

Investigations on unfolding with RooUnfold

Patrick L.S. Connor

Deutsches Elektronen-Synchrotron

29 September 2016



In the current slides, a code for unfolding studies is described¹:

- requires latest RooUnfold version and Root6
- resolution and pt spectrum to be defined by the user
→ trivial RM can also be defined
- compares different unfolding methods available in RooUnfold
→ at least two methods should give similar results

Tests on MadGraph, Pythia8 and Run2015CD can be found in the other presentation on the same slot.

Unfolding methods

Introduction

Description

Unfolding
methods

Tool

Examples

Early

conclusions

bin/bin correct central values but wrong handling of statistical uncertainties

Bayes iterative procedure, good experiences in previous analyses, converges but unknown number of iterations

Inversion best on principle, but possible instabilities if statistics is too low

TUnfold likelihood minimisation, including regularisation, recommended by statistics comitee, developer is at DESY

SVD ...

Introduction

Description

Unfolding
methods**Tool**

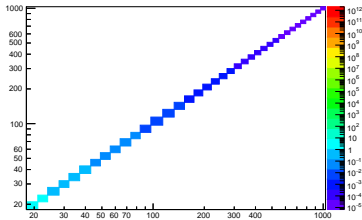
Examples

Early

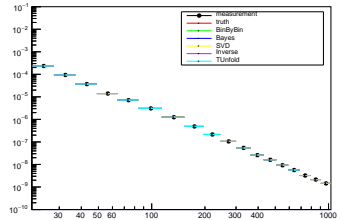
conclusions

- works with latest RooUnfold version in Root6
- produces a canvas with RM, resolution in pt bins, statistically independent reconstructed spectrum, ratio of the different unfolding methods with the "measured" spectrum
- $\sigma = 1/p_T^4$
- binning can easily be changed, shown in the following with the jet standard binning
- only a trivial and a gaussian smearing have been tried so far

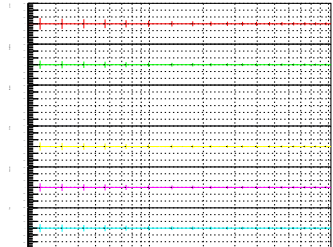
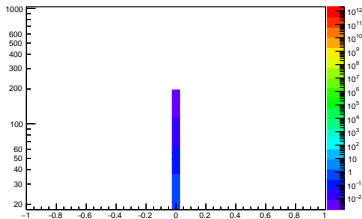
RM



measurement



resolution



Unfolding

Patrick
Connor

Introduction

Description

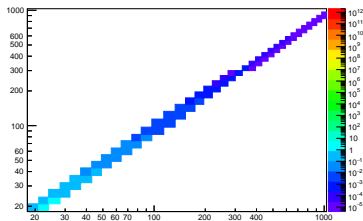
Unfolding
methods

Tool

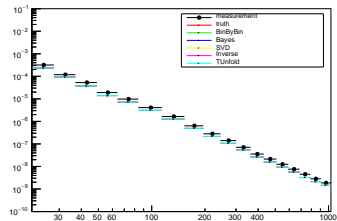
Examples

Early
conclusions

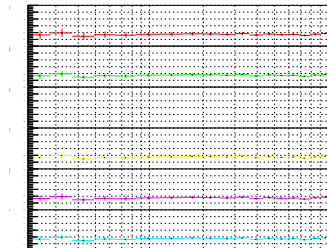
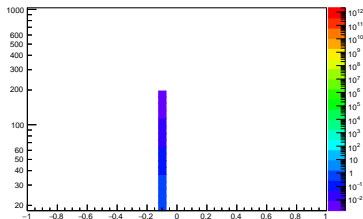
RM



measurement



resolution



Unfolding

Patrick
Connor

Introduction

Description

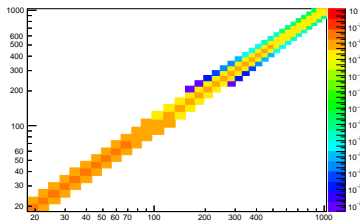
Unfolding
methods

Tool

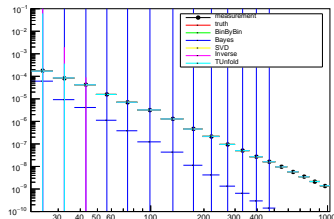
Examples

Early
conclusions

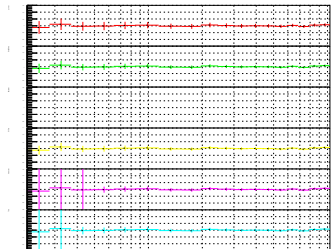
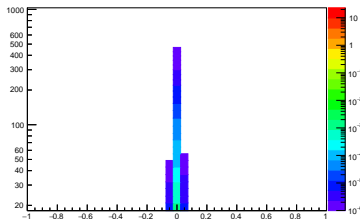
RM

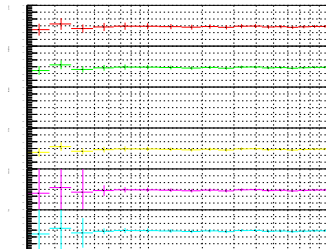
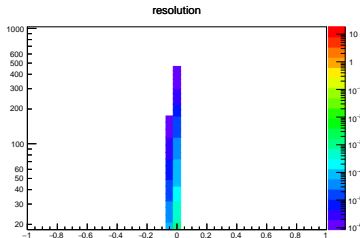
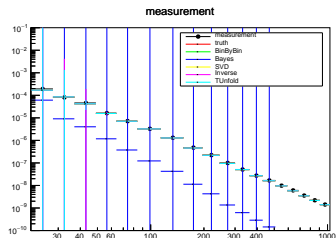
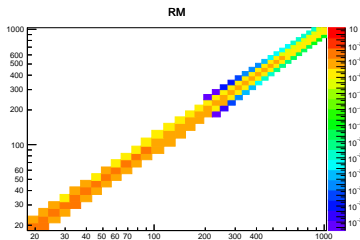


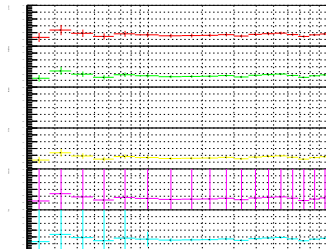
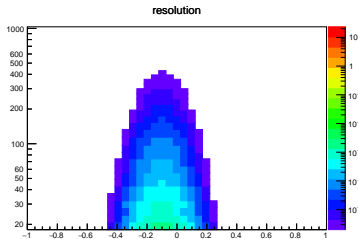
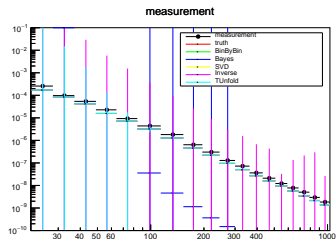
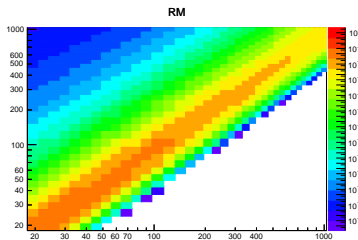
measurement



resolution







Early conclusions

Introduction

Description

Unfolding
methods
Tool
ExamplesEarly
conclusions

From this unfolding model and from previous investigations (see other attachment to the indico page)

- no unfolding method looks 100% reliable in any conditions
→ find the conditions under which they are reliable
- parameters that can matter:
 - ① normalisation (only for Bayes)
 - ② binning (especially for Bayes and Inversion)
 - ③ uncertainties (especially for Inversion and TUnfold)