



Unfolding in ALICE

Jan Fiete Grosse-Oetringhaus, CERN
for the ALICE collaboration

PHYSTAT 2011
CERN, January 2011



Content

- Unfolding methods used in ALICE
- Applications of unfolding in ALICE
 - Observations, Issues and remarks

Unfolding Methods

- χ^2 -Minimization with regularization (only 1D)

$$\chi^2(T^*) = \sum_m \left(\frac{M_m^* - \sum_t R_{mt} T_t^*}{e_m} \right)^2 + \beta R(T^*) \quad R(T) = \sum_t (a_t)^2$$

β weight

- Available regularizations

- Mostly used:
linear and constant

$$a_t = \frac{T_t'}{\sqrt{T_t}} = \frac{T_t - T_{t-1}}{\sqrt{T_t}} \quad \text{prefer constant}$$

$$a_t = \frac{T_t''}{\sqrt{T_t}} = \frac{T_{t-1} + 2T_t - T_{t+1}}{\sqrt{T_t}} \quad \text{prefer linear least curvature}$$

$$a_t = \frac{\hat{T}_t''}{\sqrt{\hat{T}_t}} \quad \hat{T}_t = \ln T_t \quad \text{prefer exp}$$

- Implementation uses
Minuit, Migrad

- Uncertainties from Minuit

$$R(T) = \sum_t T_t \ln \frac{T_t}{\varepsilon_t} \quad \text{reduced cross-entropy}$$

Unfolding Methods (2)

- Iterative Bayesian Unfolding (1D + nD)

$$\tilde{R}_{tm} = \frac{R_{mt}P_t}{\sum_{t'} R_{mt}P_{t'}} \quad U_t = \sum_m \tilde{R}_{tm}M_m$$

d'Agostini

- Optionally smoothing can be applied

$$\hat{U}_t = (1-\alpha)U_t + \frac{\alpha}{3}(U_{t-1} + U_t + U_{t+1})$$

- Uncertainty on unfolded distribution by randomization of input spectrum
 - Poisson distribution used per bin

Unfolding Methods (3)

- Unfolding methods part of the software framework
 - Main interface function (for 1D case):
Unfold(TH2* correlation, TH1* efficiency, TH1* measured, TH1* initialConditions, TH1* result)
 - nD case uses THnSparse
- Functions to evaluate bias b_t due to regularization

$$b_t = \sum_m \frac{\partial T_t}{\partial M_m} ((RT)_m - M_m) \quad \text{(see e.g. Cowan)}$$

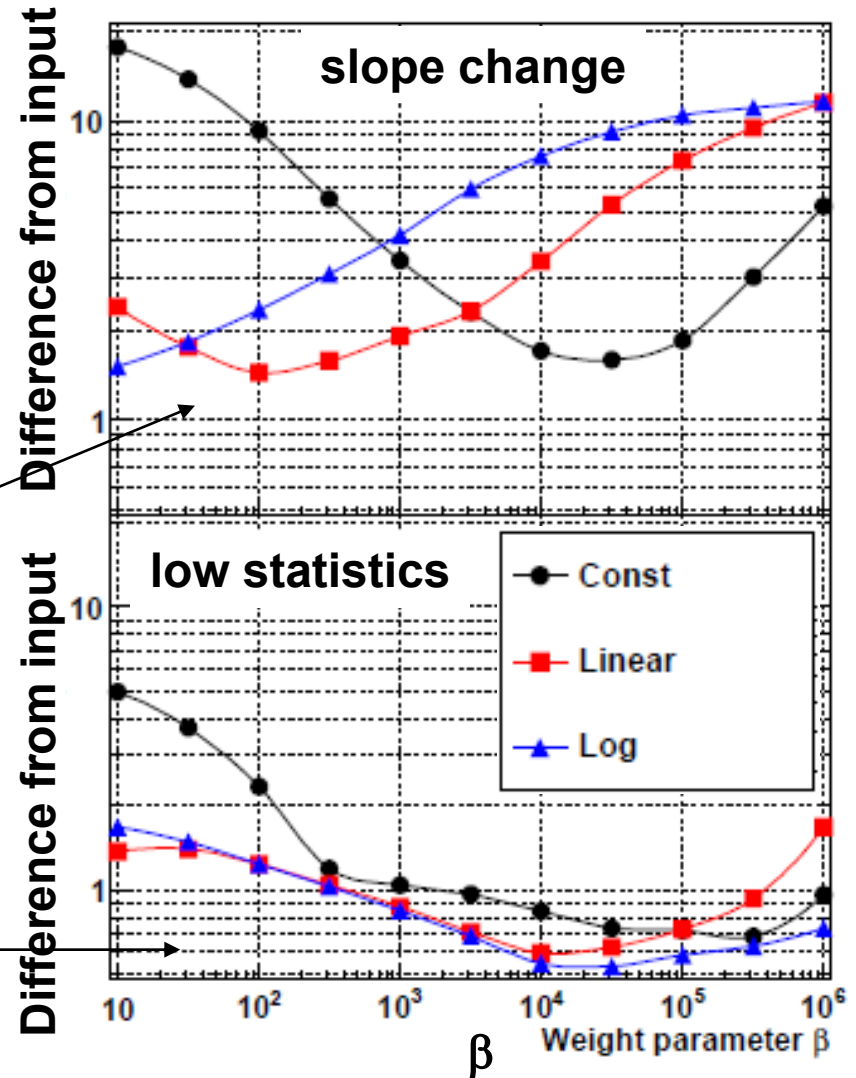
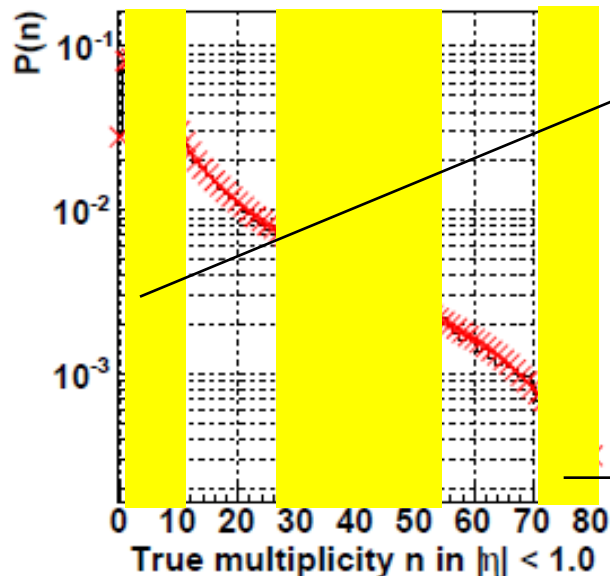
- Derivate calculated numerically

$$\frac{\partial T_t}{\partial M_m} = \frac{1}{6d} \left[8 \left(f\left(\frac{d}{2}\right) - f\left(-\frac{d}{2}\right) \right) - (f(d) - f(-d)) \right]$$

$$f(x) = T_t(M | M_m = M_m + x\sqrt{M_m})$$

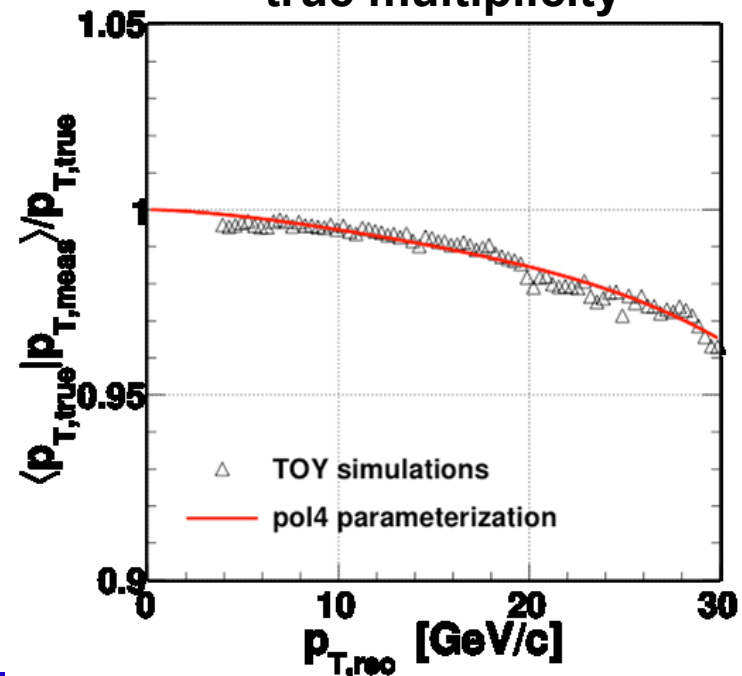
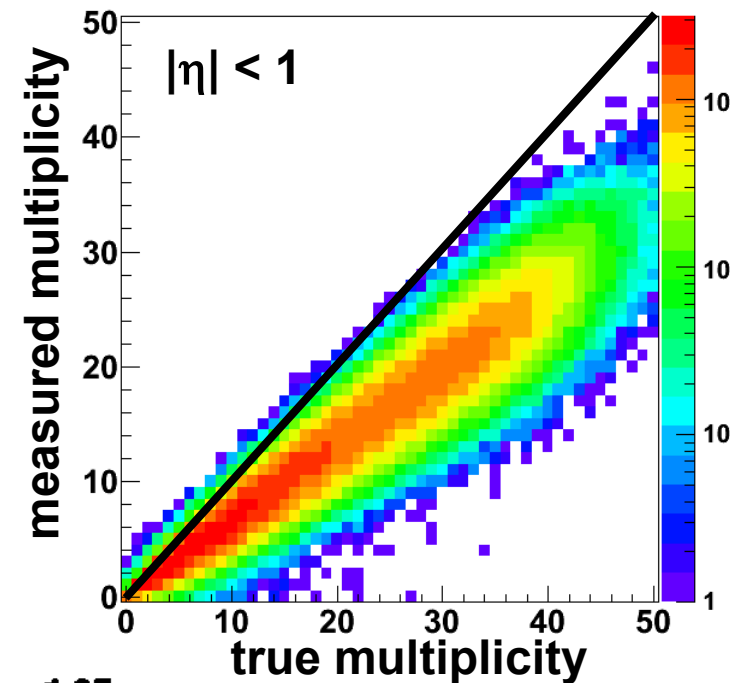
Example: Evaluate Regularization Weight with MC

- Cooking down the consistency of the unfolded solution with the MC to one number is tricky
 - Different regions, different qualitative shape



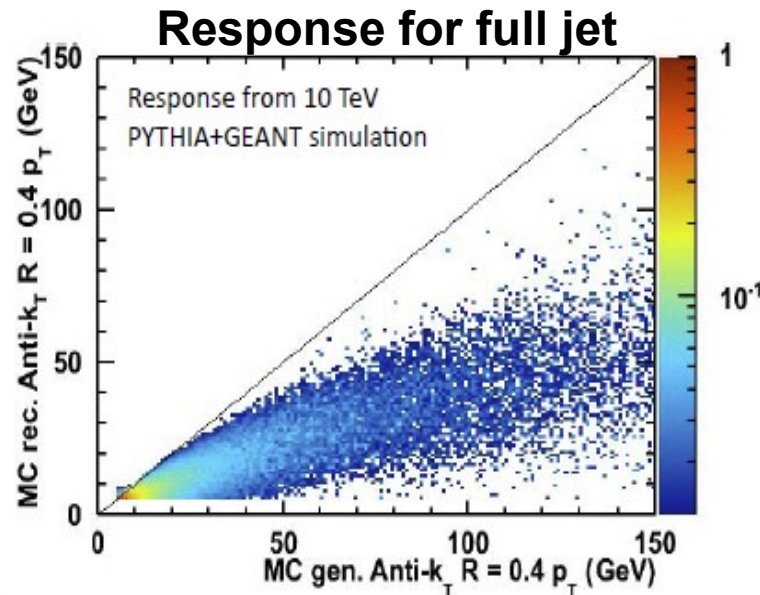
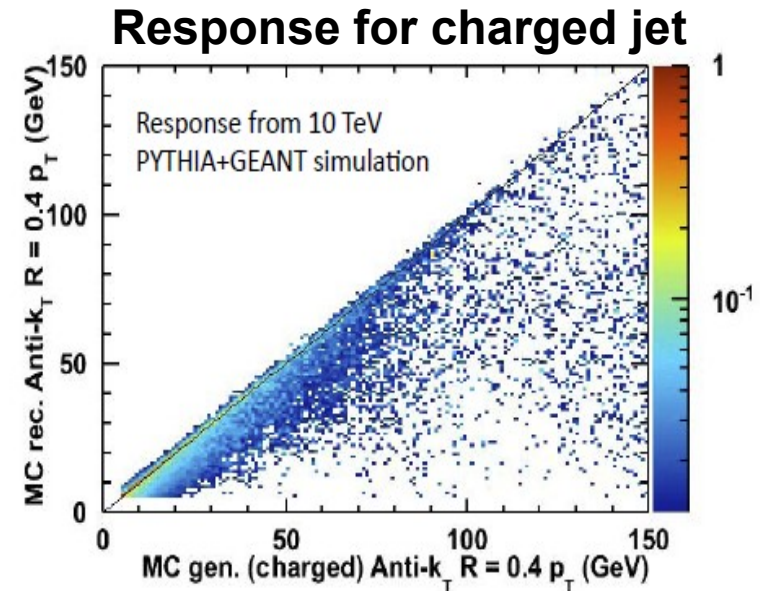
Applications

- Multiplicity distribution
 - Efficiency $\sim 70\%$ (inactive modules in pixel detector due to cooling issues)
 - far off the diagonal
 - Significant spread in response matrix → wide correlations in unfolded spectrum
- p_T spectrum
 - Shift is 2-4% for 20-30 GeV
 - Unfolding needed, response matrix not very wide



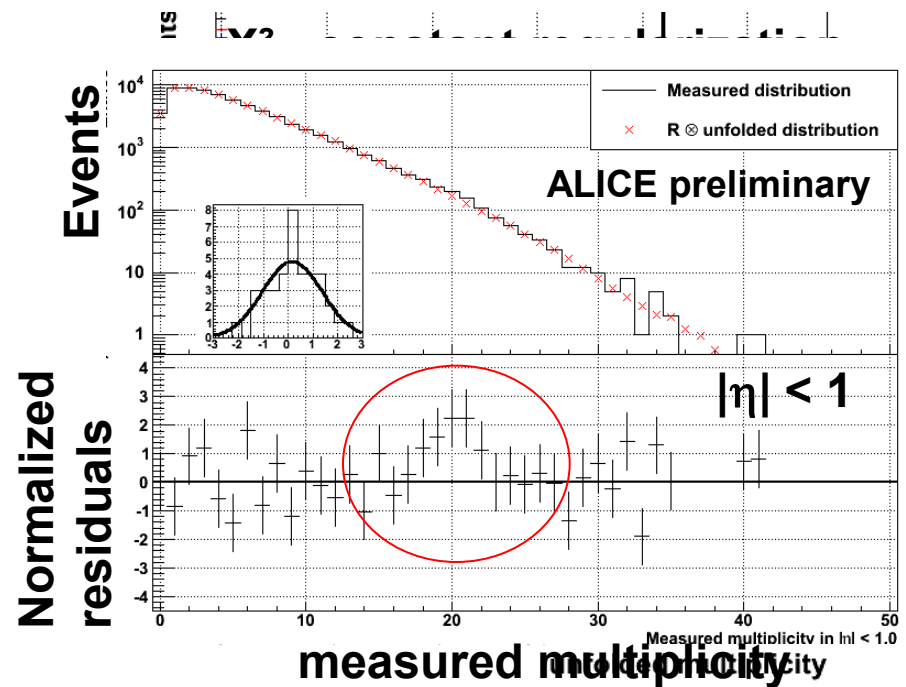
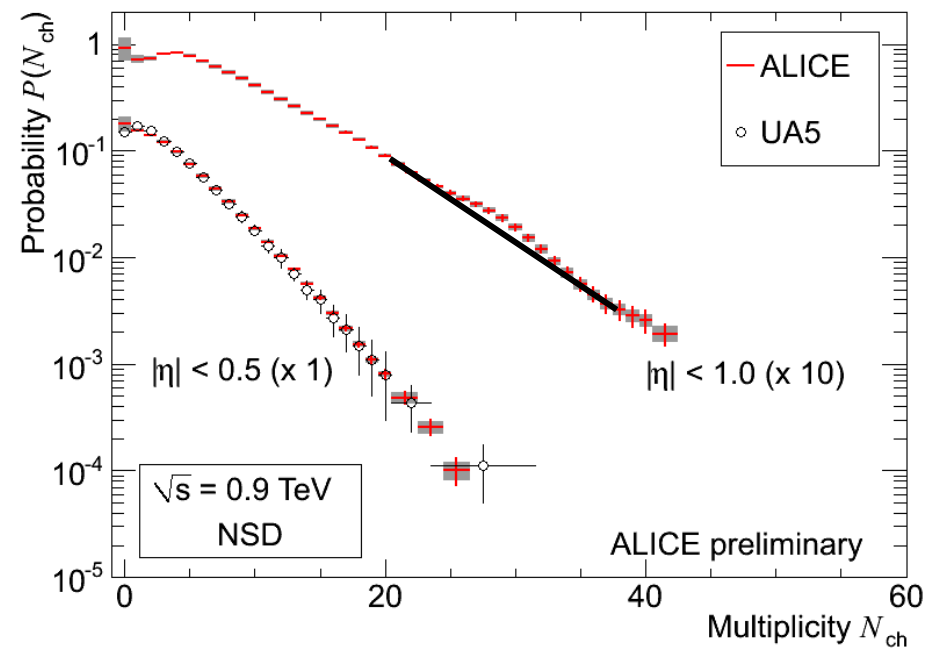
Applications (2)

- Jet spectrum
 - Wide response matrix $\sigma \sim 20\%$
 - Corrections to full (charged+neutral) jet \rightarrow significant shift because all neutrals missing (1/3)
 - Caveat: MC simulations usually done in $p_{T, \text{hard}}$ bins
 - Jet yield at low $p_T \gg$ MC at low p_T



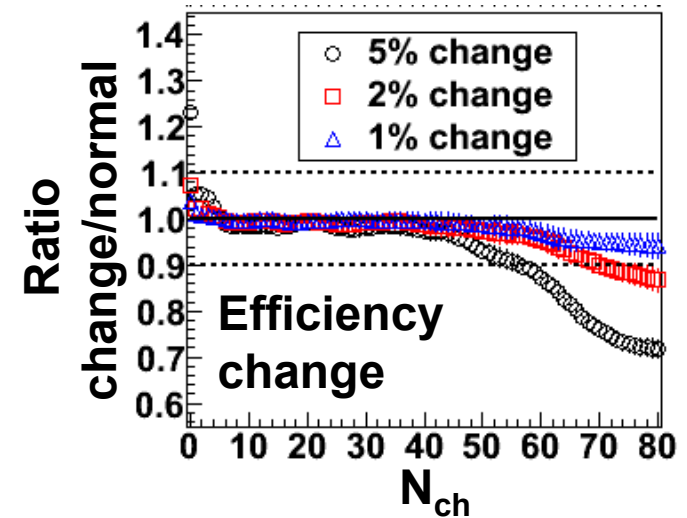
Example Issue

- Observation: nice hump in unfolded distribution (which got people immediately excited)
- Usual cross-checks → independent of
 - Regularization scheme
 - Unfolding method
- Forcing the distribution to be exponential
 - Moves the structure to the residuals
 - However, in fewer bins and (visually) much less significant
- Uncertainties of unfolded spectrum and wide response matrix can enhance fluctuations in the data significantly
- Slope change visible in measured data → transition gets spread out by regularization



Systematic Uncertainties

- From the unfolding procedure
 - Bias
 - Some fluctuations (1-5%) remain
 - Use MC to get a feeling
- Uncertainties are correlated between the bins
- Due to the response matrix
 - Uncertainties on the response matrix "propagate through" the unfolding into the unfolded distribution
 - Create different response matrices resembling the uncertainty
 - E.g. detector efficiency uncertainty of 1% → Create three response matrices with -1%, 0, +1% efficiency w.r.t. nominal
 - Use difference in unfolded distribution as systematic uncertainty → tricky because overlaid by "typical" fluctuations, small effects cannot be disentangled



Summary

- ALICE uses unfolding (up to now) for multiplicity distribution, p_T spectrum and jet spectrum
- The ALICE software framework provides
 - χ^2 -minimization with regularization
 - Iterative Bayesian unfolding
- Unfolding usually requires a lot of tuning
- Systematic uncertainties sometimes tricky
- Uncertainties on unfolded distributions are always understood (by others) as single bin uncertainties
 - Can we invent a nice way to visualize that this is not the case?

Thanks for help with preparing this talk to Christian Klein-Boesing, Jan Rak

Unfolding tutorial (introductory slides, exercises and solutions):
www.cern.ch/jgrosse/permanent/unfolding
(prepared for a Helmholtz power week)
Feel free to use it!