.



Iterative Neural Network Reweight: OmniFold.

- Continuous generalization of IBU
- Inherits all the advantages from ML reweight
- Unfolds experimental data using multiple dimensions (possibly full phase space)

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Work in progess

Drell-Yan phenomenology analysis with CMS data.

- Explore HEP analysis viavility with Python
- Unlock ML ecosystem and methods
- Validation of NanoAODRun1 format for CMS data
- Improve accuaracy of infered distributions





Figure 2: Syntetic and Natural distributions.





Future work.

- Full phase space unfolding
- Unbinned and data driven analysis
- Evaluate statistical and systematical uncertanties
- Search for more robust observables where current theoretical predictions struggle



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